Air Force Institute of Technology AFIT Scholar

Theses and Dissertations

Student Graduate Works

3-10-2006

# Evaluation of Chlorinated Solvent Removal Efficiency among Three Wetland Plant Species: A Mesocom Study

Jun Yan

Follow this and additional works at: https://scholar.afit.edu/etd

Part of the Environmental Chemistry Commons, and the Environmental Sciences Commons

## **Recommended Citation**

Yan, Jun, "Evaluation of Chlorinated Solvent Removal Efficiency among Three Wetland Plant Species: A Mesocom Study" (2006). *Theses and Dissertations*. 3395. https://scholar.afit.edu/etd/3395

This Thesis is brought to you for free and open access by the Student Graduate Works at AFIT Scholar. It has been accepted for inclusion in Theses and Dissertations by an authorized administrator of AFIT Scholar. For more information, please contact richard.mansfield@afit.edu.





# EVALUATION OF CHLORINATED SOLVENT

# **REMOVAL EFFICIENCY AMONG THREE**

# WETLAND PLANT SPECIES: A MESOCOM STUDY

THESIS

Jun Yan, 2<sup>nd</sup> Lt, USAF

AFIT/GEM/ENV/06M-18

# DEPARTMENT OF THE AIR FORCE AIR UNIVERSITY AIR FORCE INSTITUTE OF TECHNOLOGY

Wright-Patterson Air Force Base, Ohio

APPROVED FOR PUBLIC RELEASE; DISTRIBUTION UNLIMITED.



www.manaraa.com

The views expressed in this thesis are those of the author and do not reflect the official policy or position of the United States Air Force, Department of Defense, or the U.S. Government.



# EVALUATION OF CHLORINATED SOLVENT REMOVAL EFFICIENCY AMONG THREE WETLAND PLANT SPECIES: A MESOCOM STUDY

# THESIS

Presented to the Faculty

Department of Systems and Engineering Management

Graduate School of Engineering and Management

Air Force Institute of Technology

Air University

Air Education and Training Command

In Partial Fulfillment of the Requirements for the

Degree of Master of Science in Engineering Management

Jun Yan, BS

2<sup>nd</sup> Lt, USAF

March 2006

# APPROVED FOR PUBLIC RELEASE; DISTRIBUTION UNLIMITED.



# EVALUATION OF CHLORINATED SOLVENT REMOVAL EFFICIENCY AMONG THREE WETLAND PLANT SPECIES: A MESOCOM STUDY

Jun Yan, B.S. 2<sup>nd</sup> LT, USAF

Approved:

/signed/

10 March 2006

date

Charles A. Bleckmann (Chairman)

/signed/

James P. Amon (Member)

/signed/

Abinash Agrawal (Member)

10 March 2006

date

10 March 2006

date



#### AFIT/GEM/ENV/06M-18

#### Abstract

The purpose of this research was to study three different wetland plant species and to compare the chlorinated solvent removal efficiency among each species. Each plant has a different characteristic favorable for chlorinated solvent degradation. *Eleocharis erythropoda* (Spike Rush) are plants with thin tube like leaves and large root mass. Carex Comosa (longhaired Sedge) has broad leaves and Scirpus Atrovirens (Green Bulrush) are broad leafed wetland plants with a long flowering stem. Previous research had shown that wetlands were effective at degradating PCE. However, wetlands are composed of many different plant species and it is unknown which species are most effective at pollutant remediation. In order to study individual plants, twelve PVC column reactors had been built and each column has an upward flowing ground water scheme similar to both the constructed wetland and natural fen wetlands. Each column will be planted with one individual plant species and low concentrations of PCE will be injected into the plant mesocosm. It is my hypothesis that PCE will be degraded into daughter products in all the mesocosms. However, this experiment will attempt identify which plant is the most efficient at PCE degradation.



## Acknowledgments

I would like to express my sincere appreciation to my advisor Dr. Amon for his continual help, advice and guidance through out this entire thesis process. His insights were pivotal in the completion of thesis. I am also very grateful for Dr. Agrawal's proactive involvement in my research. His expertise in Gas Chromatograph analysis was a tremedous help. I would also like to thank Dr. Shelley and Dr. Bleckmann for providing the tools and encouragement that propelled me to the very end.

Lastly but not least, I would like to thank my beautiful wife, who never gave up on me and gave me the joy I needed to carryout this endeavor.

Jun Yan



Page
Abstract iv
Acknowledgmentsv
Table of Contents vi
List of Figures ix
List of Tables
I. Introduction1
Background1
Conventional Treatment Methods
Bioremediation4
Phytoremediation7
Wetlands10
Problem Statement
Research Objective and Hypothesis
Research Question
II. Literature Review
Natural Attenuation14
Thermodynamics14
Bioremediation
Phytoremediation21
III. Methods
Column Construction
Planting Timeline
Syringe Pump Injection Rate
Inflow Rate

# **Table of Contents**



	Water and Organic content:	Page30
	Standard Preparation	31
	Sampling Schedule	
	Gas Chromatograph	34
	Samples for IC	24
	Desidence Time Concelling	
	Residence Time Sampling	
	Ion Chromatograph	35
	General Supplies	
IV.	Results	
	Pre PCE Injection Baseline Results	37
	Time Series Data Sets	
	PCE Concentration:	51
	TCE Data	54
	TCE and PCE Comparison	56
	Sulfate and Nitrate Comparison	59
	Conductivity	62
	Residence Time	63
V. ]	Discussion	65
	Pre and Post Methane comparison	66
	PCE concentration Decrease	67
	6 <sup>th</sup> Week	70
	TCE:	73
	Sulfate and Nitrate comparison	76



Methane and TCE	Page
VI Conclusion	
Answers to specific research questions	
Limitations	
Appendices	
Bibliography	
Vita 108	



# List of Figures

Page
Figure 1.1 PCE is degraded to TCE in a reductive dechlorination reaction
Figure 1.2. PCE reduction pathway6
Figure 1.3. Phytoremediation mechanism9
Figure 1.4. Constructed wetland with three perforated PVC pipes at the bottom11
Figure 1.5. Cross-section of wetland soil layers12
Figure 2.1 Redox potential of possible subsurface reactions17
Figure 2.2 Underground chemical composition change with time
Figure 3.1 Mesocosm placement within greenhouse25
Figure 3.5 Experimental set up27
Figure 4.1 Background methane concentration pre-PCE injections
Figure 4.2 Average PCE Concentration for first sample set
Figure 4.4 Average methane concentration for the first sample set41
Figure 4.5 Methane comparison in (mmol/L) pre and post PCE injections
Figure 4.6 PCE comparisons 2 and 4 weeks after continuous PCE injections began43
Figure 4.7 TCE comparisons 2 and 4 weeks after continuous PCE injections began44
Figure 4.8 PCE data 2 weeks after injections began46
Figure 4.9 PCE data 4 weeks after injections began46
Figure 4.10 PCE data 6 weeks after injections began46
Figure 4.11 TCE data 2 weeks after injections began47
Figure 4.12 TCE data 4 weeks after injections began47
Figure 4.13 TCE data 6 weeks after injections began
Figure 4.14 Carex comosa methane comparison



P Figure 4.15 Scirpus atrovirens methane comparison	age 50
Figure 4.16 <i>Eleocharis erythropoda</i> methane comparison	50
Figure 4.17 Control methane comparison	50
Figure 4.18 PCE Concentration decrease between ports after 2 weeks	51
Figure 4.19 Cumulative percentage of PCE removed at 2 weeks	52
Figure 4.20 PCE Concentration decrease between ports at 6 weeks	53
Figure 4.21 Percentage of PCE removed at 6 weeks	54
Figure 4.22 TCE concentration at 2 weeks	55
Figure 4.23 TCE Concentration 6 weeks after continuous PCE injection began	56
Figure 4.24 <i>Carex comosa</i> PCE-TCE comparison	57
Figure 4.25 Scirpus atrovirens PCE-TCE comparison	57
Figure 4.26 <i>Eleocharis erythropoda</i> PCE-TCE comparison	58
Figure 4.27 Control PCE – TCE comparison	58
Figure 4.28 Average nitrate concentrations with depth	59
Figure 4.29 Average Sulfate concentrations for all four treatments	61
Figure 4.30 Average conductivity for each treatment	63
Figure 4.31 KBr breakthrough measurement	64
Figure 5.1 PCE & TCE comparison at week 2.	69
Figure 5.2 6th week comparison of PCE removal between 0" and 9"	70
Figure 5.3 6th week PCE concentration decrease between 0" to 54"	71
Figure 5.4 6th Week PCE removed between 0" and 15"	72
Figure 5.5 Average TCE concentrations from port 7	74



Figure 5.6 Overlay of nitrate trend with 6th week TCE concentration	Page 77
Figure 5.7 Overlay of sulfate trend with 6th week TCE concentration	78
Figure 5.8 Overlay of Carex comosa reactor methane and TCE concentration	81
Figure 5.9 Overlay of Scirpus atroviren reactor methane and TCE concentration	81
Figure 5.9 Overlay of Eleocharis erythropoda reactor methane and TCE concentration	on82
Figure 5.9 Overlay of control reactor methane and TCE concentration	82



# List of Tables

Table 1.1 . Physical properties of chlorinated solvents	Page
Table 1.2. EPA's MCL and potential health effects of chlorinated solvents	2
Table 2.1 Delta G of electron ACCEPTOR half cell reactions	15
Table 2.2 Delta G of electron DONOR half cell reaction	15
Table 3.1 Plant species and number of reactors	
Table 3.2 Plant and control column placement	
Table 3.3 Initial PCE injection calibration result	
Table 3.3 Tubing characteristic and Flow rate	30
Table 3.4 Effluent flow rate calculate over 36 hrs.	
Table 3.5 Water content and organic content of soil.	
Table 3.6: Stock solution Prep	
Table 3.7 Detection limits on the GC	



# EVALUATION OF CHLORINATED SOLVENT DEGRADATION EFFICIENCY AMONG THREE WETLAND PLANT SPECIES: A MESOCOM STUDY

## I. Introduction

## Background

Tetrachloroethylene (PCE) and its degradation products: Trichloroethylene (TCE), Dichloroethylene (DCE), and Vinyl chloride (VC), are chlorinated solvents belonging to a class of very persistent and toxic environmental pollutants. They were frequently used after WWII as industrial degreasing and dry cleaning agents (Pankow and Cherry, 1996). However, rampant use and improper disposal have led to widespread contamination (Freedman and Gossett, 1989). Chlorinated solvents enter the environment through dispersive loss during use, improper waste dispersal, and accidental spills (Bouwer, 1992). These chlorinated solvents are DNAPLS, allowing them to sink below the water table to form pools at the bottom of an aquifer (Masters, 1997). These pools of contaminants also have low absolute solubility. PCE for example, has an absolute solubility of around 150 mg/L (Yaws, 2004), which means that pools of chlorinated solvents will persist in groundwater aquifers for decades. However, PCE is just soluble enough to contaminate any surrounding groundwater with concentrations far exceeding EPA's Maximum Contaminant Level (MCL) of just 0.005 mg/L (Johnson and Pankow, 1992). Therefore, these properties make chlorinated solvents a group of very persistent and widespread environmental pollutants (Table 1.1).

The abundance of PCE and TCE in groundwater is of particular concern due to their carcinogenic properties (Fan, 1988). EPA started to regulate chlorinated solvents in



www.manaraa.com

the Safe Drinking Water Act of 1986 (Freedman and Gossett, 1989) and passed maximum contaminant level standards to protect public health while balancing economic feasibility (EPA, 2002). These pollution standards dictate the clean up response of chlorinated solvents and Table 1.2 below lists some of the EPA standards concerning chlorinated ethenes.

	Density	Solubility	Henry's Const	Log Kow	Vapor Pressure
Compound	(g/mL) at 25 oC	(mg/L)	(atm*m^3/mol)		(mm Hg)
Tetrachloroethylene	1.613	150	0.0269	3.4	17.8
Trichloroethylene	1.458	1,100	0.0116	2.42	57.9
cis-I,2-Dichloroethylene	1.265	3,500	0.0074	1.86	208
trans-I,2-Dichloroethylene	1.244	6,300	0.0067	2.09	324
1,1-Dichloroethylene	1.117	3,345	0.0228	2.13	600
Vinyl Chloride	0.903	2,697	0.0224	1.62	2,660

Table 1.1. Physical properties of chlorinated solvents (Yaws, 2004)

Table 1.2. EPA's MCL a	nd potential health	effects of chlorinated	solvents (	EPA.	2002)
10010 1121 2110 1102 0					/

Contaminant	<u>MCLG</u> (mg/L)	<u>MCL</u> (mg/L)	Potential Health Effects from Ingestion of Water	Sources of Contaminant in Drinking Water
Tetrachloroethylene	zero	0.005	Liver problems; increased risk of	Discharge from factories and dry cleaners
Trichloroethylene	zero	0.005	Liver problems; increased risk of	Discharge from metal degreasing sites and other
<u>.1-Dichloroethylene</u>	0.007	0.007	Liver problems	Discharge from industrial chemical factories
<u>cis-1,2-</u> Dichloroethylene	0.07	0.07	Liver problems	Discharge from industrial chemical factories
<u>trans-1,2-</u> Dichloroethylene	0.1	0.1	Liver problems	Discharge from industrial chemical factories
Vinyl chloride	zero	0.002	Increased risk of cancer	Leaching from PVC pipes; discharge from plastic

Unfortunately many military sites are in violation of EPA standards. Since 1998, 1400 groundwater contaminants sites were placed on the Nation Priorities List (NPL) and TCE was the most frequently detected chemical on these sites (Lee, 1998). Out of the 1400 installations listed on the NPL, 126 were under the direct responsibility of the DoD (USGAO, 1995). Some examples of DoD installations are current military bases,



decommissioned bases, or storage depots. The total clean up cost for the DoD is estimated to be upwards of \$30 billion (Astin and Sanders, 1996).

#### **Conventional Treatment Methods**

Previous chlorinated solvent removal processes involved the use of the "pumpand-treat" method to pump contaminated groundwater to the surface for treatment above ground. This method was similar to conventional drinking water treatment and this methodology had been well documented (EPA, 1996). However, the "pump-and-treat" method has several disadvantages. First, the physical treatments used by this method only concentrate PCE into another medium and does not promote the degradation of PCE. Second, chlorinated solvents are denser than water and have low capillary pressure, thus allowing them to seep into pore spaces along the bottom of an aquifer. The seepage into pore spaces reduces the availability of contaminants and increases the duration of cleanup to decades, possibly even centuries (Pankow and Cherry, 1996). Finally, above ground treatment facilities are expensive to construct and have continual high operating and maintenance cost throughout the lifetime of the cleanup (Vogel, 1998).

Other soil and groundwater treatments includes soil venting, in which clean gas are pumped into the contaminated subsurface. Afterwards, these gases become laden with chlorinated solvents and are then vented into the atmosphere. This method takes advantage of the high vapor pressure of chlorinated solvents to transfer contaminants from a liquid to a gaseous phase (Russell, 1992). But volatilization is another cause for concern because of the carcinogenic properties of gaseous chlorinated compounds (Lynge et al, 1997). Other treatments included soil excavation, surfactant flushing, and thermal treatments (EPA, 2004). However these methods still only convert chlorinated solvents



www.manaraa.com

into a different medium and require additional contaminant processing. A much more economic and efficient solution would be the complete destruction of chlorinated solvents.

## **Bioremediation**

Due to the need for cost effective treatment technology that could remediate contaminated sites to the standards set forth by EPA, bioremediation has become a viable alterative to conventional treatment methods. This method encourages the growth of indigenous or introduced microorganisms in the subsurface to metabolically facilitate the complete degradation of potential contaminants. Biodegradation of chlorinated solvents was first seen in 1980 at a Palo Alto groundwater recharge project (Bouwer and McCarty, 1983); subsequent laboratory studies showed that chlorinated products could be degraded under both anaerobic and aerobic conditions (Freedman and Gossett, 1989; Wilson and Wilson, 1985, Lee et al, 1998).

Field studies have also shown that aquifers have a number of indigenous microbial populations with a carrying capacity of around 10<sup>6</sup>-10<sup>7</sup> microorganisms per gram of dry soil (Bouwer, 1992). ATP extraction also indicates that around 10% of the cells in the subsurface are metabolically active (Bouwer, 1992). These microorganisms are from three main groups: prokaryotes, eukaryotes, archeabacteria. Each group has its own particular niche within the subsurface but the two most important classes of microbes, for bioremediation, appears bacteria and archeabacteria (Chapelle, 1993).

Under anaerobic conditions, bacteria obtain energy by conducting a reductive dechlorination reaction where chlorinated solvents combine with hydrogen to produce a less chlorinated product. This reaction uses highly chlorinated chemicals as electron



www.manaraa.com

acceptors and introduces a hydrogen atom to replace a chlorine atom in the compound (see Figure 1.1). Figure 1.2 below shows how PCE can be sequentially degraded into TCE, DCE, VC, and ethylene through reductive dechlorination (Vogel and McCarthy, 1985; Freedman and Gossett, 1989).



Figure 1.1 PCE is degraded to TCE in a reductive dechlorination reaction (Schmit and Vos, 2004)





Figure 1.2. PCE reduction pathway (Wiedemeier et al., 1999).

However, the rate of reductive dechlorination decreases as the number of attached chlorine atom decrease (Bouwer, 1993). So PCE, being saturated with four chlorine atoms, is a good electron acceptor and is readily degraded. TCE, with three chlorine atoms is also a good candidate for reductive dechlorination (Bouwer, 1993). VC, with only one chlorine atom attached, becomes the rate limiting step and could lead to a possible accumulation of VC in the subsurface (Chapelle, 1993).



Bacteria can also aerobically degrade chlorinated solvents through a co-metabolic process. During this reaction, bacteria did not gain any energy from the degradation. Instead contaminants were fortuitously degraded as bacteria produced an enzyme used for methane or ammonia oxidation (Wilson and Wilson, 1985; Yang et al. 1999). Studies have found that these methanotrophic bacteria, in the presence of methane, could fortuitously degrade chlorinated compounds such as TCE, DCE and VC into non-chlorinated products (Eguchi et al, 2001, Little et al, 1988). However this reaction was best for degrading less chlorinated compounds such as VC, DCE and TCE. Fully chlorinated PCE was not shown to degrade at all under co-metabolic conditions (Fogel et al., 1986). The rate of co-metabolic chlorinated solvent degradation was fastest for VC and slower for DCE and TCE (Bouwer, 1992).

Bioremediation has great potential for contaminate cleanup. However, anaerobic degradation is efficient at degrading highly chlorinated compounds, while aerobic degradation is efficient at degrading less chlorinated compounds. A better clean up solution would be to combine both degradation processes to take advantage of their respective benefits.

#### **Phytoremediation**

Phytoremediation in wetlands could be one solution allowing combined aerobic and anaerobic contaminate clean up (Lorah and Olsen, 1999, Lee et al, 1998). This process uses plants in the treatment of contaminated soils and could be a viable *in-situ* treatment to promote sequential anaerobic and aerobic microbial degradation (Cunningham, 1997). In the subsurface, plant roots interact with surrounding soil and forms a zone called the rhizosphere. Roots extrude photosynthetic products such as



amino acids, sugar, and vitamins into the rhizosphere (Walton, 1994). These products may serve as substrates to stimulate microbial facilitated degradation (Schnoor, 1995). In the process it has also been found that wetland plants possess a specialized gas transport tissue, called aerenchyma, used during oxygen transport to roots (Armstrong et al, 2000). However, some oxygen escapes and radially diffuses from the root into the surrounding rhizosphere, creating a zone of aeration millimeters wide surrounding the plant root (Armstrong et al, 2000; Bankston, 2002). The result is that an aerobic gradient is created within the rhizosphere and elsewhere the soil remains anaerobic. suggesting the potential for sequential anaerobic then aerobic chlorinated solvent degradation. Highly chlorinated contaminates, such as PCE & TCE, could be degraded in the anaerobic regions through reductive dechlorination (Vogel and McCarthy, 1985). Less chlorinated products of this reaction, such as DCE & VC, could diffuse into the aerated zones along the rhizosphere to be further degraded during co-metabolic reactions.

Some other mechanisms of phytoremediation include phytoextraction, phytodegradation, phytostabilization and phytovolatilization (Lunney, 2004). Phytoextraction is the process in which plants extract pollutants and degraded them by phytodegradation or incorporate them into plant biomass thru phytostabilization. Finally pollutants could be volatilized into the atmosphere through phytovolatilization (Lunney, 2004).

Plants could be used in many on-site remediation efforts of shallow aquifers and contaminated soil. They have been suited for degrading toxic organic contaminates such as benzene and chlorinated solvents or for accumulating inorganic heavy metals in plant



roots and shoots (Schnoor et al, 1995). Figure 1.3 below summarizes the phytoremediation mechanisms at work.



Figure 1.3. Phytoremediation mechanism (Schnoor et al, 1995)



#### Wetlands

Lorah and Olson in 1995 conducted a microbial study of a natural groundwater fed wetland at Lauderick Creek near the TCE contaminated sites of the Aberdeen Proving Ground (Lorah and Olson, 1999). TCE and tetrachloroethane (PCA) were shown to be degraded with subsequent increase of degradation products such as DCE and VC (Lorah and Olson, 1999). However these daughter products were soon degraded as well and not detectable at the surface (Lorah and Olson, 1999). Later, in 2003, Lorah and Voytec showed that dehalorespiration and methanogenic microbial populations were present in subsurface of these wetlands (Lorah and Voytek, 2003).

The observations by Lorah and Olson showed that it might be feasible to use wetlands for phytoremediation. In August 2002, constructed wetlands were built at Wright-Patterson AFB to conduct further studies. The constructed wetlands were modeled after natural upward flowing fens (Amon et al, 2002). Fens are a type of wetland fed by upward flowing groundwater source. The groundwater is recharged nearby at an area of higher elevation and it is discharged into fens where the groundwater flow reaches the surface (Amon et al. 2002). During phytoremediation contact with root surface is essential (Van der Lelie, 2001) and a fen wetland provides contaminated groundwater with significant root surface contact area and contact time (Amon, personal comm. 2005).

The constructed wetlands cells were built over an aquifer contaminated with PCE. Water was pump from the same aquifer and fed into the wetland. The cells were approximately 120 feet long, 60 feet wide, with an impermeable liner at 6 feet depth (Clemmer, 2003). Contaminated groundwater are pumped from an underground aquifer



www.manaraa.com

and enters the wetland thru three-inch diameter perforated PVC pipes, which are enclosed in a nine inch gravel layer, at the bottom of the constructed wetland (see Figure 1.4) (Clemmer, 2003). This configuration allows groundwater to enter the wetland in an upward flow schematic similar to fens. Above the gravel layer is 18 inches of 10% wood chip and hydric soil mix (Clemmer, 2003). The wood chips provide an organic carbon energy source for the microbes (Chapelle, 1993) and the hydric soils are very similar to local wetland soils with silt and clay inclusions (Amon et al, 2006 unpublished). Finally, 36 inches of wetland hydric soil are placed on top (see figure 1.5). Sampling tubes are set at depths of 9, 27, and 44 inches below the surface. The exit weir is located across from the inlet pipe and could be used to control water levels in the wetland (Clemmer, 2003). A study by Soboleweski (2004) has showed a 90% PCE degradation in the constructed wetland and subsequent increase in daughter products such as TCE, DCE and vinyl chloride.



Figure 1.4. Constructed wetland with three perforated PVC pipes at the bottom (Clemmer, 2003)





Figure 1.5. Cross-section of wetland soil layers (Soboleweski, 2003)

Many species of local wetland plants were planted in the constructed wetland. The plants were laid down in a grid pattern to determine which species might be most suitable for the treatment of PCE (Amon, person comm, 2005). Some plants such as *Eleocharis erythropoda* have the potential of being a good choice for remediation due to its large and deep root mass. Other plants such as *Carex comosa* and *Scirpus atrovirens* have broad leaves that are capable of volatilizing contaminates. However no studies have been done to compare treatment efficiencies among different plants.

## **Problem Statement**

Studies have shown the effectiveness of phytoremediation at degrading chlorinated solvents (Bankston, 2002; Schnoor, 1995; Lelie, 2001). However, plant species have a variety physical property such as: exudates quality, exudates concentration, plant metabolism, root adsorption surface area, root morphology, leaf size,



temperature tolerance and toxic compound tolerance (Shann and Boyle, 1994). When studies use a consortium of plants, difficulties often arise in characterizing the remediation efficiency of individual plants. For example at the constructed wetland at WPAFB, numerous plant species were planted in separate plots but they soon became mixed due to natural reseeding processes (Amon, personal comm. 2005).

### **Research Objective and Hypothesis**

Different plants species need to be studied individually to compare the remediation efficiency of each species. This research studied three different wetland plants species and an unplanted control, under a laboratory setting. Each plant had a different characteristic favorable for chlorinated solvent degradation. *Eleocharis erythropoda* (Spike Rush) has thin tube like leaves and large root mass. *Carex comosa* (Bearded Sedge) has broad leaves and *Scirpus atrovirens* (Green Bulrush) is broad leafed wetland plant with a long flowering stem during reproduction. PCE will be injected into the plant mesocosm and any possible PCE degradation will be observed. The hypothesis is that PCE will be degraded into daughter products in all the mesocosms; however, the question will be which plant is the most efficient at chlorinated solvent degradation and is there difference between the planted reactors and the control reactors?

## **Research Question**

- 1. Are there significant differences in the chlorinated solvent removal among the plant species and the unplanted control treatment?
- 2. Is there a difference in sulfate, nitrate and methane concentration among the mesocosm?
- 3. Do the sulfate, nitrate, and methane concentration influence PCE degradation?



#### **II. Literature Review**

## **Natural Attenuation**

Natural attenuation is the observed reduction in contaminate concentration as it migrates in the subsurface. The concentration reductions are due to the processes of dilution, dispersion, sorption, volatilization, and biotic transformation (Wiedemeier et al., 1999). Degradation mechanisms such as sorption, volatilization, and dispersion are nondestructive and only decrease the contaminant concentrations but not the contaminant mass. However, biodegradation degrades contaminants into harmless components and is the most important attenuation mechanism in the reduction of contaminant mass and concentration (Wiedemeier et al., 1999).

## Thermodynamics

To better understand biodegradation, it would be helpful to first understand the thermodynamic principals behind subsurface biological reactions. All biological reactions are constrained by the laws of thermodynamics. One of the biological reactions used during subsurface contaminate degradation is biologically mediated electron transfer during oxidation-reduction reactions (Wiedemeier et al., 1999). This reaction results in the oxidation of an electron donor and reduction of an electron acceptor to release energy. The energy produced can be quantified in terms of Gibbs free energy ( $\Delta G_r^o$ ), which is the maximum useful energy change for a chemical reaction at standard state (Zumdahl, 1997). The amount of energy available is dependent on the electron acceptor and donors used during the reaction. Table 2.1 and 2.2 below provides some examples of Gibbs free energy (Wiedemeier et al., 1999), calculated for electron acceptor half reactions and electron donor half reactions.



www.manaraa.com

## Table 2.1 Delta G of electron ACCEPTOR half cell reactions (Wiedemeier et al., 1999)

Half-Cell Reaction	$\Delta G_r^{\circ}$ (kcal/mol e <sup>-</sup> )
$4e^{-} + 4H^{+} + O_2 \implies 2H_2O$ Aerobic respiration	-18.5
$5e^{-} + 6H^{+} + NO_{3}^{-} \Rightarrow 0.5N_{2} + 3H_{2}O$ Denitrification	-16.9
$e^- + \underline{Fe^{3+}} \implies Fe^{2+}$ Fe(III) reduction	-17.8
$\begin{array}{l} 8e^- + 9.5 \mathrm{H}^+ + \mathrm{SO_4^{2-}} \implies 0.5 \mathrm{HS^-} + 0.5 \mathrm{H_2S} + 4 \mathrm{H_2O} \\ \mathrm{Sulfate\ reduction} \end{array}$	5.3
$8e^- + 8H^+ + CO_{2,g} \Rightarrow CH_{4,g} + 2H_2O$ Methanogenesis	5.9
$C_2Cl_{4,g} + H^+ + 2e^- \Rightarrow C_2HCl_3 + Cl^-$ PCE reductive dechlorination	-9.9
$C_{2}HCl_{3} + H^{+} + 2e^{-} \Rightarrow C_{2}H_{2}Cl_{2} + Cl^{-}$ TCE reductive dechlorination	-9.6
$C_2H_2Cl_2 + H^+ + 2e^- \Rightarrow C_2H_3Cl + Cl^-$ cis-DCE reductive dechlorination	-7.2
$C_2H_3Cl + H^+ + 2e^- \Rightarrow C_2H_4 + Cl^-$ VC reductive dechlorination	-8.8

## Table 2.2 Delta G of electron DONOR half cell reaction (Wiedemeier et al., 1999).

Half-Cell Reaction	$\Delta G_r^{\circ}$ (kcal/mol e <sup>-</sup> )
$1/2H_2 \Rightarrow H^+ + e^-$ Hydrogen oxidation	-9.9
$^{1/4}CH_2O + ^{1/4}H_2O \implies ^{1/4}CO_2 + H^+ + e^-$ Carbohydrate oxidation	-10.0
$4H_2O + C_2H_2Cl_2 \implies 2CO_2 + 10H^+ + 8e^- + 2C1^-$ DCE oxidation	-16.1
$4H_2O + C_2H_3CI \implies 2CO_2 + 11H^+ + 10e^- + C1^-$ Vinyl chloride oxidation	-11.4

A negative Gibbs free energy means that the reaction is exothermic and will proceed from left to right as shown on the tables above. Notice how aerobic respiration has the greatest  $\Delta G_r^o$  allowing bacteria specializing in aerobic respiration to out compete most other bacterial species if supplies are readily available (Bouwer, 1992).

If oxygen is limited, such as in underground aquifers, other electron acceptors such as nitrate, iron (III), sulfate, Mn(IV) PCE, TCE, and CO<sub>2</sub> can be used (Mitsch,



1993). However, none of these electron acceptors are as efficient as oxygen and in turn have a smaller  $\Delta G_r^o$ .

From the definition above, the  $\Delta G_r^o$  in Table 2.1 and 2.2 represents the maximum energy gained under ideal conditions and with free electron donors readily available. However, in the subsurface environment, electron donors are not readily available and microorganisms need to couple both oxidation and reduction half reactions to carry out the whole oxidation-reduction reaction. When two half reactions are combined the  $\Delta G_r^o$  of the two reactions are summed. The reaction will occur in nature only if the sum of the  $\Delta G_r^o$  results in a net energy gain (Wiedemeier et al., 1999).

One method to predict whether a reaction will occur is to examine the oxidationreduction potential or redox potential of the groundwater and then compare it to the redox potential of a chemical reaction. Redox potentials or  $E_h^o$  is measured in mV and ranges from -400 to 800mV. A high redox potential represents oxidizing zones containing high concentrations of electron acceptors, whereas a low redox potential represents reducing zones with a low concentration of electron acceptors. Most subsurface chemical reactions have a preferential redox range for reactions to take place (Mitsch, 1993). For example, Figure 2.4 below shows that oxygen and nitrate reduction occurs in high redox potential zone with an  $E_h^o$  at +820mV. Using oxygen and nitrates as electron acceptors generates a high energy yield and those microorganisms are able to compete for resources in oxidizing environments with high concentrations of the electron acceptors. On the other hand, optimum reductive dechlorination occurs in sulfate and CO<sub>2</sub> redox zones with  $E_h^o$  around -220mV. Reductive dechlorination of PCE does not generate a high energy



www.manaraa.com

yield and highly reducing conditions are needed before reductive dechlorination could take place (Bouwer, 1992).



Figure 2.1 Redox potential of possible subsurface reactions (Wiedemeier et al., 1999).

Bacteria are adapted to utilizing different electron acceptors during subsurface redox reactions and the goal of the redox reactions is to produce energy needed for cell growth. Based on thermodynamic principals, microorganisms would preferentially use oxygen and nitrate, because using these electron acceptors allow for a greater Gibbs free energy gain when compared with PCE. This process of utilizing the redox reaction with the greatest energy gain eventually changes the underground chemical composition and Figure 2.5 below shows the succession of underground chemical composition over time (Mitsch, 1993). This change in geochemistry is facilitated by microbes and an analysis of electron donor and electron acceptor in the subsurface could provide an indicator of what type of microbial population is present (Bouwer, 1992; Chapelle, 1993). So in an anaerobic environment, optimized PCE and TCE dechlorination will occur only if



competing electron acceptors with high energy yields are depleted and the subsurface environment becomes highly reductive.



Figure 2.2 Underground chemical composition change with time (Mitsch, 1993)

# Bioremediation

However, chlorinated solvent bioremediation is complex and several distinct bacterial communities may be needed for successful chlorinated solvent mineralization. Therefore it would be helpful to understand the chemical reactions that facilitate biotic degradation. The two main reactions used during biotic chlorinated solvent degradation are co-metabolic reactions and reductive dechlorination (Chapelle, 1993).

Under aerobic conditions co-metabolic chlorinated solvent degradation is possible (Wilson and Wilson, 1985; Little et all, 1988, Eguchi, 2001). During co-metabolic reactions bacteria produce monoxygenase enzymes that degrades methane or ammonium in the presence of oxygen (Yang et al., 1999). This reaction is used to generate energy and gain the carbon needed for bacteria growth and survival. Studies by Wilson and Wilson, showed that monoxygenase was a non-specific enzyme and could fortuitously degrade chlorinated contaminants along with the primary substrate. Later research



revealed that methanotrophic bacteria, such as *Methylococcus capsulatus*, used monoxygenase to degrade TCE, DCE, and VC into carbon dioxide (Fogel, 1986).

However co-metabolic degradation is best for degrading low chlorinated compounds such as VC and DCE. While TCE was shown to have some degradation and fully chlorinated PCE was not shown to degrade at all (Bouwer, 1992). Another limitation is that co-metabolic degradation will only proceed in the presence of methane and oxygen (Fogel, 1986). Yetbiotic methane generation occurs in highly anaerobic conditions and very limited amounts of methane are present in an aerobic environment. Therefore, the conditions necessary to carry out co-metabolic reactions typically do not occur in a natural environment and need to be engineered (Wiedemeier et al., 1999).

Under anaerobic conditions, bacteria could obtain energy by conducting Reductive Dechlorination Reactions (RDR), where chlorinated solvents react with hydrogen to produce a less chlorinated product. This reaction uses highly chlorinated compounds as electron acceptors and introduces a hydrogen atom to replace a chlorine atom in the compound (Vogel and McCarthy, 1985; Freeman and Gossett, 1989).

Fermentation products (lactate, methanol) and man-made chemicals( toluene) have been used to stimulate RDR (Freedman and Gossett, 1989). However, those compounds appears to be indirect electron donors that will produce dissolved hydrogen through further fermentation (Wiedemeier et al., 1999). Dissolved hydrogen is a high energy electron donor consumed by a wide variety of subsurface bacterial species (Wiedemeier et al., 1999). It is used by methanogens, nitrated reducers, sulfate reducers, and iron reducers (Wiedemeier et al., 1999). It is also the main electron donor used in anaerobic reductive dechlorination reactions (Wiedemeier et al., 1999). Subsurface



bacteria may utilize a variety of electron acceptors but they all compete for the limited amount of electron donor in the subsurface.

In a wetland, oxygen is frequently in limited supply (Mitsch, 1993). Some plants could diffuse oxygen into the soil but the gradient of aeration may only be a few millimeters thick (Bankston, 2002; Armstrong, 2000). However, many bacteria are adapted to carry out reduction reactions under anaerobic conditions (Freeman and Gossett, 1989). Each reduction reaction has a redox potential, depending on the type of terminal electron acceptor available. Chlorinated compounds are fairly electronegative and could be a good electron acceptors during reduction reactions (Smidt and Vos, 2004). One study showed that methanogenic conditions produce a favorable environment for microbes to use PCE as electron acceptors (Vogel and McCarty, 1985). In this reaction a chlorine atom is replaced by a hydrogen atom and PCE is sequentially transformed into TCE, DCE, and VC (Vogel and McCarty, 1985). Reductive dechlorination progresses rapidly for highly chlorinated compounds but as the number of attached chlorine atom decreases the reaction slows down and the final dechlorination of vinyl chloride is the rate limiting step (Freedman and Gossett, 1989). Therefore vinyl chloride, a carcinogenic substance, (Lynge et al, 1997) has been found to accumulate during reductive dechlorination (Freedman and Gossett, 1989). But a later study showed that refined methanogenic bacterial cultures that were acclimated to the subsurface environment could fully dehalogenate PCE to ethylene if sufficient electron donors were available (Smidt and Vos, 2004).

Currently studies have been focused at identifying the bacteria that could fully metabolize chlorinated solvents and a group of halorespiring bacteria (HRB) has been



www.manaraa.com

found to readily metabolize chlorinated solvents into harmless byproducts. These bacteria rely strictly on halorespiration and use hydrogen as the electron donor (Smidt and Vos, 2004). One genus in this group, *Dehalococcoides*, is able to fully dehalogenate PCE into ethane in a four step process (Smidt and Vos, 2004).

Bioremediation offers great potential for chlorinated solvent clean up. However, highly chlorinated species like PCE degrade rapidly under anaerobic conditions and less chlorinated species like DCE or VC are more readily degraded in an aerobic environment (Lee et al, 1998). An more efficient degradation process is needed to combine both anaerobic and aerobic degradation.

#### Phytoremediation

Phytoremediation is a subset of bioremediation and it uses plants in the treatment of contaminated soils, sediments, and water. Phytoremediation could be a viable in-situ treatment option to promote sequential anaerobic and aerobic microbial degradation (Cunningham, 1997). Some wetland plants could transport oxygen into the rhizosphere (Bankston, 2002; Armstrong, 200) and provide an aerobic gradient along the rhizosphere for co-metabolic chlorinated solvents degradation to take place. The rhizosphere is a specialized region of root and microbe interaction, with organic substrate and oxygen to support a microorganism community up to 100 times more abundant than non-vegetated soils (Walton, 1994). However the dimension of this zone of interaction is dependent on plant species. Competition from aerobic bacteria quick deplete any oxygen diffused by the plants (Walton, 1994) and soil just few millimeters away from the rhizosphere could be anaerobic. This is advantageous in chlorinated solvent remediation because anaerobic microbes could reductively dechlorinate PCE under methanogenic conditions (Vogel and



McCarthy, 1985). However, the rates of dechlorination under anaerobic conditions are slow (Crowley, 1997) and only specialized *Dehalococcoides* grown in labs could fully degrade PCE to ethane (Smidt and Vos, 2004). PCE daughter products not degraded under reductive dehalogenation could possibly be transported to the rhizosphere. There aerobic microbes could proceed to degrade PCE daughter products such as TCE, DCE, and VC into ethylene, methane or carbon dioxide (Wilson and Wilson, 1985; Fogel, 1985).

Plants also release organic carbon such as amino acids, vitamins, and sugars into the root rhizosphere (Walton, 1994) to further stimulate microbial degradation (Schnoor, 1995). This sequential anaerobic and aerobic bioremediation involves microbial interaction within the subsurface. But EPA studies have shown that plant themselves could also directly extrude contaminate degrading enzymes such as dehalogenase, nitroreductase, peroxidase, lactase, and nitrilase (Schnoor, 1995). Plants could also directly uptake pollutants. But direct uptake is dependent on three factors: the physicochemical properties of the compound, environmental conditions, and plant characteristics (Cunningham, 1997). Chemical properties, such as water solubility and octanol-water partition coefficient, are important in the determining the availability of the pollutant. For example, contaminant uptake studies using poplar trees have showed that a moderately hydrophobic organic chemical with Log Kow = 0.5-3 could be taken up and translocated to above ground tissues (Schnoor, 1995). The environmental conditions are also important. For example soil with concentrated ferrous iron content could be oxidized into insoluble ferric iron from oxygen in the rhizosphere. Ferric iron then coats the root surface and prevents the uptake of pollutants (Mitsch, 1993). Finally plant


characteristics are important as well because root surface area could significantly affect adsorption rate. Roots with numerous root hairs would have a higher surface area to absorb pollutants (Cunningham, 1997). Once the chlorinated solvents are taken up into the plant biomass, they undergo hydroxylation and glycosilation to change them into more soluble forms (Cunningham, 1997). These pollutants are then sequestered into plant cytoplasm or cell wall matrix.

The amount of pollutant degradation can be highly dependent on the species of plants used for remediation (Shann and Boyle, 1994). Many different type of growth properties exist such as: exudates quality and amount, plant metabolism, root adsorption characteristic and morphology, leaf size, temperature tolerance and toxic compound tolerance. So choosing the right species fit to degrade a particular type of contaminate is very important in developing a phytoremediation treatment plan (Shann and boyle, 1994)



## **III. Methods**

## **Column Construction**

My study was designed to examine three different wetland plant species and an unplanted control mesocosm subjected to continuous injection of low concentrated PCE solutions. In order to study individual plant species, twelve PVC column reactors were built to replicate the constructed wetland. Each column had an upward flowing ground water scheme similar to both the WPAFB constructed wetland and a natural fen wetland. The columns were divided into three mesocosm and each mesocosm was planted with one wetland plant species. A fourth mesocosm was set aside and reserved as a control with no plants. Each reactor was then randomly placed along a glass wall within a greenhouse. See table 3.1 and 3.2 for a list of reactors and plant species. Figure 3.1 shows a picture of the actual column setup within the greenhouse.

Table 3.1 Plant species and number of reactors

Carex comosa	Scirpus atrovirens	Eleocharis erythropoda	Control
2 reactor	4 reactors	3 reactors	3 reactors

Reactor	Plant species	Reactor	Plant species
1	Carex comosa	7	Eleocharis erythropoda
2	Carex comosa	8	Control
3	Control	9	Scirpus atrovirens
4	Eleocharis erythropoda	10	Eleocharis erythropoda
5	Scirpus atrovirens	11	Control
6	Scirpus atrovirens	12	Scirpus atrovirens

Table 3.2 Plant and control column placement





Figure 3.1 Mesocosm placement within greenhouse

The reactors were built to approximate the dimensions found at the WPAFB constructed wetland and the plant species were chosen to simulate the WPAFB wetlands. Figure 3.2 - 3.4 shows a schematic of the column reactors, sampling port and sampling port placements. The reactors were constructed from 6" diameter PVC pipes 60" in height, with an internal volume of 0.9817 ft<sup>3</sup> or 27.7 L. Along the sides of the columns, 1/2" soil extraction and water sampling ports were drilled. An influent port was located on bottom of the reactor and 7 sampling ports were placed along the length of the reactors. Ports 1- 4 were located on bottom half of the reactors and spaced 6 inches apart. Ports 5 - 7 were located along the upper half of the reactor and spaced 9 inches apart. Each sampling port consisted of a 1/2" diameter PVC pipe 5.5 inches long. The sampling pipes were perforated with slits 3/4" apart to allow water to flow through the pipes. Inside the sampling pipe, a 3/4" diameter polyethylene tube 4 inches long was inserted



half way into the sampling pipe. This was connected to a Fisher 3-way valve to allow for sampling. Each sampling pipe was place 30 degrees apart along the circumference of the column. This placement was necessary to avoid channeling within the soil. Finally an outflow port was placed 6 inches above port 7 to allow the out flow of excess water.



Figure 3.4 Sampling port placement

Water was distributed as shown on Figure 3.5. Pure distilled water could leach ions out of the soil in the mesocosm. So water in the reservoir was filled every three days using 50% tap water and 50% distilled water. Water was pumped into the column reactors using 2 Cole-palmer masterflex peristaltic L/S pumps (Cole Palmer, Vernon



Hills, IL). Each pump was connected to six column reactors and the pump rate was maintained 2.0 mL/min throughout the experiment. An 8 channel, 3 roller, pumphead, with 6 standard small cartridges was attached to each pump and used to distribute the water to columns. A KDScientific model 100 syringe pump and a 300mL mixing chamber were placed between the pump and reservoir to allow for PCE injection into the columns. Water was drawn from the mixing chamber and pumped through a manifold in an effort to ensure equal flow into all of the columns. The water was then delivered to each column using a transparent Nalgen FEP Teflon tubing with 1/8" I.D. Figure 3.5 below shows the setup used during the experiment.



Figure 3.5 Experimental set up



## **Planting Timeline**

Each column reactor has 6 inches of pea gravel placed along the bottom. The soil substrate used in the columns consisted of hydric soils from taken from the Beavercreek wetlands. On May 20<sup>th</sup>, approximately 325 liters of soil was inoculated with one top to bottom core sample from the WPAFB's constructed wetland. The core sample was 7.5 cm in diameter and ran the entire depth of the constructed wetland. This was done to try to populate the greenhouse mesocosms with microbial species that was already present in the constructed wetland. The soil substrate was then mixed to homogenize the soil and filled into the columns. On May 24<sup>th</sup>, wetland plants were transplanted into the reactors and allowed approximately 3 month to overcome the shock of transplanting and to acclimate to the new habitats within the columns. The primary purpose of this development period was to promote the growth of the root systems shown to be important during plant assisted PCE degradation (Walton, 1994).

#### **Syringe Pump Injection Rate**

The injection rate and stock PCE concentration was experimentally calibrated on September 20, 2006 to achieve the desired PCE concentration of around 50 ppb for injection into the column reactors (see Table 3.3). The syringe pump holds a 50cc Hamilton Gastight Teflon plunger syringe (Hamilton, Reno, NV) and injects a 100 ppm stock PCE solution into the mixing chamber. In Table 3.3 below, two samples were taken from the influent port of column 2 and 10 and measured twice on the GC to determine their PCE concentration. Samples were taken when the syringe pump was injecting at rate of 1ml/hr, 1.2 mL/hr, and 1.4 mL/hr. In end a syringe pump injection



28

rate of 1.6 mL/Hr was used throughout the experiment to ensure that the PCE concentration pumped into the columns did not fall below 50 ppb.

Table 5.5 Initial I CL Injection canoration result									
1 ml/Hr syringe pump injection rate.			1.2 ml/Hr syringe pu	ump injection rate.		1.4 ml/Hr syringe p	ump injection rate.		
Port	PCE Conc (ppb)		Port	PCE Conc (ppb)		Port	PCE Conc (ppb)		
2 IFF	25.60436		2 IFF	31.57188		2 IFF	53.72962		
2 IFF	23.6021		2 IFF	31.54772		2 IFF	45.92896		
10 IFF	31.36048		10 IFF	41.53788		10 IFF	58.870868		
10 IFF	30.38502		10 IFF	38.40916		10 IFF	54.95272		

Table 3.3 Initial PCE injection calibration result

### **Inflow Rate**

A flow rate measurement was conducted on September 21, 2005 by measuring the bubble velocity within a tube. As stated above in the column construction section, each column was connected to the pump using a transparent Teflon tubing with an 1/8" ID. While the column was connected to the pump, a 30 cm section of the tubing was marked off and the time it took for a bubble to traverse that 30 cm was measured with a stop watch. The velocity of the bubble and the inner area of the tubing are then calculated. The bubble velocity is used to represent the velocity of the water flowing through the tubing.

Subsequently, the flow rate for each column could be calculated from the bubble velocity and the area of the tubing. The results for flow rate are shown in Table 3.3. This method takes into account the backpressure from the columns because the pump is connected to the column while the measurements are gathered. However, some possible errors from this method could be the friction and capillary forces between the bubble and the tubing inner surface. Another error could be the compression of the air bubble as it traverses through the tube.



	Table 3.3	Tubing	characteristic	and	Flow	rate
--	-----------	--------	----------------	-----	------	------

Tubing ID 1/8":	0.3175	cm
Tubing Area:	0.0792	cm^2
Distance Travelled:	30	cm

	Col 1	Col 2	Col 3	Col 4	Col 5	Col 6	Col 7	Col 8	Col 9	Col 10	Col 11	Col 12
Bubble Travel Time (sec)	62	70	78	62	72	73	66	71	62	74	71	66
Bubble Velocity (cm/sec)	0.48	0.43	0.38	0.48	0.42	0.41	0.45	0.42	0.48	0.41	0.42	0.45
Flow rate (CC/Min)	2.30	2.04	1.83	2.30	1.98	1.95	2.16	2.01	2.30	1.93	2.01	2.16

An effluent flow rate was also gathered from the effluent port by measuring the volume of liquid accumulated within a certain time span. The results are shown below on Table 3.4. Column 3 and 12 did not have any flow at the top of the column during the measurement time period and no effluent flow rate data was gathered.

Table 3.4 Effluent flow rate calculate over 36 hrs.

Column:	Col 1	Col 2	Col 3	Col 4	Col 5	Col 6	Col 7	Col 8	Col 9	Col 10	Col 11	Col 12
Flow rate(mL/min):	2.17	2.16		1.95	2.08	2.01	1.91	2.10	2.10	1.94	2.09	

# Water and Organic content:

The water and organic content of soil in each column were measured on August 31, 2005. A soil sample was taken out of port 7 and weighted. It was then placed into a 99 °C oven for 16 hours to dry out the sample. The dried soil was then weighted again to calculate the water content of soil. The same soil sample was then baked in a 500 °C furnace for 8 hours to burn off the organic content of the soil. See Table 3.2 for the water content and organic matter for each column reactor. The water content of the soils ranged from 42% to 48% and the organic matter for each column were consistent at around 10%.



	Water Content	Organic Content
Column	(Mw/Mwet)*100	(Mo/Mdry)*100
1 Carex Comosa	41.9%	10.2%
2 Carex Comosa	46.0%	10.8%
3 Control	43.7%	10.4%
4 Eleocharis erythropoda	45.4%	10.6%
5 Scirpus atrovirens	49.0%	10.5%
6 Scirpus atrovirens	47.2%	10.5%
7 Eleocharis erythropoda	44.9%	9.7%
8 Control	47.1%	10.3%
9 Scirpus atrovirens	48.5%	10.2%
10 Eleocharis erythropoda	46.5%	9.8%
11 Control	45.3%	10.0%
5 Scirpus atrovirens	48.2%	10.0%

Table 3.5 Water content and organic content of soil. Water content is calculated from mass of water and mass of wet soil. The organic content is a percentage of the mass organics and the mass of dry soil.

# **Standard Preparation**

Standards were prepared for PCE, TCE, cisDCE and transDCE during September and October. Standards preparation were calculated using the equation  $C_1V_1 = C_2V_2$ . The stock solutions for the standards were prepared by injecting 2 µL of pure PCE into a capped 160 mL serum bottle. The bottle was then spun for 48 hrs to 72 hrs. Table 3.6 below list the calculations used to find the concentration within the stock bottles.

 $C_1$ : is the initial concentration of stock  $V_1$ : is the initial volume of stock  $C_2$ : is the desired concentration of standards  $V_2$ : is the volume of the standards



PCE Stock Prepara	ation for standards:	TCE Stock Prep	aration for standards:
Volume of pure PCE:	2 µL	Volume of pure TCE:	2 µL
Density:	1.623 mg/µL 20oC	Density:	1.458 mg/µL 20oC
Stock bottle volume:	160 mL	Stock bottle volume:	160 mL
Mass of PCE:	3.246 mg	Mass of TCE:	2.916 mg
Concentration PCE:	20.288 mg/L(ppm)	Concentration TCE:	18.225 mg/L(ppm)
cisDCE Stock Prepa	aration for standards:	transDCE Stock Pr	eparation for standards:
Volume of pure cDCE:	2 µL	Volume of pure tDCE:	2 µL
Density:	1.282 mg/µL 20oC	Density:	1.26 mg/µL 20oC
Stock bottle volume:	160 mL	Stock bottle volume:	160 mL
Mass of DCE:	2.564 mg	Mass of DCE:	2.52 mg
Concentration DCE:	16.025 mg/L(ppm)	Concentration DCE:	15.750 mg/L(ppm)

Table 3.6: Stock solution Prep

Each standard bottle was created by injecting 5, 15, 30, 150, or 250  $\mu$ L of stock solution into a 15 mL serum bottle filled with DI water. The new concentration within each standard bottle was then calculated using the equation  $C_1V_1 = C_2V_2$  and the bottles were spun for 1 hr to allow for the stock solution to completely diffuse. After 1 hr, the spinning process stopped and 5 mL was withdrawn and injected into an empty Nitrogen gas filled serum bottle. This created a standard bottle with 5mL of liquid and 10mL of headspace. Spin the newly injected standard bottle overnight for 12 hr to allow for equilibration. Afterwards initiate head space analysis of the standards on GC. This standard preparation process best approximated the sampling methods used in this experiment and allows for a more accurate calibration.

VC, methane and ethylene gas standards are prepared using gaseous standard preparation procedure. First, purge a 72 mL serum bottle with the desired gas for 30 minutes. Then use a Hamilton gastight syringe to withdraw 2, 5, 20, 50, or 200  $\mu$ L of gas and inject it into a serum bottle filled with 5 mL of DI water. In the end the serum bottle will have 10mL of headspace and 5mL of liquid volume. The bottle is then spun over



www.manaraa.com

night for 12 hrs to allow for equilibration. The concentration in the gas and water is calculated using ideal gas and mass fraction equation. After equilibration initiate head space analysis of the standards on the GC.

#### **Sampling Schedule**

The first set of water samples was collected on Aug 15, 2004 to establish a baseline condition within the column reactors. Samples were collected from ports 7, 5, and 1 for columns one through two and port 7, 6, 4, and 1 for columns three through six. The six columns sampled contained a representative sample of the different wetland plants under observation.

Starting September 18, 2005, PCE injections began. The goal was to inject 50 ppb PCE solution continuously into the column reactors over the experimental period. This injection of chlorinated solvents was designed to simulate an upward flowing contaminant plume traveling past the roots system of a wetland plant. After PCE injections began, the plants were given approximately 2 weeks to acclimate to the presence of PCE and to allow for possible development microbial colonies capable of degrading PCE.

The second of water samples from the reactors were collected on October 1, 2005. The third sample was conducted 4 weeks after PCE injections began on Oct 16, 2005. A final sample set was conducted on Nov. 3, 2005 or 6 weeks after PCE injections began. Each sampling set collected samples from seven sampling ports and one influent port per reactor. All together the twelve reactors combine to give 106 samples per sampling set. (see appendix A for sampling method)



33

## Gas Chromatograph

A Hewlett-Packard model 6890 was the only GC used during the experiment. The GC was setup with a split inlet into two columns. The column used for the FID (Flame ionization detector) was a JW Scientific Inc, Cat# 1134332, 30m x 0.320 mm. The column for ECD (Electron capture detector) was a Hewlett-Packard HP-624, 30m x 0.320mm x 1.8 µm film thickness. Table 3.7 below shows the detection limit on the GC. PCE, TCE, 1,2 cDCE, and 1,2 tDCE was analyzed on ECD, while methane was analyzed on the FID. Test on ethylene was conducted to determine the residence time, however no detection limit analysis was conducted. VC, 1,1DCE, and ethene was not analyzed in this experiment. (See appendix B for GC methods)

	1		
		Detection Limit (ECD)	Detection Limit (FID)
	R-time	Concentration (ppb)	Concentration (ppb)
PCE	7.565	0.23	none
TCE	5.485	0.56	none
1,2 cDCE	2.586	0.45	none
1,2 tDCE	3.175	2.188	none
Methane	1.533	none	0.2
Ethylene	2.422	N/A	N/A

Table 3.7 Detection limits on the GC

# Samples for IC

Nitrate, sulfate, nitrite and residence time samples were gathered for IC analysis. The ion samples were gathered from all 106 ports on twelve columns. Sampling methodology was similar to the chlorinated samples methodology stated in Appendix A.

The ions of interest during sampling were sulfate and nitrate. The samples were collected by first opening the port and allowing it to flush with 10 mL of liquid. The samples was then gathered and stored in a screw capped 10mL vial from cole-palmer.



Each sample vial was then spun for 10 min on a 10,000 rpm centrifuge to separate out the particular matter. The liquid samples were then transferred to a 10 mL Dionex autosampler vial.

### **Residence Time Sampling**

A breakthrough test using Potassium Bromide (KBr) had been conducted to determine the retention time within the columns. As water was injected into the bottom of the column, it traveled up the column and out through the outlet port. The amount of time needed for the water to progress through the entire column was the retention time (RT). The retention time was useful in determining how long chlorinated solvents are in contact with the soil and root matrix in the column. The longer the contact time the more opportunity there was for degradation or uptake into plant tissues. The breakthrough test was conducted by injecting 60 mL of 6g/L KBr solution into the inlet port. Two analysis procedures had been used to measure the breakthrough curve. The first procedure used a conductivity meter to continuously monitor any increase in conductivity at top of column 9. The resulting increase in conductivity should form a breakthrough curve allowing for the calculations of retention time. The second procedure used a timeseries method to take samples from column 9 port 4 every 30 minutes. Samples were gathered over a period of 35 hours. The KBr samples were then analyzed on an IC to determine the bromide concentrations of each sample and then plotted to form a breakthrough curve.

#### Ion Chromatograph

A Dionex Ion Chromatograph equipped with AS-50 autosampler was used for the analysis. Brad Short Improved Control Program was the methods used to analyze each



35

sample for sulfate and nitrate. Bromide Tracer JY Control Program was the IC method used to measure bromide concentration (see Appendix C for IC methods).

# **General Supplies**

All PCE, TCE, cDCE, and tDCE chemicals were reagent grade and purchased from Fisher Scientific. The chemicals have been in storage for at least 1.5 yrs at room temperature in a fire proof case. All GC gases used were ordered from WSU lab supply shop. Nitrogen gas was zero grade nitrogen (99.9%) and methane was ultra high purity (99.95%). 10% w/v KBr stock solution was acquired from Fisher Scientific.



#### **IV. Results**

### **Pre PCE Injection Baseline Results**

In order to assess the PCE degradation potential, a baseline pollutant concentration level was established. This baseline would allow for the meaningful comparisons of results and possible assessment of reactor conditions before and after PCE injection. PCE was not injected during the first sample set and consequently PCE, TCE and other chlorinated degradation products were not detected within the reactors. Figure 4.1 plots out the methane concentration pre-PCE injection. The control (unplanted) column had the highest methane concentration and the maximum was reached at 45". The planted columns had less peak methane concentration than the control columns and tended to peak at 27" or 36". Feed water from the reservoir contained no methane and the methane concentration than the control reactors, indicating that the plants within the columns were influencing the ground water chemistry.





Figure 4.1 Background methane concentration pre-PCE injections (µg /L)



## **Time Series Data Sets**

During first and second sampling, the injection methods were still under development and resulted in varying PCE concentrations at the influent ports. For all time series data, the Y-axis represents column height along the column, from 0" at the influent port to 54" at port 7 and the X-axis was the contaminant concentration in ppb.

The first sampling set was conducted 2 weeks after continuous PCE injections began and Figure 4.2 below shows the average PCE concentration. The largest PCE concentration decrease for all treatments occurred along the bottom of the columns, between 0" and 15". At the top half of the columns, PCE concentration decreased slowed for all treatments. The significant PCE concentration decrease occurred at the bottom of the reactors indicating that some type PCE removal process was taking place just 2 weeks after injections began.



Average PCE comparison 2 weeks after PCE injection





Figure 4.3 below, illustrates the average TCE concentration within the columns. TCE was not detected during background sampling and presence of TCE within the columns proves that PCE dechlorination was taking place. The primary trend for this data set shows that TCE concentration increased towards the middle of the column, with EE columns having the most TCE, followed by the control column. It is also of note that between 45" and 54" the TCE concentrations decreased for *Carex comosa* and *Eleocharis erythropoda*, while it increased for the control columns and for *Scirpus atrovirens*. The highest TCE concentration in this sampling set was only 4 ppb. So 15 days after continuous PCE injections began the column reactors may not have fully developed its chlorinated solvent degradation potential.



Average TCE comparison 2 weeks after PCE injection

Figure 4.3 Average TCE concentration for the first sample set



The third chemical under observation was methane, which was detected in both the background samples and post-PCE injection samples. Figure 4.4 below shows that the planted reactors had an increase methane concentration towards the middle of the columns and the concentration decreased towards the top. This is consistent with pre-PCE methane results shown on Figure 4.1. The only exception was the control reactors, in which the average concentration increased all the way to port 6 and then remained steady to the very top of the columns.



Figure 4.4 Average methane concentration for the first sample set



Further comparisons between methane concentration pre and post PCE injections are shown in Figure 4.5. The methane concentration for both data sets maintained similar trend of increasing towards the middle but decreasing towards the top of the columns. However, the methane concentrations pre-PCE injections were significantly greater than post-PCE injections. This shows that methanogenic activities were reduced after PCE has been introduced.



Figure 4.5 Methane comparison in (mmol/L) pre and post PCE injections



Sampling Set #2:

The second sampling set was conducted 4 weeks after continuous PCE injections began. Figure 4.6 presents the data for 2 weeks and 4 weeks data and allows for comparison between the two sampling sets. The PCE concentration for the 4<sup>th</sup> week sampling set was greater than the 2<sup>nd</sup> week samples. The trend for PCE decrease during the 4<sup>th</sup> week was much more linear and more PCE penetrated at 54" near the top of the column.



Average PCE comparison 2,4 weeks after PCE injection

Figure 4.6 PCE comparisons 2 and 4 weeks after continuous PCE injections began



In Figure 4.7 below, comparisons of TCE concentrations were made at the 2 week and 4 week point. As shown on the graph, the average TCE concentration had increased for all treatments after 4 weeks. However, the TCE concentration for all treatments still remained less than the PCE concentration. The second sampling set also showed that the TCE trend for all treatments increased to a maximum at 36" and then decreased above it. However, at 54" TCE had not been fully removed and all treatments had detectable levels remaining. At 54" The control columns had the greatest TCE concentration followed by *Eleocharis erythropoda*.



Figure 4.7 TCE comparisons 2 and 4 weeks after continuous PCE injections began



Sampling Set #3:

The third sampling set was conducted 6 weeks after continuous PCE injections began. In Figure 4.8 -4.10 the PCE results for all three data sets were compared, comparisons of the average PCE concentrations were made at the 2, 4 and 6th week point. The trend for the 6<sup>th</sup> week sampling set was similar to 2<sup>nd</sup> week samples and the majority of PCE concentration decrease occurred below 15". For all treatments, the 6<sup>th</sup> week samples had more PCE than the 2<sup>nd</sup> week samples but it had less PCE than the 4<sup>th</sup> week samples. Showing that PCE removal at 6<sup>th</sup> week was more efficient than at the 4<sup>th</sup> week.



#### Average PCE comparison 2 weeks after PCE injection



Figure 4.8 PCE data 2 weeks after injections began

Average PCE comparison 4 weeks after PCE injection



Figure 4.9 PCE data 4 weeks after injections began

Average PCE comparison 6 weeks after PCE injection



Figure 4.10 PCE data 6 weeks after injections began



Figure 4.11 - 4.13 shows that the TCE concentrations at 6<sup>th</sup> week had increased. The planted columns followed the same trend of increasing in the middle of the columns and decreasing near the top. The control columns on the other hand continued have increasing concentration through out the entire length of column. At 6h week the increase in TCE concentration below 15" correlated well to the decrease in PCE concentration PCE data shown in Figure 4.10 above.









Average TCE comparison 2, 4, 6 weeks after PCE injection

Figure 4.13 TCE data 6 weeks after injections began



Figure 4.14-4.17 compares the methane concentrations among the *Carex comosa*, *Scirpus atrovirens, Eleocharis erythropoda*, and control column respectively. In Figure 4.14, the maximum methane concentration was achieved at 27"; afterwards the concentration decreased to near zero at the top of the column. In addition, a lower methane concentration was observed in the 6<sup>th</sup> week sample set. In Figure 4.15, the *Scirpus atrovirens* maximum methane concentration was achieved at 21". It also showed a significantly lower methane concentration in the 6<sup>th</sup> week sampling set. In Figure 4.16, *Eleocharis erythropoda* maximum methane concentration was achieved at 36". Finally in Figure 4.17, the control column's methane concentration increased throughout the length of the column and the maximum concentration was found near the top of column at 54". The methane concentration for the control samples stopped increasing as it approached the top of the column, suggesting something is inhibiting methane production near the top. In the all sample sets no *cis*-DCE or *trans*-DCE or any other PCE degradation products were detected.



Figure 4.14 Carex comosa methane comparison .









Figure 4.16 Eleocharis erythropoda methane comparison







# **PCE Concentration:**

The following charts are used to compare the PCE removal effectiveness of different plants. Figure 4.18 shows PCE concentration decrease between each port at week 2. Each bar represents the amount of PCE decrease between two ports for a particular treatment. For example, the average concentration decrease for *Carex comosa* between the influent port and port 1 is about 32 ppb. The error bars on the graph represents the standard deviation of the data. Unfortunately, all data sets had a significant standard deviation due to the limited sample size of each species. Figure 4.18 shows that the greatest PCE concentration decrease occurred below 15".



Figure 4.18 PCE Concentration decrease between ports after 2 weeks of continuous PCE injection. Student t-analysis with an alpha = 0.1, showed that between 0" and 9" *Carex comosa's* concentration decrease was significantly greater than the control reactors. All other ports showed no significant difference between treatments.



51

Figure 4.19 shows the cumulative percentage of PCE removed at each port. For example at 9", the control column had about 38% PCE concentration decrease. At 15", *Carex comosa* had the greatest percentage of PCE concentration decrease, with 90% drop in PCE concentration. Near the top of the column at 54", *Carex comosa* and *Eleocharis erythropoda* had the greatest removal percentage with about 99% of PCE removed at port 7 or 54"



Figure 4.19 Cumulative percentage of PCE removed at 2 weeks



Figure 4.20 below shows the PCE removal between ports after six weeks of continuous PCE injections. Carex comosa's most active zone of PCE decrease was still between 0 and 9 inches. Between 9" and 15" all treatments had less PCE removals but Carex comosa had the greatest drop in PCE removal.



6th Week PCE concentration Decrease Between Ports

Figure 4.20 PCE Concentration decrease between ports at 6 weeks. Student tanalysis performed on the data, with an alpha = 0.1, showed that between the influent and port 1, the Carex comosa decrease was significantly greater than all of the other treatments. All other ports showed no significant difference between treatments.



Figure 4.21 below shows the cumulative percent of PCE decrease after 6 weeks. *Carex comosa* had the greatest decrease in PCE concentration with about 96% decrease at port 7. However, the other treatments also showed significant percentage of PCE removal, with every treatment removing greater than 90% of the initial PCE injection.



Figure 4.21 Percentage of PCE removed at 6 weeks

# **TCE Data**

After 2 weeks of continuous PCE injections, Figure 4.22 below compares the TCE concentration among the treatments. There were large standard deviations among each of the treatments but comparisons of the average concentrations showed that below 21", *Eleocharis erythropoda* had the highest TCE concentration. At 27" the control reactors had the greatest concentration of TCE and above 27" TCE concentration because



to decrease among all treatments. Finally at 54", the control reactors had the greatest concentration among all the treatments.



Figure 4.22 TCE concentration at 2 weeks

After 6 weeks of continuous PCE injections, Figure 4.23 below compares the TCE concentration among the treatments. TCE were detected at 9" and *Carex comosa* had the highest concentration. At 15" all treatments had higher concentration of TCE, with the control columns having the highest TCE concentration. Between 15" and 36" all planted columns had increasing concentration of TCE. Above 36" the TCE concentrations within the planted columns decreased. However the control column's TCE concentration maintained about the same level until 45" whereby it increased to have the highest concentration of TCE among all the treatments.



55



#### TCE concentration at 6 weeks



# **TCE and PCE Comparison**

Since, TCE was not detected during the initial background sampling of the reactors, any TCE detected after PCE injections began was due to the degradation of PCE. Under ideal conditions, the molar concentration of TCE detected should equal the molar concentration of PCE decrease between each port. After 6 weeks of continuous PCE injections, comparisons were made between PCE decrease and the TCE increase (see Figures 4.24 - 4.27). All concentrations were in units of mmol/L to allow for comparison between different species.

The results from all four treatments showed that initially there was a large decrease in PCE concentration between the influent 0" and 9" and the increase in TCE concentration did not match the PCE decrease. Between 9" and 15", *Scirpus atrovirens* 



and control reactors had increased TCE removal. Between, 15" and 45" the PCE concentration decreased at a slower rate but the TCE concentration increase still did not match PCE concentration decrease.







Figure 4.25 Scirpus atrovirens PCE-TCE comparison





Figure 4.26 *Eleocharis erythropoda* PCE-TCE comparison



Control PCE and TCE concentration change after 6 weeks




## **Sulfate and Nitrate Comparison**

Sulfate, nitrite and nitrate are natural occurring chemicals within the subsurface. Sulfates  $(SO_4^{-2})$  are formed from the oxidation of hydrogen sulfide. Nitrite  $(NO_2^{-})$  and nitrate  $(NO_3^{-})$  are formed from the aerobic oxidation of ammonium ions  $NH_4^+$  (Mitsch, 1993). Like PCE, both nitrate, nitrite and sulfate are electron acceptors and, in an anaerobic environment, microbes could combine these chemicals with an electron donor to produce energy (Wiedemeier et al., 1999). Microbes preferentially use the chemical that produces the most energy and Figure 2.5, shows the succession of the chemicals within the subsurface (Mitsch, 1993). So the presence of these chemicals may influence optimum PCE dechlorination within the subsurface.

Figure 4.28 and 4.29 below presents the average nitrate and sulfate concentration found within the columns. No nitrite was detected among any of the columns.



Figure 4.28 Average nitrate concentrations with depth



The nitrate concentration for all four treatments grew at a very slow rate to 36". Above 36", planted treatments had a large increase in the nitrate concentration. However, the control treatment had little change in nitrate concentration and it maintained approximately the same nitrate concentration above 36". At 45", a student-t analysis with alpha=0.1 found the nitrate concentration for *Scirpus atrovirens* to be statistically greater than both *Eleocharis erythropoda* and control columns (p-value = .003). At 54" there was no statistically difference between all treatments.



Figure 4.29 shows the average sulfate concentration found within the columns. A student-t analysis with alpha=0.1 was also performed for the sulfate concentration and it showed that *Eleocharis erythropoda* had statistically greater sulfate concentration than all other treatment at 0" and 9". Between 15" and 27" all treatments had no statistical difference in sulfate concentrations and they are slowly decreasing. At the top of the columns between 45" and 54" the sulfate concentrations for *Carex comosa, Scirpus atrovirens,* and control reactor started to increase. Comparisons showed that the control reactors had statistically greater sulfate concentration than Eleocharis erythropoda at 54" (p-value = 0.07). No significant difference were detected among the other treatments.



Figure 4.29 Average Sulfate concentrations for all four treatments



## Conductivity

Soil conductivity is a indirect measure of the ion content within the soil. Increase conductivity could be correlated to an increase in ion concentration. However, soil conductivity does not differentiate between what species or mixtures of ions are in the soil. It could also be used to measure bromide concentration during retention time analysis. Finally conductivity could be used as an indirect measure of the plant's evapotranspiration rate. If conductivity rises as water travels through the column, it could indicate that plants are actively transpiring water causing an increase in salt concentration. Such increase in salt concentration could be harmful to the plants and actions should be taken to flush out the excess salts.

The soil conductivity within the columns has been measured to give a better understanding of the ion activity within the soil Average conductivity results are shown below in Figure 4.30. All inflow was from the same reservoir tank, so at 0" there were no differences in conductivity results. Between 9" and 36" the soil had little impact on the conductivity and no significant difference was found between treatments. However, above port 36" the conductivity reading started to increase for all of the planted treatments. This suggest that evapotranspiration is influencing the salt concentration within the top of the columns. *Scirpus atrovirens* had the greatest conductivity increase followed by *Carex comosa*, both plants have broad leafs useful during evapotranspiration. Only the control treatment did not show a significant increase in conductivity. If evapotranspiration is occurring it suggests that phytovolatilization of chlorinated chemicals could also be occurring.





Figure 4.30 Average conductivity for each treatment



In order to assess the residence time for the column reactors several KBr breakthrough tests were conducted. The first test was conducted by injecting a 60 mL plug of 6g/L KBr solution into the influent port of column 9 and the conductivity was measured at the top of the column. However, the conductivity measurements fluctuated with temperature and did not produce any reliable breakthrough curve.

In another attempt to quantify the residence time within the columns a second KBr breakthrough test was performed. KBr was initially injected into the influent port of column 9 and over a period 3.5 days samples were taken out at 27" above the influent port. KBr concentrations from the samples were measured using an IC and the results are shown below in Figure 4.31. After 3.5 days the peak and tail of the curve was still not detected.



Towards the end of the sampling period, in addition to 27", more samples were gathered at 36" and 45" above the influent port (see Figure 4.26). The time for KBr to travel from 27" to 36" and from 36" to 45" could be seen on the figure below. At 27" there was a concentration of 11.5 mg/L at the 58<sup>th</sup> hr and 36" reached the same concentration 11 hrs later. The same increase could be seen at the 36" and 45" point. At 36" there was a concentration of 9.4 mg/L at the 58<sup>th</sup> hr and port 6 reached approximately the same concentration 11 hrs later. The total length from influent port to the top most port is 54 inches. If it takes 11 hrs for KBr to travel 9 inches, then residence for the entire column could be interpolated to be 66hr or 2.75 day.



Figure 4.31 KBr breakthrough measurement



#### V. Discussion

The increase in methane concentration at middle of the mesocosms indicates that methane was being produced within the lower half the columns. It has been shown that methanogenesis occurs under anaerobic conditions (Wiedemeier et al., 1999) and the increases in methane concentration suggest that pockets of anaerobic conditions were developing in the lower half of the columns. This is important because microbial reductive dechlorination of PCE appears to require anaerobic conditions (Bouwer, 1991).

As the water travels up the columns less methane was detected towards the top. This suggests that methane loss is occurring at the upper half of the columns. One possibility is that methane could be oxidized by methanotrophic microbes residing within the root's rhizosphere and within the soil matrix. Plants roots could also be uptaking methane and subsequently volatilizing it into the atmosphere, thus reducing the methane concentration near the surface. Finally, another possibility is that methane is very volatile and could be escaping through the soil pores and into the atmosphere. At port 7 (54") along the column, (Figure 4.1) each of the columns with plants had a lower methane concentration when compare to the control columns without plants. This suggests that, while methane could be escaping through the soil pores, it is occurring at a much slower rate than either plant assisted methane degradation or volatilization. This experiment did not measure the volatilization rate within the column reactors so it can not be determined whether methane degradation or methane volatilization is the dominant process.

However, previously research showed that methane degradation exists during aerobic conditions where methanotrophic bacteria use methane as an energy source



www.manaraa.com

(Chapelle, 1993). Research also showed that co-metabolic degradation of chlorinated solvent could be carried out by methanotrophic bacteria when methane is the primary substrate (Eguchi et al, 2001; Little et al 1998).

Nitrate measurements in the results showed that nitrate concentration for all planted mesocosms increased near the top of the column and the unplanted control columns had no increase in nitrate concentration. Nitrate is produced from ammonium oxidation (Mitsch, 1993) and the increase in nitrate concentration at the tops suggests that ammonium oxidation is occurring. Research has also shown that ammonium oxidizing bacteria could also co-metabolically degrade TCE (Yang et al, 1999).

So, the methane concentration decrease and nitrate concentration increase above 36" suggests that co-metabolic TCE degradation could be occurring. Previous research has suggested that during co-metabolic TCE degradation, the TCE is broken down into TCE epoxide intermediate which then are converted by heterotrophic organisms into  $CO_2$  (Little et al, 1988). However, future research is needed to determine the contaminant volatilization rate of wetland plants in order confirm TCE is being degraded instead being phyto-volatilized into the atmosphere.

#### **Pre and Post Methane comparison**

Methane concentration pre-PCE injections were greater than post-PCE injection for all treatments. The methane within the reactors was produced by using carbon dioxide as an electron acceptor during methanogenesis (Chapelle, 1993). This process requires an electron donor and may compete with reductive dechlorination of PCE. PCE reduction reactions have low  $\Delta G_r^o$  and do not generate a lot of energy (Wiedemeier et al.,



1999). However, PCE reduction reactions have a higher  $\Delta G_r^o$  than methanogenesis (See Table 2.1) and PCE reducers could be competitive with methane generating bacteria for the limited amount of electron donor present in subsurface environment. Therefore, within a natural environment PCE reduction could take up the electron donors normally used for methane production, causing a decrease in methane concentration. The reduction in methane concentration post PCE injection suggests that reductive dechlorination reactions are taking place within the reactors at expense of methane production.

## **PCE concentration Decrease**

After 2 weeks of continuous PCE injections, the water in the reservoir will have been replaced 5 to 6 times. The results of 2<sup>nd</sup> week PCE concentration decrease between different column intervals is shown in figure 4.18. Between the column intervals 0" to 9", a student-t comparison of PCE removal showed that the *Carex comosa* PCE concentration decrease was significantly greater than control reactors. No significant differences were shown between the other planted treatments and the control columns. Between 0" and 9", the 2<sup>nd</sup> week TCE data (Figure 4.22) showed that the least amount of TCE was detected within the control reactors. Therefore, this suggested that low PCE concentration removal resulted in low concentration of TCE detected.

Between 9" and 15", *Carex comosa* had a significant drop in PCE concentration removal and *Eleocharis erythropoda* reactors were better at removing PCE at those depths. This suggested that *Carex comosa* could be a better candidate for PCE removal at lower depth below ground around 50" below ground, while *Eleocharis erythropoda* 



and the control columns are better at PCE removal slightly higher up around 40" below ground.

Above port 2 (15"), there was less removal of PCE concentrations and no statistically significant difference was detected between mesocosms. The decrease in PCE removal above port 2 suggest that methane and nitrate concentrations increase within the reactors maybe influencing PCE degradation.

Optimum PCE reduction occurs in the same redox zone as methanogenesis (Wiedemeier et al., 1999), so the fact that methane concentration increased at the bottom of the reactors (Figure 4.14 -4.17) indicates that conditions are favorable for PCE reduction. However, methanogenesis reactions also competes with PCE for electron donors (Chapelle, 1993). So the increase in methane production could also inhibit PCE reduction and could be the reason why less PCE was removed above 15"

Nitrate concentration in the groundwater would also influence PCE reduction rates. Nitrate reduction produces more energy than PCE reduction (Wiedemeier et al., 1999) and bacteria would preferentially use nitrate in the subsurface (Mitsch, 1993). So the increase in nitrate concentration at the top of planted columns (Figure 4.23) could also inhibit PCE reduction.

However, this was only 2 weeks after PCE injections began and the mesocosm within the columns are only beginning to adapt to the presence of a new chemical. It may not have developed the microbial capability to degrade PCE. Figure 5.1 below shows the absolute value of PCE concentration decrease as compared to the TCE concentration increase between the influent port and port 2 (0" to 15"). It could be seen that for all treatments the PCE concentration decrease was significantly greater than the



concentration of TCE formed. No comparisons were made for PCE concentration decrease between treatments due variability among influent concentration.

Significant amounts of PCE concentration decrease occurred between the 0" and 15" and the lack of TCE increase suggests several possibilities. One, PCE could be sorbed onto soil and organic particles instead of degrading. Two, TCE maybe degraded as well and the low TCE concentration shown on Figure 5.1 could be due to the fact that TCE has been degraded to DCE or VC which are harder to detect. Three, low concentration of TCE could also be due to TCE binding to soil particles. Previous experiments showed that significant amounts of TCE could bind to hydric soils much like the one used in this experiment (Enwright, 2002).



Figure 5.1 PCE & TCE comparison at week 2. PCE influent concentration and absolute value of PCE decrease and TCE increase between influent and port 2. With a P-value < 0.0001 there are significant differences among 0-15" TCE and PCE for all treatments



# 6<sup>th</sup> Week

After 6 weeks of continuous PCE injections, the PCE concentration removed between each port is shown in figure 4.16. A student-t comparison among mesocosms showed that between the interval 0" to 9" *Carex comosa* had the greatest concentration of PCE removed (see Figure 5.2).



6th week PCE removed between 0" and 9" and Influent Concentration

Figure 5.2 6th week comparison of PCE removal between 0" and 9". Comparisons and p-values are between *Carex comosa* and other treatments.

Analysis of other ports did not show any statistical significant PCE concentration difference between treatments. Figure 5.3 shows the total PCE concentration removed within the entire column (0" to 54") and there were no significant difference in the total amount of PCE removal between each treatment. So even though *Carex comosa* had greater reduction of PCE concentration between the 0" and 9", overall along the entire length there was no difference between the removal efficiency of all the treatments...





Figure 5.3 6th week PCE concentration decrease between 0" to 54", between 0" and 9" and the influent PCE concentration. There was no significant difference among treatments for 0" to 54" PCE removal. (p>0.1)



As with week 2 data, it is important to understand how much TCE was detected and compare it with the concentration of PCE removed. Figure 5.4 shows a comparison of PCE and TCE concentration difference between the influent port and port 2 (0" and 15"). It could be seen that the 6<sup>th</sup> week PCE concentration decrease was still greater than TCE concentration detected. However, the TCE concentration at week 6 was statistically greater than the TCE concentration at week 2. This data suggests that: One, less TCE sorption was occurring because more TCE was detected during week 6. Two, microbial colonies were beginning to establish themselves and starting to become more effective at PCE degradation at week 6. Three, the discrepancy between PCE and TCE concentration could be due to TCE degrading into daughters products (DCE isomers, VC, ethene, CO<sub>2</sub>) or being phyto-volatilized by plants (Lunney, 2004).



Figure 5.4 6th Week PCE removed between 0" and 15".  $2^{nd}$  and  $6^{th}$  week comparison of TCE removed between 0" and 15". Student t-test showed significant difference between the TCE increase at week 6 with TCE increase at week 2. P-values shows comparison between  $6^{th}$  and

2<sup>nd</sup> week PCE removal



TCE:

Since the 6<sup>th</sup> week results showed no significant difference in total PCE removal among all four treatments (Figure 5.3), the next question of concern is the removal efficiency of TCE. Figure 4.13 showed that the 6<sup>th</sup> week TCE concentration for all planted columns initially increased to a maximum in the middle of the reactors and then the concentration began to decrease along the upper half of the columns. The build up of TCE concentration within the bottom half of the reactors could be due to the slow rate of TCE reduction. Studies have shown that reductive dehalogenation of PCE could occur under anaerobic conditions but as the number of chlorine atoms on the molecule decrease the rate of reductive dechlorination also decreases (Chapelle, 1993). So TCE with one less chlorine atom, would take longer to degrade, causing a temporary TCE build up in the lower half of the reactors. In the upper half of the reactors, TCE removal has begun to occur, as seen by the decreasing concentration of TCE within the planted columns (Figure 4.9). This suggests that plants have an influence on TCE concentration and possible mechanisms behind the removal will be explained below.

The control treatments also had a TCE concentration increase in the lower half of the reactors but it did not exhibit the same TCE concentration decrease in the upper half of the reactors, as seen in the planted columns. This was especially true between ports 6 and 7 (45" and 54") where the control column TCE concentration increased to a maximum level.

Therefore, a student-t test with an alpha=0.1 was conducted on the week 6 TCE concentration found at port 7 (54"). Port 7 is the highest port on the column and would be most representative of the effluent concentration. Figure 5.5 below shows that at port7



www.manaraa.com

the average TCE concentration within the control reactors was statistically greater than the TCE concentration found within the planted reactors. However, a comparison among each of the planted reactors showed no significant difference among the planted reactors.



Port 7 Average TCE concentration for week 6

Figure 5.5 Average TCE concentrations from port 7. P-values on the figure are generated from a student-t test between the control reactors and each of the planted reactors.

Since the decrease in TCE concentration occurred only for planted columns, the results supported the fact that plants have an effect on the TCE concentrations in the subsurface. The presence of roots near the top of the columns could affect TCE concentrations in several ways. Plant roots could uptake TCE through phytoextraction and then volatilize gaseous TCE into the atmosphere through phytovolatilization (Lunney, 2004). Plants could also be transporting oxygen into the soil (Armstrong et. al, 2000; Bankston, 2002) which in turn support aerobic co-metabolic degradation of TCE.



Plant roots extrude photosynthetic products such as amino acids, sugar, and vitamins into the rhizosphere (Walton, 1994) and the substrates could then be used to stimulate microbial facilitated degradation.

Plant uptake of TCE and oxygen transport within the plant root was not investigated during this experiment. Future study is also needed to assess the volatilization rate with plants to determine how much contaminant is being volatilized into the atmosphere. Plant tissues could also be examined to determine how much contaminants has been sequestered into plant tissues such as roots and leaves.

In this experiment 1,2 cisDCE, 1,2 transDCE, and ethylene was not detected. These chemicals are degradation products of PCE and the non-detection of these chemicals suggest several possibilities. One, previous research showed that *Dehalococcoides* could degrade PCE all the way to ethane(Smidt and Vos, 2004). Previous research also showed that iron reducer could degrade DCE isomers and VC (Lee, 1998). So if the conversion rate PCE degradation products are very quick, then no accumulation would be detected. Two, co-metabolic degradation of TCE results in the production of  $CO_2$  (Little et al, 1988). A study has proposed that during co-metabolic TCE degradation, TCE is first converted into TCE epoxide and then it is degraded by heterotrophic bacteria into  $CO_2$  (Little et all, 1988). The carbon dioxide could then be dissolved in water and become part of the carbonate cycle or it could escape through soil pore into the atmosphere. Therefore no DCE or VC would accumulate during cometabolic TCE degradation. Three, plants could have an influence on the concentrations of these chemicals. Phytoextraction and phytovolatilization (Lunney, 2004) could allow possible transport of these products into the atmosphere.



#### Sulfate and Nitrate comparison

Sulfate and nitrate concentration within the column reactors were investigated in this experiment. Figure 5.6 below overlaid the nitrate reading above the TCE concentration to compare the trend between nitrate and TCE (Nitrate concentration is ppm). Below 27 inches the nitrate concentration did not significantly increase. In that same zone the majority of TCE has been detected and nitrate did not appear to have any effect on TCE production. However, it is unclear whether more TCE could have been detected if the nitrate concentration was lower.

Above 36", the nitrate concentration for the planted mesocosms started to increase. This suggested that plant roots were having an influence on soil chemistry at those depths. Above 36" TCE concentrations started to decline for all planted columns. This suggested two possibilities: One, nitrate is produced from ammonia oxidation and the enzymes used during ammonia oxidation could also co-metabolize TCE (Yang, 1999). So the increased nitrate and decreased TCE concentration could indicate that TCE is being co-metabolically degraded into CO<sub>2</sub>. Two, the wetland plants could also influence TCE concentration through uptake and venting.

Above 36", the unplanted control reactors had no increase in nitrate concentration. This reiterated the suggestion that plants do have an influence on the nitrate concentration. At the same depth TCE concentration also increased within the control reactors. Again this suggest two possibilities: One, microbes reducing nitrate out competes PCE reducing microbes (Chapelle, 1993) and the presence of nitrate would inhibit microbial reductive dechlorination of PCE. Therefore, within the unplanted control there was less nitrate available to inhibit PCE reduction and TCE concentration



would increase. Two, the lack of plants could be causing TCE to accumulate within the columns, since no uptake or volatilization could occur.



Figure 5.6 Overlay of nitrate trend with 6th week TCE concentration for comparison.



Figure 5.7 below overlaid the sulfate reading above the TCE concentration to compare the trend between sulfate and TCE (sulfate concentration is in ppm) Sulfate reduction has approximately the same redox potential as PCE reduction (see Figure 2.4). The decrease in sulfate concentration suggests that sulfate reduction reactions are taking place and that PCE reduction could also occur. TCE was compared with sulfate because any TCE detected within the reactors be attributed to PCE degradation. According to the figure there is no correlation between sulfate concentration and TCE production. Along the length of the column TCE concentration are varying independent of the sulfate concentration.



Figure 5.7 Overlay of sulfate trend with 6th week TCE concentration for comparison.



## Methane and TCE

Methane production in subsurface competes with reductive dechlorination for electron donors (Wiedemeier et al., 1999). So an increase in methane concentration in the reactors could also impact the PCE removal rate. Figure 5.8 -5.11 overlays the methane with TCE concentration and compares the trend of two species. TCE was used in the comparison instead of PCE because TCE detected in the columns could be attributed to PCE degradation.

The figures below are taken from the 6th week samples and below 18 inches TCE concentration increased as methane increased. This suggests that the bottom of the column provides the necessary reductive environment for both methanogenesis and reductive dechlorination. However, between 18" and 36" TCE concentration for all treatment remained steady and did not have a large increase. At the same depth the methane concentration varied among treatments. The methane concentration for the planted columns reached a maximum at different depth: *Carex comosa* reached a maximum at 27", for *Scirpus atrovirens* at 21", for *Eleocharis erythropoda* at 36". Since the TCE concentration leveled off at the middle of the column and methane concentration increased, it suggest that methanogenesis is out competing PCE degradation. The leveling off of TCE concentration could also suggest that iron reducing microbes could be reducing TCE to a DCE isomer (Lee et al, 1998).

Above 36" inches, both methane and TCE concentration decreased for the planted columns. However the control columns continued to increase in both methane and TCE. This suggests that plants have an influence on both methane and TCE concentration. In the planted columns, co-metabolic degradation of TCE could be taking



www.manaraa.com

place or plants could directly uptaking and venting TCE. In the control column no plants are available to transport oxygen to the root rhizosphere (Armstrong, 2000) and therefore methane oxidation and TCE co-metabolic degradation may be inhibited. Another possible suggestion for the TCE increase within the control columns could be TCE accumulating within the columns because plants are not available for phytovolatilization.





Figure 5.8 Overlay of Carex comosa reactor methane and TCE concentration for comparison



Figure 5.9 Overlay of Scirpus atroviren reactor methane and TCE concentration for comparison





Figure 5.9 Overlay of Eleocharis erythropoda reactor methane and TCE concentration for comparison



Figure 5.9 Overlay of control reactor methane and TCE concentration for comparison



#### **VI** Conclusion

The purpose of this research was to study three different wetland plant species and an unplanted control, in order to determine which plant species was best for chlorinated solvent removal. A low concentrated PCE solution will be injected into the plant mesocosm and then samples will be gathered and analyzed to determine the concentrations of chlorinated ethenes within the mesocosms. This studied was modeled after the constructed wetland at WPAFB and column reactors were constructed to determine the PCE removal efficiency among each plant species. A total of seven sampling ports were spaced along the length of the columns to allow for better assessment of contaminant concentration at different depth within the columns. A sampling methodology was developed for the column reactors and samples were gathered for conductivity, chlorinated solvent, nitrate and sulfate measurements.

#### Answers to specific research questions

1. Is there a significant difference in chlorinated solvent degradation among the plant species and the unplanted control treatment?

The presence of TCE in the columns demonstrated that PCE was being degraded. However, after 6 weeks of continuous PCE injections there were no significant differences in total PCE removal among any of the treatments. Between the influent port and port 7 (0" to 54"): *Carex comosa* removed 48.14 +/- 0.797 ppb, *Scirpus atrovirens* removed 47.14 +/- 3.054, *Eleocharis erythropoda* removed 47.19 +/- 7.304 ppb, and the control column removed 47.139 +/- 2.062 ppb. All treatments including the control reactors removed approximately the same concentration of PCE. However, the PCE concentration removed does not differentiate between removal mechanisms. Sorption,



degradation, phyto-volatilization, and phyto-assimilation are all combined to produce the removal results shown.

The total PCE degradation could be determined by analyzing the concentration of TCE formed. At port 7 ( 54") the TCE concentration are as follows: *Carex comosa* TCE concentration was 6.463 +/- 2.822 ppb, *Scirpus atrovirens* TCE concentration was 6.510 +/- 1.383, *Eleocharis erythropoda* TCE concentration was 6.037 +/- 2.118 ppb, and the control column TCE concentration was 12.833 +/- 5.191 ppb.

There was no significant TCE concentration difference between the planted treatments. However, the control columns had significantly greater TCE concentration at the 54" near the top the column. This suggests several possibilities: One, without plants, phytovolatilization will not occur and thus TCE may be accumulating within the mesocosms. Two, the control mesocosm had high methane concentrations (Figure 4.13) at the top of the reactors. The accumulation in methane suggested that both methane oxidation and co-metabolic TCE degradation maybe inhibited . Thus, TCE would not accumulate within the columns. Third, the control columns had no increase in nitrate concentration suggesting that ammonia oxidation was not occurring and again co-metabolic TCE degradation was not occurring. Thus leading to an increase in TCE concentration.

TCE is monitored by the EPA and any treatment option would also require the removal of TCE from the system. So even though there were no differences in PCE degradation among mesocosms, the unplanted control mesocosm was not efficient at TCE removal. Therefore plants are needed in any constructed wetland pollution



treatment option. However, the planted columns showed no significant difference in total PCE removed and it also showed no difference in the TCE concentration at port 7. More studies are needed to determine possible contaminant removal mechanism with each plant. But based on just my results all three wetland plants species under observation could be viable candidate for wetland pollution treatments.

2. Is there a difference in sulfate, nitrate and methane concentration among the reactors?

There was a difference in the sulfate concentration among the reactors. At the bottom of the column at 0" and 9" *Eleocharis erythropoda* had significantly greater sulfate concentration. Between 9" and 45" there was no significant difference in sulfate concentration among all treatments. At 54" the control columns had significantly greater sulfate concentration than *Eleocharis erythropoda* and a comparison of the other treatments did not show any significant difference. Even though sulfate had some differences between treatments, the results were not consistent with expectations and no conclusions could be drawn.

Between 0" and 36" there was no significant difference in nitrate concentration among all the treatments. However, at 45", a student-t analysis with alpha=0.1 found the nitrate concentration for *Scirpus atrovirens* to be statistically greater than both *Eleocharis erythropoda* and control columns (p-value = .003). All other treatments have no statistical difference. At 54" there was no statistical difference among all treatments.

The methane concentration below 18" increased for all treatments. This suggests that the bottom of the column provides the necessary reductive environment for both



methanogenesis. However, between 18" and 36" the methane concentration varied among treatments. The methane concentration for the planted columns reached a maximum at different depth: *Carex comosa* reached a maximum at 27", for *Scirpus atrovirens* at 21", for *Eleocharis erythropoda* at 36".

Above 36" inches, methane decreased for the planted columns. However the control columns continued to have methane increase. This suggests that plants are influencing methane concentration, methane are removed either through aerobic methane oxidation or plants themselves are uptaking methane.

#### 3. Do the sulfate, nitrate, and methane concentration influence PCE degradation?

Results showed that sulfate concentration did not influence PCE degradation (Figure 5.7). However, nitrate concentrations did have an influence on PCE degradation. Figure 5.6 showed that above 36", the nitrate concentration for the planted columns increased, suggesting that plant roots were having an influence on groundwater chemistry at those depths. At those same depths TCE concentration started to decrease for the planted columns. So increasing nitrate concentration suggests that less TCE was formed or that TCE was being co-metabolically degraded to CO<sub>2</sub>.

The control reactors maintained the same nitrate concentration between 36" and 54 " and TCE concentrations in the control reactors started to increase at those depth. This suggested that the lack of nitrate increase was causing TCE to accumulate within the mesocosm.



Methane also influence TCE degradation and below 18" the methane and TCE concentration increased together. However, above 18", TCE concentration remained steady while methane concentration for all treatments increased. This suggests that when methane concentration increased over a certain threshold it inhibited PCE degradation. Finally, above 36" methane and TCE both decreased for the planted columns. Suggesting that methane could be oxidized at those depth and co-metabolic degradation of TCE could be taking place.

## Limitations

Some benefits that may be associated with plant assisted contaminate remediation include direct removal of chlorinated solvents through the mechanism of phytoextraction, phytodegradation, and phytostabilization. This laboratory study was conducted under controlled conditions and does not fully reflect natural conditions. The temperature inside the greenhouse laboratory never dropped below 15 °C and lighting was controlled to give at least 13 hrs of light each day. These conditions were quite different from the constructed wetland, where the climate varies considerably during each season.



# Appendices

# Appendix A: Chlorinated Ethene Sampling Method

All samples for the gas chromatograph were gathered with a Hamilton Gastight

10mL syringes. Samples are stored in a 10 mL serum bottles and capped with a 20mm

Teflon coated serum stopper and aluminum crimps. The following list goes over the

sampling procedure used during chlorinated ethene sampling.

Sample bottle preparation:

- 1. Cap each serum bottle with Teflon coated caps and crimp tightly.
- 2. Flush each serum bottle with approximately 60 mL of nitrogen gas

# Sampling Methodology:

- 1. Start at the column one and move laterally, sampling the top ports (port 7) of each consecutive column.
- 2. After port 7 of all the columns have been sampled, proceed to the next lower port (port 6) and again move laterally until all of the columns have been sampled. Proceed in this pattern until all desired ports are sampled.

# Port sampling procedure:

- 1. Open desired port and purge the dead space in the sampling tubing by draining and discarding 15 mL of liquid. Measure with graduated cylinder. This flushes out the dirt blockage inside the sampling tube, as well as any liquid that accumulated in the dead space.
- 2. Afterwards, attach a 10mL gastight syringe to the port and fill syringe by slowly drawing back on the plunger.
- 3. After sampling tilt the syringe with needle facing up and eject any bubbles inside the syringe.
- 4. If sample fluid contains dirt particles and the fluid is not transparent, keep the syringe inverted and allow dirt particles to settle for 10 minutes.
- 5. Keeping the syringe inverted, inject 5mL into an inverted15 mL serum bottle. Do not inject any of the soil particles from the bottom of the syringe.
- 6. Keep the serum bottle inverted, to be transported back to lab



- 7. Spin sample bottle overnight for 12 hrs to achieve equilibrium. Keep the bottles inverted at all times to maintain a water seal and prevent gas from escaping.
- 8. After 12 hrs perform head space analysis on the GC.



# Appendix B: GC 6890 Run Method

		6890	GC	METHOD			
OVEN Initial	temp:	50	'C	Maximum	temp:	240	min
Ramos	une.	2	111111	Lyuiibration	une.	1	
#	Rate	Final Temp		Final Time			
1	10	100		1.25			
2	0.0(Off)						
Post	temp:	50	'C				
Post	time:	0	min				
Run	time:	8.25	min				
FRONT INLET				BACK INLET			
Mode:	Splitless			Mode:	Split		
Initial Temp:	200	'C	(On)	Initial Temp:	50	'C	
Pressure:	5.45	psi	(On)	Pressure:	0	psi	(Off)
Purge flow	19.7	mL/min		Total Flow:	45	mL/min	
Purge Time:	0	min		Gas saver:	Off		
Total Flow:	24.2	mL/min		Gas type:	Helium		
Gas Saver:	Off						
Gas Type:	Helium						
COLUIVIN I	Column			COLUIVIN 2			
Model Number		112/222		Model Number	Agilopt	0001\/ 41	2
	J&VV	1134332		Wodel Number:	Agilent	90910-41	3
GS Gas Plo	220			HP-624 Special analy		10	
Nominal Longth:	230	C		Nominal Longth:	200		
Nominal Diamatari	30	111		Nominal Length.	30	111	
Nominal film thickness:	320	um		Inominal Diameter.	320	um	
Mode:	U Constant flow		r	Modo:	1.8	um	
INIOUE.		ml /min		Ividue.	2	ml /min	
Initial FIOW:	2.1	mL/min			2	mL/min	
Nominal Init pressure:	9.85	psi em/eee			9.84	psi om/ooo	
	54 Eront Inlot	CIII/Sec		Average velocity.	Eropt Inlot	CIII/SEC	
Outlot:	Front dotoctor			Outlot:	Profit Inlet		
Outlet.	ambiont			Outlot Prossuro:	ambiont		
Oullet Flessule.	amplent			Oullet Flessule.	anden		
FRONT	DETECTOR	(FID)		BACK	DETECTOR	(ECD)	
Temperature:	250	'C	(On)	Temperature:	250	'C	(On)
hydrogen flow	40	mL/min	(On)	Anode Purge flow:	6	mL/min	(On)
Air flow:	450	mL/min	(On)	Mode:	Constant Makeup Flow		
Mode:	istat makeup flo	W		Makeup Flow:	60	mL/min	(On)
Makeup flow:	45	mL/min		Makeup Gas Type	Nitrogen		
Make up gas type:	Nitrogen			Adjust offset	60		
Flame:	On						
Electrometer:	On						
Lit Offset:	2						
SIGNAL	1			SIGNAL	2		
Data Rate	20	Hz		Data Rate	20	H7	
Type:	Front Detector			Type:	back detector		
Save Data:	On			Save Data:	On		
Zero	0	(Off)		Zero <sup>.</sup>	0	(Off)	
Range:	õ	(0)		Range:	õ	(0)	
Fast Peaks	Off			Fast Peaks	Off		
Attenuation:	0			Attenuation:	0		
COMP	1			COLUMN	COMP	2	
Derive	from	front	detector	Derive	from	back	detector



www.manaraa.com

Brad Improved Short Program:	Bromide Tracer JY
Cartridge serial number: 040602435015 type: EluGen-OH.	Cartridge serial number: 040602435015 type: EluGen-OH.
Pressure.LowerLimit = 200	Pressure.LowerLimit = 200
Pressure.UpperLimit = 2500	Pressure.UpperLimit = 2500
%A.Equate = "Water"	%A.Equate = "Water"
%B.Equate = "%B"	%B.Equate = "%B"
%C.Equate = "%C"	%C.Equate = "%C"
%D.Equate = "%D"	%D.Equate = "%D"
Flush Volume = 100	Flush Volume = 100
Wait FlushState	Wait FlushState
Wait finished	Wait finished
NeedleHeight = 5	NeedleHeight = 5
CutSegmentVolume = 10	CutSegmentVolume = 10
SyringeSpeed = 3	SyringeSpeed = 3
ColumnTemperature = 30	ColumnTemperature = 30
Cycle = 0	Cycle = 0
Data_Collection_Rate = 2.0	Data_Collection_Rate = 2.0
Temperature_Compensation = 1.7	Temperature_Compensation = 1.7
Oven_Temperature = 30	Oven_Temperature = 30
Suppressor_Type = SRS	Suppressor_Type = SRS
Suppressor_Current = 100	Suppressor_Current = 135
Flow = 1.50	Concentration = 35.00
%B = 0.0	EluentGenerator.Curve = 5
%C = 0.0	Flow = 1.50
%D = 0.0	%B = 0.0
Pump.Curve = 5	%C = 0.0
WaitForTemperature = False	%D = 0.0
Wait SamplePrep	Pump.Curve = 5
Wait finished	WaitForTemperature = False
Concentration = 1.00	Wait SamplePrep
EluentGenerator.Curve = 5	Wait finished

Appendix C: IC run method



# Appendix D: Student-t P-value calculations

2<sup>nd</sup> week PCE p-value comparison of PCE removal between ports

Port 6 to 7					
Level	Minus Level	Difference	Lower CL	Upper CL	p-Value
10-1 Con PCE	10-1 CC PCE	0.749667	-1.49438	2.993716	0.463224
10-1 SA PCE	10-1 CC PCE	0.6045	-1.52439	2.733393	0.530974
10-1 EE PCE	10-1 CC PCE	0.375	-1.86905	2.61905	0.71002
10-1 Con PCE	10-1 EE PCE	0.374667	-1.63247	2.381806	0.678226
10-1 SA PCE	10-1 EE PCE	0.2295	-1.64801	2.107007	0.785191
10-1 Con PCE	10-1 SA PCE	0.145167	-1.73234	2.022673	0.862922

5 to 6	_				
Level	Level	Difference	Lower CL	Upper CL	p-Value
10-1 CC PCE	10-1 EE PCE	2.032667	-2.89966	6.964993	0.369767
10-1 SA PCE	10-1 EE PCE	1.858917	-2.26776	5.985597	0.329286
10-1 Con PCE	10-1 EE PCE	1.623	-2.78861	6.034607	0.420893
10-1 CC PCE	10-1 Con PCE	0.409667	-4.52266	5.341993	0.852882
10-1 SA PCE	10-1 Con PCE	0.235917	-3.89076	4.362597	0.898373
10-1 CC PCE	10-1 SA PCE	0.17375	-4.50547	4.852966	0.933867

4 to 5					
Level	Level	Difference	Lower CL	Upper CL	p-Value
10-1 EE PCE	10-1 CC PCE	2.507	-3.59177	8.605768	0.363423
10-1 EE PCE	10-1 SA PCE	1.824	-2.20996	5.857956	0.32046
10-1 EE PCE	10-1 Con PCE	1.645667	-2.66681	5.958147	0.396855
10-1 Con PCE	10-1 CC PCE	0.861333	-5.23743	6.960101	0.748193
10-1 SA PCE	10-1 CC PCE	0.683	-5.22211	6.588107	0.792364
10-1 Con PCE	10-1 SA PCE	0.178333	-3.85562	4.212289	0.919677

3 to 4					
Level	Level	Difference	Lower CL	Upper CL	p-Value
10-1 EE PCE	10-1 Con PCE	2.83	-2.06682	7.726824	0.219337
10-1 CC PCE	10-1 Con PCE	1.987833	-3.48698	7.462649	0.426739
10-1 EE PCE	10-1 SA PCE	1.637667	-2.94289	6.218226	0.433573
10-1 SA PCE	10-1 Con PCE	1.192333	-3.38823	5.772893	0.564945
10-1 EE PCE	10-1 CC PCE	0.842167	-4.63265	6.316982	0.731967
10-1 CC PCE	10-1 SA PCE	0.7955	-4.39837	5.989366	0.733071



Port 2 to 3

			Lower	Upper	
Level	Minus Level	Difference	CL	CL	p-Value
10-1 Con	10-1 CC		-		
PCE	PCE	1.322333	4.01346	6.658131	0.583369
10-1 Con	10-1 EE		-		
PCE	PCE	0.943333	3.82915	5.715816	0.660647
10-1 Con	10-1 SA		-		
PCE	PCE	0.722833	3.74142	5.187082	0.718567
	10-1 CC		-		
10-1 SA PCE	PCE	0.5995	4.46248	5.661482	0.791691
	10-1 CC				
10-1 EE PCE	PCE	0.379	-4.9568	5.714798	0.873955
	10-1 EE		-		
10-1 SA PCE	PCE	0.2205	4.24375	4.684749	0.912124

#### Port 1 to 2

			Lower	Unner	
Level	Minus Level 10-1 CC	Difference	CL	CL	p-Value
10-1 EE PCE 10-1 Con	PCE 10-1 CC	7.085833	-7.2229	21.3946	0.286498
PCE	PCE 10-1 CC	4.6595	-9.6493	18.96826	0.474204
10-1 SA PCE	PCE 10-1 SA	4.03075	-9.5437	17.60523	0.512857
10-1 EE PCE	PCE 10-1 Con	3.055083	-8.9165 -	15.02665	0.572445
10-1 EE PCE 10-1 Con	PCE 10-1 SA	2.426333	10.3718	15.22448	0.67354
PCE	PCE	0.62875	11.3428	12.60032	0.906589

#### Port iff to 1

			Lower	Linner	
Level	Minus Level 10-1 Con	Difference	CL	CL	p-Value
10-1 CC PCE	PCE 10-1 SA	20.06317	-2.6238	42.75012	0.075755
10-1 CC PCE	PCE 10-1 Con	16.267	-5.2557	37.78973	0.119518
10-1 EE PCE	PCE 10-1 SA	12.42267	-7.8692 -	32.71449	0.195718
10-1 EE PCE	PCE 10-1 EE	8.6265	10.3548 -	27.60777	0.325257
10-1 CC PCE	PCE 10-1 Con	7.6405	15.0465 -	30.32745	0.45973
10-1 SA PCE	PCE	3.79617	15.1851	22.77743	0.656942



# 6<sup>th</sup> week PCE p-value comparison of PCE removal between ports

Port 6 to 7 Level 11-3 EE PCE 11-3 EE PCE 11-3 EE PCE 11-3 CC PCE 11-3 CC PCE 11-3 SA PCE	Minus Level 11-3 Control PCE 11-3 SA PCE 11-3 CC PCE 11-3 Control PCE 11-3 SA PCE 11-3 Control PCE	Difference L 2.0625 2.02225 1.1745 0.888 0.84775 0.04025	Lower CL U -0.975 -0.85938 -2.15292 -2.1495 -2.03388 -2.50111	Jpper CL p 5.100004 4.90388 4.501919 3.925504 3.72938 2.581609	-Value 0.152395 0.140991 0.431471 0.511653 0.509076 0.971171
Port 5-6 Level 11-3 Control PCE 11-3 SA PCE	Minus Level 11-3 EE PCE 11-3 EE PCE	Difference L 6.750667 5.0885	ower CL l 0.03737 -1.28029	Jpper CL p 13.46396 11.45729	o-Value 0.049041 0.100783
11-3 CC PCE 11-3 Control PCE 11-3 Control PCE 11-3 SA PCE	11-3 EE PCE 11-3 CC PCE 11-3 SA PCE 11-3 CC PCE	4.0715 2.679167 1.662167 1.017	-3.28255 -4.03413 -3.95458 -5.35179	11.42555 9.39246 7.27891 7.38579	0.231817 0.376754 0.506656 0.71692
Port 4-5 Level 11-3 EE PCE 11-3 Control PCE 11-3 SA PCE 11-3 EE PCE 11-3 Control PCE 11-3 EE PCE	Minus Level 11-3 CC PCE 11-3 CC PCE 11-3 CC PCE 11-3 SA PCE 11-3 SA PCE 11-3 Control PCE	Difference L 2.813 2.296167 1.657 1.156 0.639167 0.516833	ower CL l -3.1187 -3.11871 -3.48 -3.981 -3.89124 -4.89804	Jpper CL p 8.744701 7.711044 6.794004 6.293004 5.169578 5.931711	0.299116 0.349394 0.470537 0.611107 0.748445 0.827886
Port 3-4 Level 11-3 SA PCE 11-3 CC PCE 11-3 EE PCE 11-3 SA PCE 11-3 SA PCE 11-3 CC PCE	Minus Level 11-3 Control PCE 11-3 Control PCE 11-3 Control PCE 11-3 EE PCE 11-3 CC PCE 11-3 EE PCE	Difference L 1.73525 1.3955 1.0625 0.67275 0.33975 0.333	Lower CL U -3.32276 -4.64997 -4.98297 -5.06249 -5.39549 -6.28948	Jpper CL p 6.793256 7.440973 7.107973 6.407989 6.074989 6.955484	-Value 0.443937 0.602132 0.690157 0.789507 0.892544 0.908694
Port 2-3 Level 11-3 EE PCE 11-3 EE PCE 11-3 EE PCE 11-3 CC PCE 11-3 SA PCE 11-3 CC PCE	Minus Level 11-3 Control PCE 11-3 SA PCE 11-3 CC PCE 11-3 Control PCE 11-3 Control PCE 11-3 SA PCE	Difference L 3.294667 2.68075 2.397 0.897667 0.613917 0.28375	Lower CL U -11.8225 -11.6606 -14.163 -14.2195 -12.034 -14.0576	Jpper CL p 18.41183 17.02215 18.95702 16.01483 13.26184 14.62515	-Value 0.622179 0.671819 0.742194 0.892289 0.911845 0.963991
Port 1-2 Level 11-3 EE PCE 11-3 Control PCE 11-3 SA PCE	Minus Level 11-3 CC PCE 11-3 CC PCE 11-3 CC PCE	Difference L 7.007 6.805333 5.4095	ower CL l -12.8348 -11.3077 -11.774	Jpper CL p 26.84881 24.91834 22.59301	-Value 0.431263 0.403806 0.480871


### Week 6, port 7 TCE Student t test results

Level	Minus level	Difference Low	er CL	Upper CL	p-Value
Con TCE	EE TCE	4.74E-05	-8.01E-07	9.55E-05	0.052979
Con TCE	CC TCE	4.44E-05	-4.00E-06	9.25E-05	0.065837
Con TCE	SA TCE	4.41E-05	3.80E-06	8.43E-05	0.036177
SA TCE	EE TCE	3.30E-06	-4.20E-05	4.90E-05	0.869194
CC TCE	EE TCE	3.00E-06	-5.00E-05	5.57E-05	0.8968
SA TCE	CC TCE	3.00E-07	-4.50E-05	4.60E-05	0.988043

2<sup>nd</sup> week Comparison of PCE decrease and TCE increase between 0" and 15"

Level	minus Level	Diff.	Lower CL	Upper CL	P-value
CC PCE	CC TCE	0.000234	0.000148	0.00032	0.0000291
SA PCE	SA TCE	0.000155	0.000094	0.0002161	0.000058
EE PCE	EE TCE	0.00021	0.00014	0.0002806	0.0000097
CON PCE	Con TCE	0.000129	0.000058	0.0001989	0.0013278

2<sup>nd</sup> and 6<sup>th</sup> week Comparison TCE increase between 0" and 15"

Level	minus Level	Difference	Lower CL	Upper CL	p-Value
CC TCE6	CC TCE2	4.44E-05	2.30E-05	6.55E-05	0.0004401
SA TCE6	SA TCE2	5.09E-05	3.60E-05	6.58E-05	0.0000028
EE TCE6	EE TCE2	2.66E-05	7.30E-06	4.59E-05	0.0100556
Con TCE6	Con TCE2	5.55E-05	3.80E-05	7.28E-05	0.0000053



Appendix	E.	$2^{nd}$	week	data
----------	----	----------	------	------

						Col 1	CC3 Col	umn 1-6	6						
RT:	7.565	5.	485	2.5	588 DOF	3. Tron	171 •DCE	,		C th	vlana	E th		1.5	33
ports	P.A. Cond	P.A.	CE Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
54	56.34 0.87	13.58	0.9723											19.77	4.93
45	40.19 0.62	23.19	1.6604											299.63	74.79
36	182.63 2.83	25.61	1.8337											377.46	94.21
21	260.1 4.03	38.4	2.7494											253.34	63.23
15	188.63 2.92	19.09	1.3668											158.92	39.67
9	569.11 8.82	6.9	0.494											70.33	17.55
0	2910.7 45.12	0	0											0	0.00
RT	7 565	5	485	1		Col 2	CC2 (	Jolumn				1		15	33
	PCE	T	CE	Cis	DCE	Tran	sDCE	Ň	VC	Eth	ylene	Eth	nane	Meth	nane
ports	P.A. Conc	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
6	7.474 0.12	7.82	0.5599											0 470 57	0.00
5	190.64 2.95	70.1	5.0192											328.3	81.94
4	219.31 3.40	24.67	1.7664											576.53	143.90
3	308.43 4.78	19.9	1.4248											520.87	130.01
2	376.43 5.83 756.44 11.7	19.2	1.3747											372.77 52.18	93.04 13.02
IFF	2530.6 39.2	0	0											0	0.00
			- '			Col 3	Control	3 column	n						
RT:	7.565	5.	485											1.5	33
porte	PCE	Т	CE	Cis	DCE	Tran	sDCE		VC	Eth	ylene	Eth	nane	Meth	nane
7	178.11 2.76	25.96	1.8587	г. <del>А</del> .	CONC.	г. <del>А</del> .	CONC.	г. <del>А</del> .	COILC.	г.А.	CONC.	г. <del>А</del> .	COILC.	846.75	211.35
6	327.89 5.08	23.97	1.7163											966.07	241.13
5	215.74 3.34	15.94	1.1413											684.44	170.84
4	446.54 6.92	72.88	5.2182											278.27	69.46 70.40
2	252.33 3.91	20.1	1.4392											308.81	79.49
1	676.39 10.48	3.94	0.2821											21.39	5.34
IFF	2252.6 34.92	0	0											0	0.00
			-			0.14	550							Ű	
RT:	7,565	5.	485			Col 4	EE3 (	column				I 		1.5	33
RT:	7.565 PCE	5. T	485 CE	Cis	DCE	Col 4 Tran	EE3 ( sDCE	column	VC	Eth	ylene	Eth	nane	1.5 Meth	i33 nane
RT:	7.565 PCE P.A. Conc	5. T P.A.	485 CE Conc.	Cis P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A.	i33 nane Conc.
RT: ports 7 6	7.565 PCE P.A. Conc 14.96 0.23 16.33 0.25	5. T P.A. 14.93 23.42	485 CE Conc. 1.069 1.6769	Cisl P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 172.42 687.37	533 nane Conc. 43.04 171.57
RT: ports 7 6 5	7.565 PCE P.A. Conc 14.96 0.23 16.33 0.25 111.5 1.73	5. T P.A. 14.93 23.42 34.36	485 CE Conc. 1.069 1.6769 2.4602	Cis P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 o sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 172.42 687.37 455.55	i33 nane Conc. 43.04 171.57 113.71
RT: ports 7 6 5 4	7.565         PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37	5. T P.A. 14.93 23.42 34.36 26.9	485 CE Conc. 1.069 1.6769 2.4602 1.926	Cisl P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 172.42 687.37 455.55 454.84	533 hane Conc. 43.04 171.57 113.71 113.53
RT: ports 7 6 5 4 3	7.565 PCE P.A. Conc 14.96 0.23 16.33 0.25 111.5 1.73 282.07 4.37 733.6 11.3	5. T P.A. 14.93 23.42 34.36 26.9 60.44	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275	Cisl P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 172.42 687.37 455.55 454.84 124.01	533 Conc. 43.04 171.57 113.71 113.53 30.95 50.95
RT: ports 7 6 5 4 3 2 1	7.565 PCE P.A. Conc 14.96 0.23 16.33 0.25 111.5 1.73 282.07 4.37 733.6 11.3 473.19 7.33 468.38 7 26	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069	Cisl P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83	533 hane Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34
RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.33           473.19         7.32           468.38         7.26           2273.6         35.2'	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069 0	Cisl P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0	i33 nane Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00
RT: 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.33           468.38         7.26           2273.6         35.24	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069 0	Cisl P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0	i33 Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT:	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.33           473.19         7.33           468.38         7.26           2273.6         35.24	5. T P.A. 14.93 23.426 34.36 26.9 60.44 15.5 15.46 0	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069 0	Cisl P.A.	DCE Conc.	Col 4 Tran P.A. Col 5	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0	i33 conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT:	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.3'           473.19         7.33           468.38         7.26           2273.6         35.2'           7.565         PCE           PA         Conc	5. T P.A. 14.93 23.426 34.366 26.9 60.44 15.5 15.46 0 5. T	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069 0 485 CE	Cisi P.A. Cisi	DCE Conc.	Col 4 Tran P.A. Col 5 Tran	EE3 of sDCE Conc. SA3 of sDCE	P.A.	VC Conc.	Eth P.A. Eth	ylene Conc. ylene	Ett P.A. Ett	nane Conc.	1.5 Mett P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0	33 hane Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: 7 Ports 7	7.565 PCE P.A. Conc 14.96 0.23 16.33 0.25 111.5 1.73 282.07 4.37 733.6 11.33 468.38 7.26 2273.6 35.2 7.565 PCE P.A. Conc 80.68 1.25	5. P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 T P.A. 12.98 P.A.	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069 0 485 CE Conc. 0.9294	Cis P.A. Cis P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Etr P.A. Etr P.A.	nane Conc. nane Conc.	1.5 Meth P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Meth P.A. 0	33 hane Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 33 hane Conc. 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.33           468.38         7.26           2273.6         35.24           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 T P.A. 12.98 16.13	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549	Cis P.A. Cis P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	p.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc. nane Conc.	1.5 Meth P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Meth P.A. 0 66.23	33 hane Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 333 hane Conc. 0.00 16.53
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.3'           473.19         7.33           468.38         7.26           2273.6         35.2'           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 T P.A. 12.98 16.13 28.33	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284	Cis P.A. Cis P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc. nane Conc.	1.5 Meth P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Meth P.A. 0 66.23 364.01	33 hane Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 33 hane Conc. 0.00 16.53 90.86
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.3'           473.19         7.33           468.38         7.26           2273.6         35.2'           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85           292.88         4.54	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 T P.A. 12.98 16.13 28.33 28.33 23.83 15.58	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.1549 2.0284 1.1549	Cis P.A. Cis P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc. nane Conc.	1.5 Meth P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Meth P.A. 0 66.23 364.01 609.4 611 95	33 Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 33 aane Conc. 0.00 16.53 90.86 152.11 152.71
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 2 2 2 2 2 3 2 3 2 3 2 3 2 3 2 3 2 3 2 3 2 3 2 3 2 3 2 3 2 3 2 3 2 3 3 2 5 5 4 3 2 5 5 5 4 3 2 5 5 5 5 5 5 5 5 5 5 5 5 5	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.33           473.19         7.33           468.38         7.26           2273.6         35.24           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85           292.88         4.54           306.65         4.75	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 T P.A. 12.98 16.13 28.33 23.83 15.58 15.58	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.1751 1.1145	Cis P.A. Cis P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc. nane Conc.	1.5 Meth P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Meth P.A. 0 66.23 364.01 609.4 611.85 721.98	33 Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 33 aane Conc. 0.00 16.53 90.86 152.11 152.72 180.21
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.3           473.19         7.33           468.38         7.26           2273.6         35.2           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85           292.88         4.54           306.65         4.75           687.74         10.60	5. P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 7 P.A. 12.98 16.13 28.33 23.83 15.58 15.58 15.58 15.21	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.1549 2.0284 1.1155 1.11141 1.089	Cis P.A. Cis P.A.	DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc. nane Conc.	1.5 Meth P.A. 172.42 687.37 455.55 454.84 124.01 276.24 233.83 0 1.5 Meth P.A. 0 66.23 364.01 609.4 611.85 721.98 84.18	33 Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 333 ane Conc. 0.00 16.53 90.86 152.71 180.21 180.21 21.01
RT: ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.3           473.19         7.33           468.38         7.26           2273.6         35.2           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85           292.88         4.54           306.65         4.75           687.74         10.60           2645.8         41.0	5. P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 F.A. 12.98 16.13 28.33 23.83 15.56 15.21 0	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.1549 2.0284 1.1549 2.0284 1.1155 1.11141 1.089 0	Cis P.A. Cis P.A.	DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett	nane Conc. nane Conc.	1.5 Meth P.A. 172.42 687.37 455.55 454.84 124.01 276.24 233.83 0 1.5 Meth P.A. 0 66.23 364.01 609.4 611.85 721.98 84.18 0	33 Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 33 mane Conc. 0.00 16.53 90.86 152.11 152.12 180.21 21.01 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.33           468.38         7.26           2273.6         35.2'           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85           292.88         4.54           306.65         4.75           687.74         10.60           2645.8         41.0	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 5. T P.A. 12.98 16.13 28.33 28.33 28.33 15.56 15.21 0	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.7062 1.1155 1.1141 1.089 0	Cis P.A. Cis P.A.	DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	column P.A. column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett	nane Conc. nane Conc.	1.5 Meth P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Meth P.A. 0 66.23 364.01 609.4 611.85 721.98 84.18 0	33 Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 333 Conc. 0.00 16.53 90.86 152.11 152.72 180.21 21.01 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 8 7	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.3           473.19         7.33           468.38         7.26           2273.6         35.2           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85           292.88         4.54           351.2         5.44           306.65         4.75           687.74         10.61           2645.8         41.0           7.565         PCE	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 5. T P.A. 12.98 16.13 28.33 28.33 15.58 15.56 15.21 0	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.7062 1.1155 1.1141 1.089 0	Cis P.A. Cis P.A.	DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran	EE3 ( sDCE Conc. SA3 ( sDCE Conc. SA2 ( sDCE	column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc.	1.5 Mett P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Mett P.A. 0 66.23 364.01 609.4 611.85 721.98 84.18 0	33 hane Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 333 hane Conc. 0.00 16.53 90.86 152.12 180.21 21.01 0.00 333 hane
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 7 6 5 4 7 6 5 4 7 6 5 4 7 6 5 4 7 6 5 4 7 6 5 6 5 4 7 6 5 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 8 7 8 7 7 6 5 7 8 7 8 7 7 6 5 7 8 7 7 8 8 8 8 8 8 8 8 8 8 8 8 8	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.3           473.19         7.33           468.38         7.26           2273.6         35.2           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85           292.88         4.54           351.2         5.44           306.65         4.75           687.74         10.60           2645.8         41.0           7.565         PCE           P.A.         Conc	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 5. T P.A. 7 P.A.	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.7062 1.1155 1.1141 1.089 0 485 CE CE CCE Conc.	Cis P.A. Cis P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc. SA2 ( sDCE Conc.	column P.A.	VC Conc. VC Conc. VC	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc.	1.5 Mett P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Mett P.A. 0 66.23 364.01 609.4 611.85 721.98 84.18 0	33 hane Conc. 43.04 171.57 113.71 113.53 30.95 73.34 0.00 333 hane Conc. 0.00 16.53 90.86 152.11 152.72 180.21 21.01 0.00 333 hane Conc.
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 7 6 5 4 7 6 5 7 6 5 4 7 6 5 4 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 7 6 5 7 7 7 7 6 5 7 7 7 7 7 7 7 7 7 7 7 7 7	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.3           473.19         7.33           468.38         7.26           2273.6         35.2           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85           292.88         4.54           351.2         5.44           306.65         4.75           687.74         10.60           2645.8         410           7.565         PCE           P.A.         Conc           265.74         10.61           265.97         2.11	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 5. T P.A. 12.98 16.13 28.33 28.33 15.58 15.56 15.21 0 T P.A. 14.93 23.82 5.5 15.46 0 T P.A. 14.93 23.82 15.5 15.46 0 T P.A. 14.93 23.82 15.5 15.46 0 T P.A. 15.5 15.56 15.57 15.57 15.58 15.58 15.58 15.58 15.58 15.58 15.58 15.58 15.58 15.58 15.58 15.58 15.58 15.58 15.58 15.58 15.58 15.57 15.78 15.58 15	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.7062 1.1155 1.1141 1.089 0 485 CE Conc. 1.0475 0 0	Cisi P.A. Cisi P.A. Cisi P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran P.A.	EE3 of sDCE Conc. SA3 of sDCE Conc. SA2 of sDCE Conc.	column P.A.	VC Conc. VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Mett P.A. 0 66.23 364.01 609.4 611.85 721.98 84.18 0 0	33 hane Conc. 43.04 171.57 113.71 113.53 30.95 73.34 0.00 333 hane Conc. 0.00 16.53 90.86 152.11 152.72 180.21 21.01 0.00 333 hane Conc. 3.34 152.72 180.21 21.01 0.00 333 hane Conc. 3.34 21.01 0.00 333 hane Conc. 3.34 21.01 0.00 333 hane Conc. 3.34 21.01 0.05 152.72 180.21 21.01 0.00 152.72 180.21 21.01 0.00 152.72 180.21 21.01 0.00 152.72 180.21 21.01 0.00 152.72 180.21 21.01 0.00 152.72 180.21 21.01 0.00 152.72 180.21 21.01 0.00 152.72 180.21 21.01 0.00 152.72 180.21 21.01 0.00 152.72 180.21 21.01 0.00 183.75 152.72 180.21 21.01 0.00 183.75 152.72 180.21 21.01 0.00 183.75 172.72 180.21 21.01 0.00 183.75 172.72 180.21 21.01 0.00 183.75 172.72 180.21 21.01 0.00 183.75 172.72 180.21 21.01 0.00 183.75 172.75 180.21 21.01 0.00 183.75 172.75 180.21 21.01 0.00 183.75 1
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 7 6 5 4 7 6 5 7 7 6 5 7 6 5 7 7 6 5 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 6 7 7 7 7 7 7 7 7 7 7 7 7 7	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.3           473.19         7.33           468.38         7.26           2273.6         35.2           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85           292.88         4.51           306.65         4.75           687.74         10.61           2645.8         41.0           7.565         PCE           P.A.         Conc           306.55         4.75           687.74         10.61           2645.8         41.0           7.565         PCE           P.A.         Conc           135.97         2.11           275.71         4.27           276.35         4.33	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 5. T P.A. 12.98 16.13 28.33 28.33 15.58 15.56 15.21 0 T P.A. 14.63 8.79 19.22	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.7062 1.1155 1.1141 1.089 0 485 CE Conc. 1.0475 0.6294 1.3762	Cis P.A. Cis P.A. Cis	DCE Conc. DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran P.A.	EE3 of sDCE Conc. SA3 of sDCE Conc. SA2 of sDCE Conc.	column P.A. Column P.A.	VC Conc. VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	hane Conc.	1.5 Mett P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Mett P.A. 0 66.23 364.01 609.4 611.85 721.98 84.18 0 1.5 Mett P.A. 12.99 298.62 329.09	33 Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 333 mane Conc. 0.00 16.53 90.86 152.12 180.21 21.01 0.00 333 mane Conc. 3.24 74.54 82.14
RT: ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 7 6 5 4 7 6 5 4 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 7 6 5 4 7 7 8 7 7 8 7 7 8 7 7 8 7 7 8 7 7 8 7 7 8 7 7 8 7 7 8 7 7 8 7 7 8 7 7 8 7 7 8 7 7 8 7 7 7 8 7 7 8 7 7 8 7 7 8 7 7 8 8 7 8 8 7 8 8 8 8 8 8 8 8 8 8 8 8 8	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.3           473.19         7.33           468.38         7.26           2273.6         35.2           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85           292.88         4.54           306.65         4.75           687.74         10.6           2645.8         41.0           7.565         PCE           P.A.         Conc           306.65         4.75           687.74         10.6           2645.8         41.0           7.565         PCE           P.A.         Conc           135.97         2.11           275.71         4.27           279.35         4.33           374.45         5.80	5. 7 P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 7 7 7 7 7 7 8 15.58 15.56 15.21 0 7 7 7 7 7 7 7 7 7 7 7 7 7	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.7062 1.1155 1.1141 1.089 0 485 CE Conc. 1.0475 0.6294 1.3762 1.3475	Cis P.A. Cis P.A. Cis	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran P.A.	EE3 of sDCE Conc. SA3 of sDCE Conc. SA2 of sDCE Conc.	column P.A. column P.A.	VC Conc. VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Mett P.A. 0 66.23 364.01 609.4 611.85 721.98 84.18 0 1.5 Mett P.A. 12.99 298.62 329.09 558.45	33 Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 333 mane Conc. 0.00 16.53 90.86 152.12 180.21 21.01 0.00 333 mane Conc. 3.24 74.54 82.14 139.39
RT: ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.3           473.19         7.33           468.38         7.26           2273.6         35.2           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85           292.88         4.54           306.65         4.75           687.74         10.6           2645.8         41.0           7.565         PCE           P.A.         Conc           306.65         4.75           687.74         10.6           2645.8         41.0           7.565         PCE           P.A.         Conc           135.97         2.11           275.71         4.27           279.35         4.33           374.45         5.80           501.28         7.77	5. T P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 5. T P.A. 12.98 16.13 28.33 28.33 15.58 15.56 15.21 0 T P.A. 14.63 8.79 19.22 18.82 17.96 14.63 15.79 19.22 17.96 14.63 15.79 19.22 17.96 14.63 15.79 19.22 17.96 14.63 15.79 19.22 17.96 14.63 15.79 19.22 17.96 14.63 15.79 19.22 17.96 14.63 15.79 19.22 19.25 19.25 19.25 19.25 19.25 19.25 19.25 19.25 19.26 19.27	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.7062 1.1155 1.1141 1.089 0 485 CE Conc. 1.0475 0.6294 1.3762 1.3475 1.3475	Cisl P.A. Cisl P.A. Cisl P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran P.A.	EE3 of sDCE Conc. SA3 of sDCE Conc. SA2 of sDCE Conc.	column P.A.	VC Conc. VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	hane Conc.	1.5 Mett P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Mett P.A. 0 66.23 364.01 609.4 611.85 721.98 84.18 0 1.5 Mett P.A. 12.99 298.62 329.09 258.45 744.69	33 Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 333 mane Conc. 0.00 16.53 90.86 152.12 180.21 21.01 0.00 333 mane Conc. 3.24 74.54 82.14 139.39 185.87 186.87 186.87 189.39 185.87 186.87 186.87 189.39 185.87 186.87 189.39 185.87 186.87 189.39 185.87 185.87 186.87 186.87 189.39 185.87 185.87 186.87 186.87 185.87
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc           14.96         0.23           16.33         0.25           111.5         1.73           282.07         4.37           733.6         11.3           473.19         7.33           468.38         7.26           2273.6         35.2           7.565         PCE           P.A.         Conc           80.68         1.25           81.66         1.27           312.62         4.85           202.88         4.54           306.65         4.75           687.74         10.60           2645.8         41.0           7.565         PCE           P.A.         Conc           306.65         4.75           687.74         10.60           2645.8         41.0           7.565         PCE           P.A.         Conc           135.97         2.11           275.71         4.27           374.45         5.80           501.28         7.77           488.87         7.58           1074.2 <td>5. 7 P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 7 7 7 7 7 7 8 15.58 15.56 15.21 0 7 7 7 7 7 7 7 7 7 7 7 7 7</td> <td>485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.7062 1.1155 1.1141 1.089 0 485 CE Conc. 1.0475 0.6294 1.3475 1.3475 1.3475</td> <td>Cis P.A. Cis P.A. Cis</td> <td>DCE Conc.</td> <td>Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran P.A.</td> <td>EE3 of sDCE Conc. SA3 of sDCE Conc. SA2 of sDCE Conc.</td> <td>column P.A.</td> <td>VC Conc. VC Conc. VC Conc.</td> <td>Eth P.A. Eth P.A.</td> <td>ylene Conc. ylene Conc.</td> <td>Ett P.A. Ett P.A.</td> <td>hane Conc.</td> <td>1.5 Mett P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Mett P.A. 0 66.23 364.01 609.4 611.85 721.98 84.18 0 1.5 Mett P.A. 12.99 298.62 329.09 558.45 744.69 660.94 35 31</td> <td>33 Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 333 mane Conc. 0.00 16.53 90.86 152.12 180.21 21.01 0.00 333 mane Conc. 3.24 74.54 82.14 139.39 185.87 164.97 8.81</td>	5. 7 P.A. 14.93 23.42 34.36 26.9 60.44 15.5 15.46 0 7 7 7 7 7 7 8 15.58 15.56 15.21 0 7 7 7 7 7 7 7 7 7 7 7 7 7	485 CE Conc. 1.069 1.6769 2.4602 1.926 4.3275 1.1098 1.1098 1.1069 0 485 CE Conc. 0.9294 1.1549 2.0284 1.7062 1.1155 1.1141 1.089 0 485 CE Conc. 1.0475 0.6294 1.3475 1.3475 1.3475	Cis P.A. Cis P.A. Cis	DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran P.A.	EE3 of sDCE Conc. SA3 of sDCE Conc. SA2 of sDCE Conc.	column P.A.	VC Conc. VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	hane Conc.	1.5 Mett P.A. 172.42 687.37 455.55 454.84 124.01 276.24 293.83 0 1.5 Mett P.A. 0 66.23 364.01 609.4 611.85 721.98 84.18 0 1.5 Mett P.A. 12.99 298.62 329.09 558.45 744.69 660.94 35 31	33 Conc. 43.04 171.57 113.71 113.53 30.95 68.95 73.34 0.00 333 mane Conc. 0.00 16.53 90.86 152.12 180.21 21.01 0.00 333 mane Conc. 3.24 74.54 82.14 139.39 185.87 164.97 8.81



				Col	7 EE2 C	olumn us	sed 10/8	stds						
RT:	7.565	5.	485	2.588	3.	171		0	Edu	. da a a	E di		1.5	533
norte	PCE PA Conc		Conc	CISDCE P.A. Conc	I ran	SDCE		Conc	Eth	ylene	Etr	Conc	Meth P A	nane
7	43.33 0.67	10.59	0.7582	F.A. Conc.	г.д.	COILC.	F.A.	CONC.	г.А.	CONC.	г.А.	COILC.	68.4	17.07
6	35.96 0.56	20.97	1.5015										276.36	68.98
5	90.27 1.40	37.32	2.6721										763.98	190.69
4	130.5 2.02	33.5	2.3986										525.39	131.14
3	202.56 3.14	85.75	6.1397										339.26	84.68
2	1467 3 22 74	17 67	1 2652										442.71	25 21
IFF	3307.1 51.20	0	0										0	0.00
-	-		-		Col 8	Control	2 Colum	n						
RT:	7.565	5.	485										1.5	533
	PCE	Т	CE	CisDCE	Tran	sDCE	١	/C	Eth	ylene	Eth	nane	Meth	nane
ports	P.A. Conc	P.A.	Conc.	P.A. Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
7	24.52 0.38	29.09	2.0828										764.61	190.85
5	42.95 0.67	41.25	2.9535										548.48 433.96	136.90
4	348.94 5.41	48.46	3.4697										321.36	80.21
3	493.79 7.65	27.63	1.9783										136.87	34.16
2	671.36 10.4	31.94	2.2869										37.52	9.36
1	1381.8 21.42	0	0										0	0.00
IFF	1613.2 25.0	0	0										0	0.00
					Col 9	SA1 d	column							
RI:	7.565 PCF	5. T	485 CE	CisDCE	Tran	sDCF	、 、	/C	Fth	vlene	Fth	ane	1.5 Meth	o33 nane
ports	P.A. Conc	P.A.	Conc.	P.A. Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
7	29.9 0.46	13.22	0.9466										18.39	4.59
6	99.2 1.54	23.84	1.7069										134.54	33.58
5	382.34 5.93	45.49	3.2571										651.64	162.65
4	587 43 9 11	38.0	2.7638										802.81 744 53	200.38
2	648.78 10.06	25.37	1.8165										313.51	78.25
1	1583.5 24.54	0	0										0	0.00
IFF	1561.9 24.2	0	0										0	0.00
													-	
RT	7 565	5	485		Col 10	EE1	column						1.5	33
RT:	7.565 PCE	5. T	485 CE	CisDCE	Col 10 Tran	EE1 sDCE	column \	/C	Eth	ylene	Eth	nane	1.5 Meth	i33 nane
RT: ports	7.565 PCE P.A. Conc	5. T P.A.	485 CE Conc.	CisDCE P.A. Conc.	Col 10 Tran P.A.	EE1 sDCE Conc.	column \ P.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A.	533 nane Conc.
RT: ports 7	7.565 PCE P.A. Conc	5. T P.A.	485 CE Conc.	CisDCE P.A. Conc.	Col 10 Tran P.A.	EE1 sDCE Conc.	column \ P.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A.	i33 nane Conc.
RT: ports 7 6 5	7.565 PCE P.A. Conc 102.75 1.59	5. T P.A. 26.72	485 CE Conc. 1.9132	CisDCE P.A. Conc.	Col 10 Tran P.A.	EE1 sDCE Conc.	column \ P.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 611.78	533 nane Conc. 152.70
RT: ports 7 6 5 4	7.565 PCE P.A. Conc 102.75 1.59 360.3 5.58	5. T P.A. 26.72 44.46	485 CE Conc. 1.9132 3.1833	CisDCE P.A. Conc.	Col 10 Tran P.A.	EE1 sDCE Conc.	column \ P.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 611.78 443.3	533 nane Conc. 152.70 110.65
RT: ports 7 6 5 4 3	7.565 PCE P.A. Conc 102.75 1.59 360.3 5.58 523.51 8.11	5. T P.A. 26.72 44.46 39.78	485 CE Conc. 1.9132 3.1833 2.8482	CisDCE P.A. Conc.	Col 10 Tran P.A.	EE1 sDCE Conc.	column ۲.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 611.78 443.3 350.42	533 nane Conc. 152.70 110.65 87.46
RT: ports 7 6 5 4 3 2	7.565 PCE P.A. Conc 102.75 1.59 360.3 5.58 523.51 8.11 530.55 8.22	5. T P.A. 26.72 44.46 39.78 82.16	485 CE Conc. 1.9132 3.1833 2.8482 5.8827	CisDCE P.A. Conc.	Col 10 Tran P.A.	EE1 sDCE Conc.	column ۲.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 611.78 443.3 350.42 253.75	533 nane Conc. 152.70 110.65 87.46 63.34
RT: ports 7 6 5 4 3 2 1	7.565 PCE P.A. Conc 102.75 1.59 360.3 5.58 523.51 8.11 530.55 8.22 2104.3 32.66	5. T P.A. 26.72 44.46 39.78 82.16 0	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0	CisDCE P.A. Conc.	Col 10 Tran P.A.	EE1 sDCE Conc.	Column	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 611.78 443.3 350.42 253.75 1.4	533 hane Conc. 152.70 110.65 87.46 63.34 0.35 0.20
RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.63           3154.1         48.89	5. T P.A. 26.72 44.46 39.78 82.16 0 0	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0	CisDCE P.A. Conc.	Col 10 Tran P.A.	EE1 sDCE Conc.	Column	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 611.78 443.3 350.42 253.75 1.4 0	33 hane Conc. 152.70 110.65 87.46 63.34 0.35 0.00
RT: 7 6 5 4 3 2 1 IFF	7.565 PCE P.A. Conc 102.75 1.59 360.3 5.58 523.51 8.11 530.55 8.22 2104.3 32.67 3154.1 48.89	5. T P.A. 26.72 44.46 39.78 82.16 0 0	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0	CisDCE P.A. Conc.	Col 10 Tran P.A.	EE1 sDCE Conc.	Column	/C Conc. n	Eth P.A.	ylene Conc.	Etr P.A.	nane Conc.	1.5 Meth P.A. 611.78 443.3 350.42 253.75 1.4 0	333 hane Conc. 152.70 110.65 87.46 63.34 0.35 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT:	7.565           PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.63           3154.1         48.89           7.565         PCE	5. T P.A. 26.72 44.46 39.78 82.16 0 0	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE	CisDCE P.A. Conc. CisDCE	Col 10 Tran P.A. Col 11 Tran	EE1 sDCE Conc.	Column	/C Conc. n /C	Eth P.A. Eth	ylene Conc.	Eth P.A. Eth	nane Conc.	1.5 Meth P.A. 611.78 443.3 350.42 253.75 1.4 0	33 hane Conc. 152.70 110.65 87.46 63.34 0.35 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: ports	7.565           PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.63           3154.1         48.89           7.565         PCE           P.A.         Conc	5. T P.A. 26.72 44.46 39.78 82.16 0 0 5. T P.A.	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Ce Conc.	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	EE1 <u>sDCE</u> Conc. <u>Control</u> <u>sDCE</u> Conc.	Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A.	33 hane Conc. 152.70 110.65 87.46 63.34 0.35 0.00 333 hane Conc.
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7	7.565           PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.62           3154.1         48.89           7.565         PCE           P.A.         Conc           57.21         0.89	5. T P.A. 26.72 44.46 39.78 82.16 0 0 5. T P.A. 42.29	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE CE Conc. 3.028	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	EE1 <u>sDCE</u> Conc. <u>Control</u> <u>sDCE</u> Conc.	Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41	333 hane Conc. 152.70 110.65 87.46 63.34 0.35 0.00 333 hane Conc. 187.55
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6	7.565           PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.62           3154.1         48.89           7.565         PCE           P.A.         Conc           57.21         0.89           58.23         0.90	5. T P.A. 26.72 44.46 39.78 82.16 0 0 5. T P.A. 42.29 32.02 0 10	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.028 2.2926	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	EE1 <u>sDCE</u> Conc. <u>Control</u> <u>sDCE</u> Conc.	Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9	333 Torrest Conc. 152.70 110.65 87.46 63.34 0.35 0.00 333 Torrest Conc. 187.55 212.63 400 00
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.63           3154.1         48.89           7.565         PCE           P.A.         Conc           57.21         0.89           58.23         0.90           196.71         3.05	5. T P.A. 26.72 44.46 39.78 82.16 0 0 5. T P.A. 42.29 32.02 66.16	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.028 2.2926 4.7371 2.716	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	EE1 <u>sDCE</u> Conc. <u>Control</u> <u>sDCE</u> Conc.	Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72	333 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 333 Dane Conc. 187.55 212.63 120.96 120.96
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.62           3154.1         48.89           7.565         PCE           P.A.         Conc           57.21         0.89           58.23         0.90           196.71         3.05           247.09         3.83           459.79         7.13	5. T P.A. 26.72 44.46 39.78 82.16 0 0 5. T P.A. 42.29 32.02 66.16 51.9 43.26	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.028 2.2926 4.7371 3.7974	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	EE1 <u>sDCE</u> Conc. <u>Control</u> <u>sDCE</u> Conc.	Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72 528.52	333 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 333 Conc. 187.55 212.63 120.96 179.89 131.92
RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.63           3154.1         48.89           7.565           PCE         P.A.           Conc         57.21         0.89           58.23         0.90         196.71         3.05           247.09         3.83         459.79         7.13           508.73         7.89         368.23         7.89	5. T P.A. 26.72 44.46 39.78 82.16 0 0 0 T P.A. 42.29 32.02 66.16 51.9 43.26 38.54	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.028 2.2926 4.7371 3.716 3.0974 2.7595	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	EE1 <u>sDCE</u> Conc. <u>Control</u> <u>sDCE</u> Conc.	Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72 528.52 234.17	33 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 33 ane Conc. 187.55 212.63 120.96 179.89 131.92 58.45
RT: ports 7 6 5 4 3 2 1 IFF 7 7 6 5 4 3 2 1	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.63           3154.1         48.89           7.565           PCE         P.A.           Conc         57.21         0.89           58.23         0.90         196.71         3.05           247.09         3.83         459.79         7.13           508.73         7.89         1416.8         21.96	5. T P.A. 26.72 44.46 39.78 82.16 0 0 0 5. T P.A. 7 82.29 32.02 66.16 51.9 43.26 38.54 0	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.028 2.2926 4.7371 3.716 3.0974 2.7595 0	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	EE1 <u>sDCE</u> Conc. <u>Control</u> <u>sDCE</u> Conc.	Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72 528.52 234.17 0	33 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 33 Conc. 187.55 212.63 120.96 179.89 131.92 58.45 0.00
RT: ports 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.62           3154.1         48.89           7.565           PCE         P.A.           Conc         57.21         0.89           58.23         0.90         196.71         3.05           247.09         3.83         459.79         7.13           508.73         7.89         1416.8         21.96           1899.6         29.44         1459.49         29.44	5. T P.A. 26.72 44.46 39.78 82.16 0 0 0 T P.A. 42.29 32.02 66.16 51.9 43.26 38.54 0 0 0	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.028 2.2926 4.7371 3.716 3.0974 2.7595 0 0	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	EE1 sDCE Conc. Control sDCE Conc.	Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72 528.52 234.17 0 0	33 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 33 ane Conc. 187.55 212.63 120.96 179.89 131.92 58.45 0.00 0.00
RT: ports 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.62           3154.1         48.89           7.565         PCE           P.A.         Conc           57.21         0.89           58.23         0.90           196.71         3.05           247.09         3.83           459.79         7.13           508.73         7.89           1416.8         21.96           1899.6         29.44	5. T P.A. 26.72 44.46 39.78 82.16 0 0 5. T P.A. 42.29 32.02 66.16 51.9 43.26 38.54 0 0	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.028 2.2926 4.7371 3.716 3.0974 2.7595 0 0	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 11	EE1 sDCE Conc. Control sDCE Conc. SDCE Conc.	Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72 528.52 234.17 0 0	33 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 33 Conc. 187.55 212.63 120.96 179.89 131.92 58.45 0.00 0.00
RT: ports 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.63           3154.1         48.89           7.565         PCE           P.A.         Conc           57.21         0.89           58.23         0.90           196.71         3.05           247.09         3.83           459.79         7.13           508.73         7.89           1416.8         21.94           7.565         PCE	5. T P.A. 26.72 44.46 39.78 82.16 0 0 5. T P.A. 42.29 32.02 66.16 51.9 43.26 38.54 0 0	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.028 2.2926 4.7371 3.716 3.0974 2.7595 0 0 0	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 12	EE1 sDCE Conc. Control sDCE Conc. SDCE SDCE	Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72 528.52 234.17 0 0	33 hane Conc. 152.70 110.65 87.46 63.34 0.35 0.00 333 Conc. 187.55 212.63 120.96 179.89 131.92 58.45 0.00 0.00 333
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 7 7 8 8 8 7 8 8 8 8 8 8 8 8 8 8 8 8 8	7.565           PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.63           3154.1         48.89           7.565         PCE           P.A.         Conc           57.21         0.89           58.23         0.90           196.71         3.05           247.09         3.83           459.79         7.13           508.73         7.89           1416.8         21.94           7.565         PCE           P.A.         Conc	5. T P.A. 26.72 44.46 39.78 82.16 0 0 5. T P.A. 42.29 32.02 66.16 51.9 43.26 38.54 0 0 5. T P.A. 7 T P.A.	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.028 2.2926 4.7371 3.716 3.0974 2.7595 0 0 485 CE CE Conc.	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 12 Tran	EE1 sDCE Conc. Control sDCE Conc. SDCE Conc. SA0 sDCE Conc.	Column P.A. 1 colum P.A. column	/C Conc. n /C Conc. /C	Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72 528.52 234.17 0 0	333 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 333 bane Conc. 187.55 212.63 120.96 179.89 131.92 58.45 0.00 0.00 333 bane Conc. 187.55 212.63 120.96 179.89 131.92 58.45 0.00 0.00 Conc. 133.92 58.45 0.00 0.00 Conc. 133.92 58.45 0.00 Conc. 133.92 58.45 0.00 Conc. 133.92 58.45 0.00 Conc. 133.92 58.45 0.00 Conc. 133.92 58.45 0.00 Conc. 133.92 58.45 0.00 Conc. 133.92 58.45 0.00 Conc. 133.92 58.45 0.00 Conc.
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 7 7 7 7 7 7 7 7 7 7 7	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.63           3154.1         48.89           7.565         PCE           P.A.         Conc           57.21         0.89           58.23         0.90           196.71         3.05           247.09         3.83           459.79         7.13           508.73         7.89           1416.8         21.94           7.565         PCE           P.A.         Conc           63.42         0.98	5. T P.A. 26.72 44.46 39.78 82.16 0 0 5. T P.A. 42.29 32.02 66.16 51.9 43.26 38.54 0 0 5. T P.A. 69.41	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.0974 2.7595 0 0 485 CE CE Conc. 4.7371 3.716 3.0974 2.7595 0 0 0	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 12 Tran P.A.	EE1 sDCE Conc. Control sDCE Conc. SA0 sDCE Conc.	Column P.A. 1 colum P.A. column	/C Conc. n /C Conc. /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A. Ett	hane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72 528.52 234.17 0 0	333 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 333 Conc. 187.55 212.63 120.96 179.89 131.92 58.45 0.00 0.00 333 120.96 179.89 131.92 58.45 0.00 0.00 333 120.96 179.89 131.92 58.45 0.00 0.00 58.45 0.00 0.00 58.45 0.00 0.00 58.45 0.00 0.00 58.45 0.00 0.00 58.45 0.00 0.00 58.45 0.00 0.00 58.45 0.00 0.00 58.45 0.00 0.00 58.45 0.00 0.00 58.45 0.00 0.00 58.45 0.00 0.00 58.45 0.00 0
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 7 6 5 4 7 6 5 4 7 6 5 4 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 7 6 5 7 6 7 6 7 6 7 6 7 6 7 6 7 7 6 7 7 6 7 7 6 7 7 6 7 7 6 7 7 6 7 7 6 7 7 7 6 7 7 7 6 7 7 7 7 6 7 7 7 7 7 7 7 7 7 7 7 7 7	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.63           3154.1         48.89           7.565         PCE           P.A.         Conc           57.21         0.89           58.23         0.90           196.71         3.05           247.09         3.83           459.79         7.13           508.73         7.89           1416.8         21.96           1899.6         29.44           F.A.           Conc         63.42           9.83         41.68	5. T P.A. 26.72 44.46 39.78 82.16 0 0 5. T P.A. 42.29 32.02 66.16 51.9 43.26 38.54 0 0 0 T T P.A. 5. T P.A. 5. T P.A.	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.0974 2.7595 0 0 485 CE CE Conc. 4.7371 3.716 3.0974 2.7595 0 0 0	CisDCE P.A. Conc. CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 12 Tran P.A.	EE1 sDCE Conc. Control sDCE Conc. SDCE SA0 sDCE Conc.	column P.A. 1 colum P.A. column N P.A.	/C Conc. n /C Conc. /C Conc.	Eth P.A. Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A. Ett P.A.	nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72 528.52 234.17 0 0 0	33 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 33 Conc. 187.55 212.63 120.96 179.89 131.92 58.45 0.00 0.00 33 258.45 0.00 0.00 33 258.45 0.00 0.00 33 258.45 0.00 0.00 33 258.45 0.00 0.00 33 258.45 0.00 0.00 33 258.45 0.00 0.00 33 258.45 0.00 0.00 33 258.45 0.00 0.00 33 258.45 0.00 0.00 33 258.45 0.00 0.00 33 258.45 0.00 0.0
RT: ports 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 IFF 7 6 5 7 6 5 7 6 5 7 6 5	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.62           3154.1         48.89           7.565         PCE           P.A.         Conc           57.21         0.89           58.23         0.90           196.71         3.05           247.09         3.83           459.79         7.13           508.73         7.89           1416.8         21.96           1899.6         29.44           F.A.           Conc         63.42           9.63.42         0.98           41.68         0.65           65.95         1.02           0.92 57         1.93	5. T P.A. 26.72 44.46 39.78 82.16 0 0 0 5. T P.A. 42.29 32.02 66.16 51.9 43.26 38.54 0 0 0 T P.A. 69.41 28.26 34.52	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.0974 2.7595 0 0 485 CE Conc. 4.7371 3.716 3.0974 2.7595 0 0 0	CisDCE P.A. Conc. CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 12 Tran P.A.	EE1 SDCE Conc. Control SDCE Conc. Conc. SDCE Conc. Conc. Conc.	column P.A. 1 colum P.A. column	/C Conc. n /C Conc. /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72 528.52 234.17 0 0 0	333 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 333 Conc. 187.55 212.63 120.96 179.89 131.92 58.45 0.00 0.00 333 120.96 179.89 131.92 58.45 0.00 0.00 333 120.96 133.92 58.45 0.00 0.00 333 120.96 133.92 58.45 0.00 0.00 333 120.96 133.92 131.92 58.45 0.00 0.00 133.92 134.92 134.92 135.92 1
RT: ports 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 2 1 S 7 6 5 4 3 2 2 1 S 7 6 5 4 3 2 7 8 7 8 7 8 7 8 7 8 7 8 7 8 7 8 7 8 7	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.62           3154.1         48.89           7.565         PCE           P.A.         Conc           57.21         0.89           58.23         0.90           196.71         3.05           247.09         3.83           459.79         7.13           508.73         7.89           1416.8         21.96           1899.6         29.44           F.A.           Conc         63.42           9.83         41.68           0.65         5.06           PCE         P.A.           84.1.68         0.65           65.95         1.02           326.55         5.06           384.34         5.96	5. T P.A. 26.72 44.46 39.78 82.16 0 0 5. T P.A. 42.29 32.02 66.16 51.9 43.26 38.54 0 0 0 T P.A. 69.41 28.26 34.52 37.33 25.17	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 0 485 CE Conc. 3.0974 2.7595 0 0 0 485 CE Conc. 4.7371 3.716 3.0974 2.7595 0 0 0	CisDCE P.A. Conc. CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 12 Tran P.A.	EE1 sDCE Conc. Control sDCE Conc. SDCE Conc. SA0 sDCE Conc.	column P.A. 1 colum P.A. column	/C Conc. n /C Conc. /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72 528.52 234.17 0 0 0 1.5 Mett P.A. 31.77 31.63 175.21 278.98 292.92	33 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 33 Conc. 187.55 212.63 120.96 179.89 131.92 58.45 0.00 0.00 33 120.96 179.89 131.92 58.45 0.00 0.00 33 120.96 133.92 58.45 0.00 0.00 33 120.96 133.92 131.92 58.45 0.00 0.00 33 120.96 131.92 58.45 0.00 0.00 33 120.96 133.92 131.92 58.45 0.00 0.00 0.00 133.92 135.92 15
RT: ports 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 2 1 IFF 7 6 5 4 3 2 2 1 S 7 6 5 4 3 2 2 1 S 7 6 5 5 4 3 2 7 8 7 8 7 8 7 8 7 8 7 8 7 8 7 8 7 8 7	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.62           3154.1         48.83           7.565         PCE           P.A.         Conc           57.21         0.89           58.23         0.90           196.71         3.05           247.09         3.83           459.79         7.13           508.73         7.89           1416.8         21.96           1899.6         29.44           7.565           PCE         P.A.           Conc         63.42         0.98           41.68         0.65         65.95         1.02           326.55         5.06         384.34         5.96           527.68         8.18         18	5. T P.A. 26.72 44.46 39.78 82.16 0 0 0 T P.A. 42.29 32.02 66.16 51.9 43.26 38.54 0 0 0 T P.A. 42.29 32.02 66.16 51.9 43.26 38.54 1.2 8 43.26 38.54 1.2 8 2 5.7 7 8 7 8 7 8 7 8 7 8 7 8 7 8 7 8 7 8 7	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 0 485 CE Conc. 3.028 2.2926 4.7371 3.716 3.0974 2.7595 0 0 0 485 CE Conc. 4.7371 3.716 3.0974 2.7595 0 0 0	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 12 Tran P.A.	EE1 SDCE Conc. Control SDCE Conc. SA0 SDCE Conc.	column 1 colum P.A. P.A. column V P.A.	/C Conc. n /C Conc. /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	hane Conc. hane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 Mett P.A. 751.41 851.9 484.6 720.72 528.52 234.17 0 0 0 1.5 Mett P.A. 31.77 131.63 175.21 278.98 292.92 161.25	333 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 333 bane Conc. 187.55 212.63 120.96 179.89 131.92 58.45 0.00 0.00 333 bane Conc. 7.93 32.85 43.73 69.63 73.11 40.25
RT: ports 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 IFF 7 6 5 4 3 2 1 1 IFF 7 6 5 4 3 7 1 1 IFF 7 6 5 7 1 1 IFF 7 6 5 7 1 1 IFF 7 6 5 7 1 1 IFF 7 6 5 7 1 1 IFF 7 6 5 7 1 1 IFF 7 6 5 7 1 1 IFF 7 6 5 7 1 1 IFF 7 6 5 7 1 1 IFF 7 1 1 IFF 7 1 1 IFF 7 1 1 IFF 7 1 1 I I I I I I I I I I I I I I I I I	7.565         PCE           P.A.         Conc           102.75         1.59           360.3         5.58           523.51         8.11           530.55         8.22           2104.3         32.62           3154.1         48.83           7.565         PCE           P.A.         Conc           57.21         0.89           58.23         0.90           196.71         3.05           247.09         3.83           459.79         7.13           508.73         7.89           1416.8         21.90           1899.6         29.44           7.565         PCE           P.A.         Conc           63.42         0.98           41.68         0.65           65.95         1.02           326.55         5.06           384.34         5.96           527.68         8.18           1187.7         18.4'	5. T P.A. 26.72 44.46 39.78 82.16 0 0 0 F.A. 42.29 32.02 66.16 51.9 43.26 38.54 0 0 0 7 T P.A. 42.29 32.02 66.16 51.9 43.26 38.54 0 0 0 7 T P.A. 12 5. T P.A. 12.25 12 5. T P.A. 12.25 12.02 12.02 12.02 12.02 13.55 1.9 13.55 1.9 14.26 10 10 10 10 10 10 10 10 10 10 10 10 10	485 CE Conc. 1.9132 3.1833 2.8482 5.8827 0 0 485 CE Conc. 3.028 2.2926 4.7371 3.716 3.074 2.7595 0 0 0 485 CE Conc. 4.9698 2.0234 2.4716 2.6728 1.8022 1.654 0.9702	CisDCE P.A. Conc. CisDCE P.A. Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 12 Tran P.A.	EE1 sDCE Conc. Control sDCE Conc. SA0 sDCE Conc.	column 1 colum P.A. P.A. column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	hane Conc. hane Conc.	1.5 Mett P.A. 611.78 443.3 350.42 253.75 1.4 0 1.5 751.41 851.9 484.6 720.72 528.52 234.17 0 0 0 0 1.5 Mett P.A. 31.77 131.63 175.21 278.98 292.92 161.25 15.87	333 Conc. 152.70 110.65 87.46 63.34 0.35 0.00 333 nane Conc. 187.55 212.63 120.96 179.89 131.92 58.45 0.00 0.00 0.00 333 nane Conc. 7.93 32.85 43.73 69.63 73.11 40.25 3.96



## Appendix F. 4th week data

					Col 1	CC3 Co	olumn <b>Al</b>	Conc l	n (µg /L)						
RT:	7.565	5. T	485	2. Cio	588 DCE	3.1 Trop	171 •DCE	,		E+b	viono	C+k	2000	1.5 Mot	533
ports	P.A. Conc.	P.A.	CE Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
7	115.5 1.79	29.8	2.1337											1.2	0.30
6	88.1 1.37	74.3	5.3199											202.2	50.47
5	754.6 11.70	99.6 80.9	7.1314											521.9 549.5	130.27
3	921.7 14.29	85.3	6.1075											330.9	82.59
2	1438.8 22.30	75.2	5.3843											327.3	81.69
1	1829 28.35	23.9	1.7112											36.3	9.06
	2400.1 30.44	0	- 0				-	L						0	0.00
RT.	7 565	5	485	2	588	Col 2	CC2 (	Jolumn		ſ				1 4	33
	PCE	T	CE	Cis	DCE	Tran	sDCE	١	VC	Eth	ylene	Eth	nane	Met	hane
ports	P.A. Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
6	62.5 0.97 200 7 3 11	23.3 95.3	1.6683											20.9	5.22 65.17
5	419.5 6.50	93.7	6.7089											480.4	119.91
4	714 11.07	83.8	6.0001											381.2	95.15
3	991.1 15.36	86.7	6.2077											356.3	88.93
2	1634.4 25.33	43.7	3.1289											236.2	9.48
IFF	2609.2 40.44	0	0											0	0.00
						Col 3	Control	3 column	1						
RT:	7.565	5.	485	2.	588	3.1	171							1.5	533
porte	PCE	Т	CE	Cis	DCE	Tran	SDCE		VC Cono	Eth	ylene	Eth	nane	Met	hane
7	861.2 13.35	64	4.5824	г. <del>д</del> .	Conc.	Г. <del>Л</del> .	Conc.	г. <del>л</del> .	Conc.	Γ.Λ.	Conc.	F .A.	Conc.	543.7	135.71
6	1025 15.89	58.5	4.1886											474.9	118.54
5	1222.3 18.95	100.9	7.2244											409.6	102.24
4	1146.5 17.77	60.2	4.6683											384.2	95.90
2	1360.5 21.09	52.3	3.7447											311.1	77.65
1	2068 32.05	34.5	2.4702											81	20.22
IFF	3158.4 48.96	0	0											0	0.00
						Col 4	EE3 (	olumn							
RT:	7.565	5.	485	2.	588	Col 4 3.1	EE3 (	column						1.5	533
RT:	7.565 PCE	5. T	485 CE	2. Cis	588 DCE	Col 4 3. Tran	EE3 ( 171 sDCE	column \	VC	Eth	ylene	Eth	nane	1.t Meti	533 hane
RT: ports	7.565 PCE P.A. Conc.	5. T P.A.	485 CE Conc.	2. Cis P.A.	588 DCE Conc.	Col 4 3. Tran P.A.	EE3 ( 171 sDCE Conc.	column ۲.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A.	533 hane Conc.
RT: ports 7 6	7.565 PCE P.A. Conc. 209 3.24 232.1 3.60	5. T P.A. 88.1 116.9	485 CE Conc. 6.308 8.37	2. Cis P.A.	588 DCE Conc.	Col 4 3. <sup>-</sup> Tran P.A.	EE3 ( 171 sDCE Conc.	column ۲.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A. 18.9 316.1	533 hane Conc. 4.72 78.90
RT: ports 7 6 5	7.565 PCE P.A. Conc. 209 3.24 232.1 3.60 630.1 9.77	5. T P.A. 88.1 116.9 110.9	485 CE Conc. 6.308 8.37 7.9404	2. Cis P.A.	588 DCE Conc.	Col 4 3. Tran P.A.	EE3 ( 171 sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Met P.A. 18.9 316.1 403.4	533 hane Conc. 4.72 78.90 100.69
RT: ports 7 6 5 4	7.565 PCE P.A. Conc. 209 3.24 232.1 3.60 630.1 9.77 1007.6 15.62	5. T P.A. 88.1 116.9 110.9 95	485 CE Conc. 6.308 8.37 7.9404 6.802	2. Cis P.A.	588 DCE Conc.	Col 4 3. Tran P.A.	EE3 ( 171 sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A. 18.9 316.1 403.4 447.9	533 hane Conc. 4.72 78.90 100.69 111.80
RT: ports 7 6 5 4 3 2	7.565 PCE P.A. Conc. 209 3.24 232.1 3.60 630.1 9.77 1007.6 15.62 1104.9 17.13 976 5 15.14	5. T P.A. 88.1 116.9 110.9 95 60 71.7	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337	2. Cis P.A.	588 DCE Conc.	Col 4 3. Tran P.A.	EE3 ( 171 sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A. 18.9 316.1 403.4 447.9 235.5 479.6	533 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71
RT: ports 7 6 5 4 3 2 1	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60	5. T P.A. 88.1 116.9 110.9 95 60 71.7 30.9	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124	2. Cis P.A.	588 DCE Conc.	Col 4 3. Tran P.A.	EE3 ( 171 sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Etr P.A.	nane Conc.	1.5 Mett P.A. 18.9 316.1 403.4 447.9 235.5 479.6 110	533 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46
RT: 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99	5. T P.A. 88.1 116.9 110.9 95 60 71.7 30.9 0	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0	2. Cis P.A.	588 DCE Conc.	Col 4 3. <sup>.</sup> Tran P.A.	EE3 ( 171 sDCE Conc.	Column	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Metl P.A. 18.9 316.1 403.4 447.9 235.5 479.6 110 0	533 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00
RT: ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99	5. T P.A. 88.1 116.9 95 60 71.7 30.9 0	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0	2. Cis P.A.	588 DCE Conc.	Col 4 3. Tran P.A. Col 5	EE3 ( 171 sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Ett P.A.	nane Conc.	1.5 Metl P.A. 18.9 316.1 403.4 447.9 235.5 479.6 110 0	533 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT:	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99	5. T P.A. 88.1 116.9 95 60 71.7 30.9 0 5.	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE	2. Cis P.A.	588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE	column P.A.	VC Conc.	Eth P.A.	ylene Conc.	Ett P.A.	nane Conc.	1.5 Mett P.A. 18.9 316.1 403.4 447.9 235.5 479.6 110 0	533 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: ports	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565         PCE           P.A.         Conc.	5. T P.A. 88.1 116.9 95 60 71.7 30.9 0 5. T P.A.	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE Conc.	2. Cis P.A. 2. Cis	588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc.	column P.A.	VC Conc.	Eth P.A. Eth	ylene Conc. ylene Conc.	Ett P.A. Ett	nane Conc. nane Conc.	1.5 Metl P.A. 18.9 316.1 403.4 447.9 235.5 479.6 110 0 1.5 Metl P.A.	533 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565         PCE           P.A.         Conc.           209.9         3.25	5. T P.A. 88.1 116.9 110.9 95 60 71.7 30.9 0 5. T P.A. 29.7	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE Conc. 2.1265	2. Cis P.A. 2. Cis P.A.	588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. P.A. Col 5 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc.	p.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Metl P.A. 18.9 316.1 403.4 447.9 235.5 479.6 110 0 1.5 Metl P.A. 0.8	533 hane Conc. 4.72 78.90 110.69 111.80 58.78 119.71 27.46 0.00 533 hane Conc. 0.20
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565         PCE           P.A.         Conc.           209.9         3.25           394.4         6.11	5. T P.A. 88.1 116.9 95 60 71.7 30.9 0 5. T P.A. 29.7 59.6 70.6	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE Ce Conc. 2.1265 4.2674 5.2674	2. Cis P.A. 2. Cis P.A.	588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc.	xolumn P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Meti P.A. 18.9 316.1 403.4 447.9 235.5 170 0 1.5 Meti P.A. 0.8 37.2	333 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00 333 hane Conc. 0.20 9.29
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565         PCE           P.A.         Conc.           209.9         3.25           394.4         6.11           722.4         11.20           121.9         18.78	5. T P.A. 88.1 116.9 95 60 71.7 30.9 0 5. T P.A. 29.7 59.6 76.8 77.4	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE Conc. 2.1265 4.2674 5.4989 5.5418	2. Cis P.A. Cis P.A.	588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc.	column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Meti P.A. 18.9 316.1 403.4 447.9 235.5 110 0 1.5 Meti P.A. 0.8 37.2 195.4 523.4	333 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00 333 hane Conc. 0.20 9.29 48.77 130.64
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565           PCE           P.A.         Conc.           209.9         3.25           394.4         6.11           722.4         11.20           1221.9         18.78           1122.6         17.40	5. T P.A. 88.1 116.9 110.9 95 60 71.7 30.9 0 5. T P.A. 29.7 59.6 76.8 77.4 60.4	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE CE Conc. 2.1265 4.2674 5.5418 4.3246	2. Cis P.A. Cis P.A.	588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc.	column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Meti P.A. 18.9 316.1 403.4 447.9 235.5 110 0 1.5 Meti P.A. 0.8 37.2 195.4 523.4 730.9	333 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00 333 Conc. 0.20 9.29 48.77 130.64 182.43
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565           PCE           P.A.         Conc.           209.9         3.25           394.4         6.11           722.4         11.20           121.1.9         18.78           1122.6         17.40           1293.8         20.05	5. T P.A. 88.1 116.9 110.9 95 60 71.7 30.9 0 5. T P.A. 29.7 59.6 76.8 77.4 60.4 55.2	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE CE Conc. 2.1265 4.2674 5.5418 4.3246 3.9523	2. Cis P.A. Cis P.A.	588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc.	column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Meti P.A. 18.9 316.1 403.4 447.9 235.5 170 0 1.5 Meti P.A. 0.8 37.2 195.4 523.4 730.9 543.9	333 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00 333 Conc. 0.20 9.29 48.77 130.64 182.43 135.76
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565           PCE           P.A.         Conc.           209.9         3.25           394.4         6.11           722.4         11.20           121.19         18.78           1122.6         17.40           2032.2         31.34           2042.3         38.18	5. TP.A. 88.1 116.9 95 60 71.7 30.9 0 5. TP.A. 29.7 59.6 76.8 77.4 60.4 55.2 46.0 4 0	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE Ce Conc. 2.1265 4.2674 5.4418 5.4418 4.3246 3.9523 3.3523 3.350	2. Cis P.A. Cis P.A.	588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc.	column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Meti P.A. 18.9 316.1 403.4 447.9 235.5 170 0 1.5 Meti P.A. 0.8 37.2 195.4 523.4 730.9 543.9 94.9 0	333           hane           Conc.           4.72           78.90           100.69           111.80           58.78           119.71           27.46           0.00           333           hane           Conc.           0.20           9.29           48.77           130.64           182.43           135.76           23.69           0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565         PCE           P.A.         Conc.           209.9         3.25           394.4         6.11           722.4         11.20           121.1.9         18.78           1122.6         17.40           1293.8         20.05           2022.2         31.34           2463.3         38.18	5. T P.A. 88.1 116.9 110.9 95 60 71.7 30.9 0 5. T P.A. 29.7 59.6 76.8 77.4 60.4 55.2 46.9 0	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE CE Conc. 2.1265 4.2674 5.5418 4.3246 3.9523 3.358 0	2. Cis P.A. Cis P.A.	588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A.	EE3 0 171 sDCE Conc. SA3 0 171 sDCE Conc.	xolumn	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	hane Conc. hane Conc.	1.5 Meti P.A. 18.9 316.1 403.4 447.9 235.5 110 0 1.5 Meti P.A. 0.8 37.2 195.4 730.9 543.9 94.9 0	333           hane           Conc.           4.72           78.90           100.69           111.80           58.78           119.71           27.46           0.00           333           hane           Conc.           0.20           9.29           48.77           130.64           132.76           23.69           0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565         PCE           P.A.         Conc.           209.9         3.25           394.4         6.11           72.2.4         11.20           121.1.9         18.78           1122.6         17.40           1293.8         20.05           2022.2         31.34           2463.3         38.18	5. TP.A. 88.1 116.9 95 60 71.7 30.9 0 5. TP.A. 29.7 59.6 76.8 77.4 60.4 55.2 46.9 0	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE Conc. 2.1265 4.2674 5.4989 5.5418 4.3246 3.9523 3.358 0	2. Cis P.A. Cis P.A.	588 DCE Conc. 588 DCE Conc. 588	Col 4 3. Tran P.A. Col 5 3. Tran P.A. Col 6	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc. SA2 ( 171	xolumn	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Metl P.A. 18.9 316.1 403.4 447.9 235.5 110 0 1.5 Metl P.A. 0.8 37.2 195.4 523.4 730.9 543.9 94.9 0	333 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00 533 hane Conc. 0.20 9.29 48.77 130.64 182.43 135.76 23.69 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF RT: RT: RT: RT: RT: RT: RT: RT:	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565         PCE           P.A.         Conc.           209.9         3.25           394.4         6.11           72.4         11.20           1211.9         18.78           1122.6         17.40           1293.8         20.05           2022.2         31.34           2463.3         38.18           7.565         PCE	5. TP.A. 88.1 116.9 95 60 71.7 30.9 0 5. TP.A. 29.7 59.6 76.8 77.4 60.4 55.2 46.9 0 5. T	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE Conc. 2.1265 4.2674 5.4989 5.5418 4.3246 3.9523 3.358 0	2. Cis P.A. Cis P.A.	588 DCE Conc. 588 DCE Conc. 588 DCE	Col 4 3. Tran P.A. Col 5 3. Tran P.A. Col 6 3. Tran	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc. SA2 ( 171 sDCE	xolumn	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	1.5 Metl P.A. 18.9 316.1 403.4 447.9 235.5 110 0 1.5 Metl P.A. 0.8 37.2 195.4 523.4 730.9 543.9 94.9 0	333 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00 533 hane Conc. 0.20 9.29 48.77 130.64 182.43 135.76 23.69 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565         PCE           P.A.         Conc.           209.9         3.25           394.4         6.11           722.4         11.20           1211.9         18.78           1122.6         17.40           1293.8         20.05           2022.2         31.34           2463.3         38.18           7.565         PCE           P.A.         Conc.           2022.0         51.34           7.565         PCE           P.A.         Conc.           2463.3         38.18	5. TP.A. 88.1 116.9 95 60 71.7 30.9 0 5. TP.A. 29.7 59.6 76.8 77.4 60.4 55.2 46.9 0 5. TP.A. 29.7	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE Conc. 2.1265 4.2674 5.4989 5.5418 4.3246 3.9523 3.358 0 0	2. Cis P.A. Cis P.A. 2. Cis P.A.	588 DCE Conc. 588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A. Col 6 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc. SA2 ( 171 sDCE Conc.	column P.A. column P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	hane Conc.	1.5 Metl P.A. 18.9 316.1 403.4 447.9 235.5 110 0 1.5 Metl P.A. 0.8 37.2 195.4 730.9 543.9 94.9 0 1.5 Metl P.A.	333 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00 533 hane Conc. 0.20 9.29 48.77 130.64 182.43 135.76 23.69 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565         PCE           P.A.         Conc.           209.9         3.25           394.4         6.11           72.2.4         11.20           1211.9         18.78           1122.6         17.40           1293.8         20.05           2022.2         31.34           2463.3         38.18           7.565         PCE           P.A.         Conc.           359.6         5.57	5. TP.A. 88.1 116.9 95 60 71.7 30.9 0 5. TP.A. 29.7 59.6 76.8 77.4 60.4 55.2 46.9 0 5. TP.A. 47.3 61.1	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE Conc. 2.1265 4.2674 5.4989 5.5418 4.3246 3.9523 3.358 0 485 CE Conc. 3.3867 Conc. 3.3867 Conc.	2. Cis P.A. Cis P.A. 2. Cis P.A.	588 DCE Conc. 588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A. Col 6 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc. SA2 ( 171 sDCE Conc.	xolumn P.A.	VC Conc. VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. P.A. P.A.	hane Conc.	1.5 Metl P.A. 18.9 316.1 403.4 447.9 235.5 110 0 1.5 Metl P.A. 0.8 37.2 195.4 730.9 543.9 94.9 0 1.5 Metl P.A. 523.4 730.9 543.9 94.9 0	333 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00 533 hane Conc. 0.20 9.29 48.77 130.64 182.43 135.76 23.69 0.00 533 hane Conc. 1.32 3.34
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565           PCE         P.A.           Conc.         209.9         3.25           394.4         6.11         72.4         11.20           1211.9         18.78         1122.6         17.40           1293.8         20.05         2022.2         31.34           2463.3         38.18         2         9           7.565           PCE         P.A.         Conc.           359.6         5.57         134.8         2.09           437.7         6.78         2.09	5. TP.A. 88.1 116.9 95 60 71.7 30.9 0 5. TP.A. 29.7 59.68 77.4 60.4 55.2 46.9 0 5. TP.A. 47.3 61.1 72.6	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE Conc. 2.1265 4.2674 5.4989 5.5418 4.3246 3.9523 3.358 0 0 485 CE Conc. 3.3867 4.3748 5.1982	2. Cis P.A. Cis P.A. 2. Cis P.A.	588 DCE Conc. 588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A. Col 6 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc. SA2 ( 171 sDCE Conc.	xolumn P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. P.A. Ett P.A.	hane Conc.	1.5 Metl P.A. 18.9 316.1 403.4 447.9 235.5 110 0 1.5 Metl P.A. 0.8 37.2 195.4 730.9 543.9 94.9 0 1.5 Metl P.A. 523.4 730.9 543.9 94.9 0 1.5 Metl P.A. 13.4 417.5	333 hane Conc. 4.72 78.90 100.69 111.80 58.78 119.71 27.46 0.00 533 hane Conc. 0.20 9.29 48.77 130.64 182.43 135.76 23.69 0.00 533 hane Conc. 1.32 3.34 hane Conc. 1.32 3.34 104.21
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565           PCE         P.A.           Conc.         209.9         3.25           394.4         6.11         72.4         11.20           1211.9         18.78         1122.6         17.40           1293.8         20.05         2022.2         31.34           2463.3         38.18         2463.3         38.18           7.565           PCE         P.A.         Conc.           359.6         5.57         134.8         2.09           437.7         6.78         624.5         9.68	5. TP.A. 88.1 116.9 95 60 71.7 30.9 0 5. TP.A. 29.7 59.6 76.8 77.4 60.4 55.2 46.9 0 5. TP.A. 47.3 61.1 72.6 73.3	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE Conc. 2.1265 4.2674 5.4989 5.5418 4.3246 3.9523 3.358 0 485 CE Conc. 3.3867 4.3748 5.1982 5.2483	2. Cis P.A. Cis P.A. 2. Cis P.A.	588 DCE Conc. 588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A. Col 6 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc. SA2 ( 171 sDCE Conc.	xolumn P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. P.A. Ett P.A.	hane Conc.	1.5 Meti P.A. 18.9 316.1 403.4 447.9 235.5 110 0 1.5 Meti P.A. 0.8 37.2 195.4 730.9 543.9 94.9 0 1.5 Meti P.A. 523.4 730.9 543.9 94.9 0 1.5 Meti P.A.	333           hane           Conc.           4.72           78.90           100.69           111.80           58.78           119.71           27.46           0.00           533           hane           Conc.           0.20           9.29           48.77           130.64           182.43           135.76           23.34           0.00           533           hane           Conc.           1.32           3.34           104.21           156.40
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         39.99           7.565           PCE         P.A.           Conc.         209.9         3.25           394.4         6.11         72.4           1211.9         18.78         1122.6           122.4         11.20         2022.2           203.3         38.18           7.565           PCE         P.A.           Conc.         359.6           306.5         5.77           134.8         2.09           437.7         6.78           624.5         9.68           954.8         14.80           1381 2         24.41	5. T P.A. 88.1 116.9 95 60 71.7 30.9 0 5. T P.A. 29.7 59.6 76.8 77.4 60.4 55.2 46.9 0 5. T P.A. 47.3 61.1 72.6 73.3 54.1 72.6 73.3	485 CE Conc. 6.308 8.37 7.9404 6.802 4.296 5.1337 2.2124 0 485 CE Conc. 2.1265 4.2674 5.4989 5.5418 4.3246 3.9523 3.358 0 485 CE Conc. 3.3867 4.3748 5.1982 5.2483 3.376 3.3956	2. Cis P.A. Cis P.A. 2. Cis P.A.	588 DCE Conc. 588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A. Col 6 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc. SA2 ( 171 sDCE Conc.	xolumn	VC Conc. VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. P.A. Ett P.A.	hane Conc.	1.5 Meti P.A. 18.9 316.1 403.4 447.9 235.5 110 0 1.5 Meti P.A. 0.8 37.2 195.4 730.9 94.9 0 1.5 Meti P.A. 523.4 730.9 94.9 0 1.5 Meti P.A. 33.4 730.9 543.9 94.9 0	333           hane           Conc.           4.72           78.90           100.69           111.80           58.78           119.71           27.46           0.00           533           hane           Conc.           0.20           9.29           48.77           130.64           182.43           135.76           23.34           0.00           533           hane           Conc.           133.64           182.43           135.76           23.34           104.21           156.40           121.01           96 77
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: 2 1 1 1 1 1 1 1 1 1 1 1 1 1	7.565         PCE           P.A.         Conc.           209         3.24           232.1         3.60           630.1         9.77           1007.6         15.62           1104.9         17.13           976.5         15.14           1780.7         27.60           2579.7         3999           7.565           PCE         P.A.           Conc.         209.9         3.25           394.4         6.11         72.4         11.20           1211.9         18.78         1122.6         17.40           1293.8         20.05         2022.2         31.34           2463.3         38.18         20.05         2022.2           9.6         5.57         PCE         P.A.         Conc.           306.6         5.57         7134.8         2.09         437.7         6.78           624.5         9.68         954.8         14.80         1381.2         21.41           210.0.4         32.56         5.57         334.8         2.09	5. P.A. 88.1 116.9 95 60 71.7 30.9 0 5. T P.A. 29.7 59.6 76.8 77.4 60.4 55.2 46.9 0 T P.A. 47.3 60.7 7.4 60.4 55.2 46.9 0 T P.A. 55.2 46.9 0 5.5 T P.A. 55.2 55.2 75.6 75.6 76.8 77.4 60.4 55.2 75.6 76.8 77.4 60.4 55.2 75.6 76.8 77.4 60.4 55.2 75.6 75.6 76.8 77.4 60.4 55.2 46.9 0 75.5 75.6 75.6 76.8 77.4 60.4 55.2 46.9 0 75.5 75.6 75.6 75.6 77.4 60.4 55.2 75.6 75.6 75.6 75.6 75.6 75.6 75.6 75.6 75.6 75.2 46.9 0 75.5 75.6 75.2 75.6 75.2 75.6 75.2 75.6 75.2 75.6 75.2 75.2 75.2 75.2 75.6 75.2 75.2 75.2 75.2 75.2 75.5 75.2 75.5 75.2 75.5 75.2 75.5 75.2 75.5 75.2 75.5	485 CE Conc. 6.308 8.37 7.9404 4.296 5.1337 2.2124 0 485 CE Conc. 2.1265 4.2674 5.4989 5.5418 4.3246 3.9523 3.358 0 485 CE Conc. 3.3867 4.3748 5.1982 5.2483 3.378 0 0 3.3873 6.39953 3.378 0 0.3723	2. Cis P.A. Cis P.A. 2. Cis P.A.	588 DCE Conc. 588 DCE Conc. 588 DCE Conc.	Col 4 3. Tran P.A. Col 5 3. Tran P.A. Col 6 3. Tran P.A.	EE3 ( 171 sDCE Conc. SA3 ( 171 sDCE Conc. SA2 ( 171 sDCE Conc.	xolumn P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. P.A. Ett P.A.	hane Conc.	1.5 Meti P.A. 18.9 316.1 403.4 447.9 235.5 110 0 1.5 Meti P.A. 0.8 37.2 195.4 730.9 543.9 94.9 0 1.5 Meti P.A. 523.4 730.9 543.9 94.9 0 1.5 Meti P.A. 5.3 13.4 417.5 626.6 484.8 387.7 5.6	333           hane           Conc.           4.72           78.90           100.69           111.80           58.78           119.71           27.46           0.00           533           hane           Conc.           0.20           9.29           48.77           130.64           182.43           135.76           23.34           0.00           533           hane           Conc.           1.32.3.69           0.00           533           hane           Conc.           1.32           3.34           104.21           156.40           121.01           96.77           1.40



ĺ						Col 7	7 EE2 C	olumn us	sed 10/8	stds						
RT:	7.565		5.485		2.588	-	3.171	-					Eu		1.533	
norto	PCE	Cono	ICE	Cono		Cono	I ransDC	E Cono		Cono	Ethylene	Cono	Ethane	Cono	Methane	Cana
pons 7	P.A. 549.4	8 52	P.A. 73.5	5 2626	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A. 54	13 48
6	723.8	11.22	87.2	6.2435											265.7	66.32
5	670.4	10.39	106	7.5896											410.9	102.56
4	1142.1	17.70	102.5	7.339											367.8	91.80
3	1010.6	15.66	74.6	5.3414											237.8	59.35
2	1272.8	19.73	59.4	4.253											104.4	26.06
1	2444.2	37.89	17.7	1.2673											5	1.25
	1200	10.00	0	_ 0			0-1-0	Original							0	0.00
RT:	7.565		5.485					Control		n			1		1.533	
	PCE		TCE		CisDCE	-	FransDC	E	VC		Ethylene		Ethane		Methane	
ports	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
7	134.5	2.08	150.6	10.783											360.1	89.88
6	296.7	4.60	145	10.382											374.7	93.53
5	754	11.90	89.7	6.4225											400.9 281.1	70.16
3	1324.2	20.53	77 1	5 5204											70.6	17 62
2	1394.6	21.62	70.2	5.0263											24.1	6.02
1	1996.9	30.95	0	0											0	0.00
IFF	1572.2	24.37	0	0											0	0.00
							Col 9	SA1 d	column							
RT:	7.5	65	5.	485	2.5	588	3.1	171							1.5	33
	PC	CE	T	CE	Cis	DCE	Tran	sDCE	`	/C	Ethy	lene	Eth	ane	Meth	nane
ports	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
6	103.8	1.01	39.4	2.821											14.0	3.04
5	956.4	14 82	123.6	8 8498											374	93.35
4	953.3	14.78	105.3	7.5395											570.9	142.50
3	884.2	13.71	76.5	5.4774											549.1	137.06
2	1424.5	22.08	66.8	4.7829											167.4	41.78
1	2182	33.82	3.1	0.222											0	0.00
IFF	1720.7	26.67	0	_ 0			Col 10	EE1	oolumn						0	0.00
RT.	7.5	65	5	485	2.5	588	20110	171	column				r		1.5	33
	PC	CE	T	CE	Cis	DCE	Tran	sDCE	`	/C	Ethy	lene	Eth	ane	Meth	nane
ports	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
7	463.9	7.19	53	3.7948											36.4	6.81
6	1053.2	16.32	92.6	6.6302											274.6	68.54
5	501.8	7.78	91.7	6.5657											573.3	143.10
4	1226.2	10.16	80.5 64 9	0.1218											308.4	91.95
2	2445	37.90	153.2	10.969											159.6	39.84
1	593	9.19	30.2	2.1623											0	0.00
IFF	1808.7	28.03	0	0											0	0.00
				_			Col 11	Control	1 colum	in						
RT:	7.5	65	5.	485	2.5	588	3.*	171		10					1.5	33
norto	PC	Cono	T D A	CE	Cis	DCE	Tran	sDCE		VC Cono	Ethy	lene	Eth	ane	Meth	nane
7	105 Q	1 64	77 3	5 5347	<b>Р.</b> А.	Conc.	<b>г</b> .А.	Conc.	<b>г</b> .А.	Cond.	г.A.	Conc.	г. <b>А</b> .	Conc.	900 4	224 74
6	121.5	1.88	142.4	10,196											639.3	159 57
5	411.6	6.38	119.8	8.5777											429.9	107.30
4	593.1	9.19	105	7.518											428.7	107.00
3	706.4	10.95	77	5.5132											200.5	50.04
2	1095.3	16.98	77.3	5.5347											73.4	18.32
	1609.4	24.95	0	0			I								0	0.00
11°E	1002.3	ZJ.29	0	U											U	0.00

Week 4 column 12 data was an outlier due to the low PCE readings within the columns. column 12 data was not used during average calculations.



# Appendix G. 6th week data

				Co	11 CC:	3 Columr	1-6 ALL	CONC	IN ppb(µg	g /L)					
RT:	7.565	5.	485	2.5	588	3.1	171	,	10	E.I.	1	<b>E</b> .1		1.5	33
porte	PCE PA Con		Conc		Conc		SDCE		Conc	Eth	yiene Conc	Etr	Conc	Metr	Conc
7	38.8 0.60	62.4	4.4678	г. <del>л</del> .	Conc.	Г. <del>Л</del> .	Conc.	г. <del>д</del> .	Conc.	г. <del>д</del> .	Conc.	г. <del>л</del> .	Conc.	3.93	0.98
6	35.9 0.56	138.25	9.8987											94.77	23.65
5	125.07 1.94	129.55	9.2758											370.12	92.38
4	154.87 2.40	134.97	9.6639											505.84	126.26
3	491.06 7.61	136.66	9.7849											174.9 90.5	43.66
1	1123.6 17.4	40.94	2.9313											19.9	4.97
IFF	3180.9 49.3	0	0											0	0.00
			_			Col 2	CC2 C	Column							
RT:	7.565	5.	485											1.5	33
	PCE	Т	CE	Cis	DCE	Tran	SDCE	`	VC	Eth	ylene	Eth	nane	Meth	nane
ports	P.A. Cond	. P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
6	171 04 2 65	134.4	0.4500 9.623											30.7 80.97	20.21
5	333.71 5.17	174.3	12.48											201.7	50.34
4	457.1 7.09	112.18	8.0321											219.49	54.78
3	547.17 8.48	100.47	7.1937											192.3	48.00
2	1036.3 16.0	5 114.98	8.2326											143.57	35.84
IFF	1574.5 24.4 3135.9 48.6	0	5.5755 0											27.3	0.00
	0100.0 40.0	Ŭ				Cal 2	Control							Ŭ	0.00
RT:	7,565	5.	485				Control							1.5	33
	PCE	Т	CE	Cis	DCE	Tran	sDCE	Ň	VC	Eth	ylene	Eth	nane	Meth	nane
ports	P.A. Cond	. P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
7	191.34 2.97	142.19	10.181											790.7	197.36
6	243.44 3.77	95.74	0.000											485.92	121.29
4	935.54 14.5	113.74	8.1438											350.16	87.40
3	1188.7 18.4	103.01	7.3755											233.34	58.24
2	1017.8 15.7	121.09	8.67											274.06	68.41
1	1855.4 28.7	15.45	1.1062											5.3	1.32
IFF	3126.9 48.4	0	0												0.00
		-	-			Col 4	EE3 /	olumn						0	0.00
RT:	7.565	5.	485			Col 4	EE3 (	column						1.5	i33
RT:	7.565 PCE	5. T	485 CE	Cis	DCE	Col 4 Tran	EE3 ( sDCE	column	VC	Eth	ylene	Eth	nane	0 1.5 Meth	i33 nane
RT:	7.565 PCE P.A. Conc	5. T . P.A.	485 CE Conc.	Cisi P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 o sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	0 1.5 Meth P.A.	i33 nane Conc.
RT: ports 7 6	7.565 PCE P.A. Cono 36.04 0.56	5. T P.A. 63.4 122.08	485 CE Conc. 4.5394 8.7409	Cis P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 4.7	533 nane Conc. 1.17
RT: ports 7 6 5	7.565 PCE P.A. Cond 36.04 0.56 30.59 0.47 101.2 1.57	5. T P.A. 63.4 122.08 154.95	485 CE Conc. 4.5394 8.7409 11.094	Cisl P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Meth P.A. 4.7 421.69 431.26	533 nane Conc. 1.17 105.25 107.64
RT: ports 7 6 5 4	7.565 PCE P.A. Cond 36.04 0.56 30.59 0.47 101.2 1.57 487.75 7.56	5. T P.A. 63.4 122.08 154.95 149.2	485 CE Conc. 4.5394 8.7409 11.094 10.683	Cisl P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A. 4.7 421.69 431.26 476.82	Conc. 1.17 105.25 107.64 119.01
RT: ports 7 6 5 4 3	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.566           707.88         10.9	5. T P.A. 63.4 122.08 154.95 149.2 7 156.9	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234	Cis P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A. 4.7 421.69 431.26 476.82 359.87	533 hane Conc. 1.17 105.25 107.64 119.01 89.82
RT: ports 7 6 5 4 3 2	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           707.88         10.9           1008.4         15.6	5. T P.A. 63.4 122.08 154.95 149.2 156.9 3 111.42	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234 7.9777	Cis P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	iane Conc.	1.5 Mett P.A. 4.7 421.69 431.26 476.82 359.87 197.05	Conc. 1.17 105.25 107.64 119.01 89.82 49.18
RT: 7 6 5 4 3 2 1	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           707.88         10.9           1008.4         15.6           2148         33.2	5. T P.A. 63.4 122.08 154.95 149.2 156.9 111.42 5.03 0 0	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234 7.9777 3.94077 0	Cis P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A. 4.7 421.69 431.26 476.82 359.87 197.05 28.8 0	Conc. 1.17 105.25 107.64 119.01 89.82 49.18 7.19 0.00
RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9	5. T P.A. 63.4 122.08 154.95 149.2 156.9 111.42 5.03 0	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234 7.9777 3.9401 0	Cisl P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A. 421.69 431.26 476.82 359.87 197.05 28.8 0	Conc.           1.17           105.25           107.64           119.82           49.18           7.19           0.00
RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9	5. T P.A. 63.4 122.08 154.95 149.2 156.9 111.42 55.03 0	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234 7.9777 3.9401 0	Cis P.A.	DCE Conc.	Col 4 Tran P.A.	EE3 ( sDCE Conc.	P.A.	VC Conc.	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A. 4.7 421.69 431.26 476.82 359.87 197.05 28.8 0	0.00           333           nane           Conc.           1.17           105.25           107.64           119.01           89.82           49.18           7.19           0.00
RT: ports 7 6 5 4 3 2 1 IFF RT:	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           7.565         PCE	5. T P.A. 63.4 122.08 154.95 149.2 156.9 111.42 55.03 0 5. T	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234 7.9777 3.9401 0 485 CE	Cisi P.A. Cisi	DCE Conc.	Col 4 Tran P.A. Col 5 Tran	EE3 ( sDCE Conc. SA3 ( sDCE	P.A.	VC Conc.	Eth P.A. Eth	ylene Conc.	Ett P.A.	nane Conc.	1.5 Mett P.A. 4.7 421.69 431.26 476.82 359.87 197.05 28.8 0	Conc. 1.17 105.25 107.64 119.01 89.82 49.18 7.19 0.00 333 hane
RT: ports 7 6 5 4 3 2 1 IFF RT: ports	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond	5. T P.A. 63.4 122.08 154.95 149.2 156.9 111.42 55.03 0 5. T P.A.	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234 7.9777 3.9401 0 485 CE Ce Conc.	Cisi P.A. Cisi P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett	aane Conc. hane Conc.	0 1.5 Mett P.A. 4.7 421.69 431.26 431.26 359.87 197.05 28.8 0 1.5 Mett P.A.	Conc. 1.17 105.25 107.64 119.01 89.82 49.18 7.19 0.00 333 hane Conc.
RT: ports 7 6 5 4 3 2 1 IFF RT: 7 7	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond           33.37         5.17	5. T P.A. 63.4 122.08 154.95 149.2 156.93 111.42 55.03 0 5. T T P.A. 63.4 106.65	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361	Cisl P.A. Cisl P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	p.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	0 1.5 Mett P.A. 4.7 421.69 431.26 431.26 359.87 197.05 28.8 0 1.5 Mett P.A. 7 47 47 47 47 47 47 47 47 47	Conc. 1.17 105.25 107.64 119.01 89.82 49.18 7.19 0.00 333 hane Conc. 1.75
RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.77	5. T P.A. 63.4 122.08 154.95 149.2 156.99 111.42 55.03 0 5. T T P.A. 106.65 183.75 205 10	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.620	Cisl P.A. Cisl P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 of sDCE Conc. SA3 of sDCE Conc.	xolumn P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc.	0 1.5 Mett P.A. 4.7 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 149.55	Conc. 1.17 105.25 107.64 119.01 89.82 49.18 7.19 0.00 333 hane Conc. 1.75 4.57 3.57 3.57
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.77           335.59         5.20	5. T P.A. 63.4 122.08 154.95 149.2 156.95 111.42 55.03 0 5. T T P.A. 0 6 5. T T P.A. 106.65 183.75 205.16 138.67	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288	Cisl P.A. Cisl P.A.	DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 of sDCE Conc. SA3 of sDCE Conc.	p.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc.	0 1.5 Mett P.A. 4.7 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06	0.00 333 hane Conc. 1.17 105.25 107.64 119.01 89.82 49.18 7.19 0.00 333 hane Conc. 1.75 4.57 37.08 119.07
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.565           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.72           335.59         5.20           635.03         9.84           890.59         13.8	5. T P.A. 63.4 122.08 154.95 144.2 156.9 111.42 55.03 0 T F.A. 106.65 183.75 205.16 138.67 129.31	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.2586	Cisl P.A. Cisl P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	xolumn P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett	nane Conc. nane Conc.	0 1.5 Mett P.A. 4.7 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13	333 hane Conc. 1.17 105.25 107.64 119.01 89.82 49.18 7.19 0.00 333 hane Conc. 1.75 4.57 37.08 119.07 125.58
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.565           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           P.A.         Cond           333.37         5.17           303.13         4.70           335.59         5.26           635.03         9.84           890.59         13.8           932.17         14.4	5. T P.A. 63.4 122.08 154.95 149.2 156.9 0 111.42 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.16 183.75 205.16 138.67 129.31 142.06	485 CE Conc. 4.5394 10.084 10.083 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.2586 10.171	Cis P.A. Cis P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	xolumn P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc.	0 1.5 Mett P.A. 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45	Conc.           1.17           105.25           107.64           119.01           89.82           49.18           7.19           0.00           333           nane           Conc.           1.75           4.57           37.08           1125.58           74.24
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 1 5	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.565           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.70           335.59         5.20           635.03         9.82           932.17         14.4           1726.9         26.7	5. T P.A. 63.4 122.08 154.95 149.2 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.18 183.75 205.16 138.67 129.31 142.06 138.67 129.31 142.06 138.67 129.31 142.06 138.67 129.51 143.75 129.51 143.75 145.75 145.75 145.75 145.75 145.75 145.75 145.75 145.75 145.75 145.75 145.75 145.75 145.75 145.75 145.75 1	485 CE Conc. 4.5394 10.04 10.083 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.2586 10.171 4.4285	Cis P.A. Cis P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	xolumn P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett	nane Conc. nane Conc.	0 1.5 Mett P.A. 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8	Conc.           1.17           105.25           107.64           119.01           89.82           49.18           7.19           0.00           333           nane           Conc.           1.75           4.57           37.08           119.07           125.58           74.24           15.92
RT: ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.565           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.70           335.59         5.22           635.03         9.84           932.17         14.4           1726.9         26.7           3097         48.0	5. 7. 7. 7. 7. 7. 7. 7. 7. 7. 7	485 CE Conc. 4.5394 10.683 11.094 10.683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.2586 10.171 4.4285 0	Cis P.A. Cis P.A.	DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	olumn P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Vene Conc.	Etr P.A. Etr P.A.	nane Conc. nane Conc.	0 1.5 Mett P.A. 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0	333 hane Conc. 1.17 105.25 107.64 119.01 89.82 49.18 7.19 0.00 333 hane Conc. 1.75 4.57 37.08 119.07 125.58 74.24 15.92 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF	7.565         PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.565           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.72           355.9         5.20           635.03         9.84           890.59         13.8           932.17         14.4           1726.9         26.7           3097         48.0	5. T P.A. 63.4 122.08 154.95 144.2 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.16 183.75 205.16 138.67 129.31 142.06 138.67 0 5.205.16 138.67 129.31 142.06 138.67 129.31 142.06 138.67 129.51 142.06 138.67 129.51 142.07 159.95 100 100 100 100 100 100 100 100 100 10	485 CE Conc. 4.5394 10.083 11.094 10.0683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.2586 10.171 4.4285 0	Cis P.A. Cis P.A.	DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 of sDCE Conc. SA3 of sDCE Conc.	p.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett	nane Conc. nane Conc.	0 1.5 Mett P.A. 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0	Conc.           1.17           105.25           107.64           119.01           89.82           49.18           7.19           0.00           333           nane           Conc.           1.75           4.57           37.08           1125.58           74.24           15.92           0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: RT:	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.72           635.03         9.84           890.59         13.8           932.17         14.4           1726.9         26.7           3097         48.0           7.565         PCE	5. T P.A. 63.4 122.08 154.95 141.42 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.03 0 55.16 138.67 129.31 142.06 61.85 0 5. 129.31 142.06 5. 5. 5. 5. 5. 5. 5. 5. 5. 5. 5. 5. 5.	485 CE Conc. 4.5394 11.094 10.683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.2586 10.171 4.4285 0	Cis P.A. Cis P.A.	DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6	EE3 ( sDCE Conc. SA3 ( sDCE Conc.	xolumn P.A. xolumn P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	nane Conc.	0 1.5 Mett P.A. 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0 1.5 Mett	Conc.           1.17           105.25           107.64           119.01           89.82           49.18           7.19           0.00           333           nane           Conc.           1.75           1.75           1.75           1.75           1.75           1.75           37.08           1125.58           74.24           15.92           0.00           333
RT: ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 7 7 8 8 8 7 8 8 8 8 8 8 8 8 8 8 8 8 8	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.72           355.9         5.20           635.03         9.84           890.59         13.8           932.17         14.4           1726.9         26.7           3097         48.0           7.565         PCE           P.A.         Cond	5. T P.A. 63.4 122.08 154.95 149.2 156.9 0 111.42 55.03 0 T P.A. 106.65 183.75 205.16 138.67 129.31 142.06 61.85 0 5. T 142.08 13.42 5. 140.2 5. 0 T 129.31 120.88 13.42 5. 0 T 129.51 13.42 5. 0 T 129.51 13.69 5. 149.2 5. 149.2 5. 149.2 5. 149.2 5. 149.2 5. 149.2 5. 149.2 5. 149.2 5. 149.2 5. 149.2 5. 149.2 5. 15.9 5. 149.2 5. 149.2 5. 149.2 5. 149.2 5. 15.9 5. 149.2 5. 15.9 5. 149.2 5. 149.2 5. 15.9 5. 149.2 5. 15.9 111.42 5. 15.9 5. 149.2 5. 149.2 5. 15.9 5. 15.9 5. 15.9 5. 15.9 5. 16.9 5. 183.75 205.16 5. 142.20 5. 142.20 5. 142.20 5. 5. 129.31 5. 142.20 5. 7. 129.31 5. 7. 7. 9. 7. 129.31 5. 7. 7. 7. 7. 7. 7. 7. 7. 7. 7	485 CE Conc. 4.5394 10.083 11.094 10.0683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.2586 10.171 4.4285 0 485 CE CE Conc. CE Conc.	Cis P.A. Cis P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran	EE3 ( sDCE Conc. SA3 ( sDCE Conc. SA2 ( sDCE Conc.	xolumn P.A. xolumn P.A.	VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene ylene Conc.	Ett P.A. Ett P.A.	nane Conc. nane Conc.	0 1.5 Mett P.A. 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0 1.5 Mett P.A.	Conc.           1.17           105.25           107.64           119.01           89.82           49.18           7.19           0.00           333           nane           Conc.           1.75           4.57           37.08           119.07           125.58           74.24           15.92           0.00           333           nane           Conc.           Conc.           Conc.           Conc.           Conc.           Conc.
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 7 6 5 4 3 2 1 IFF RT: ports 7 7 7 7 7 8 8 9 9 9 9 9 9 9 9 9 9 9 9 9	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.70           335.59         5.20           635.03         9.8           932.17         14.4           1726.9         26.7           3097         48.0           7.565         PCE           P.A.         Cond           90.89         1.41	5. 7. 7. 7. 7. 7. 7. 7. 7. 7. 7	485 CE Conc. 4.5394 10.683 11.094 10.683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.2586 10.171 4.4285 0 485 CE CE Conc. 7.3247	Cisl P.A. Cisl P.A. Cisl P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran P.A.	EE3 of sDCE Conc. SA3 of sDCE Conc. SA2 of sDCE Conc.	xolumn P.A. xolumn P.A.	VC Conc. VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	hane Conc.	0 1.5 Mett P.A. 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 63.8 0 1.5 63.8 0 1.5 Mett P.A. 4.7 148.55 63.8 0 1.5 Mett P.A. 4.7 18.3 148.55 63.8 0 1.5 63.8 1.5 63.8 1.5 63.8 1.5 63.8 1.5 63.8 1.5 63.8 1.5 63.8 1.5 63.8 1.5 63.8 1.5 63.8 1.5 1.5 1.5 1.5 1.5 1.5 1.5 1.5	Conc.           1.17           105.25           107.64           119.01           89.82           49.18           7.19           0.00           333           nane           Conc.           1.75           4.57           37.08           119.07           125.58           74.24           15.92           0.00           333           nane           Conc.           1.33           Conc.           1.33           Conc.           1.175
RT: ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.70           335.59         5.20           635.03         9.8           8932.17         14.4           1726.9         26.7           3097         48.0           7.565         PCE           P.A.         Cond           90.89         1.41           36.4         0.56	5. 7. 7. 7. 7. 7. 7. 7. 7. 7. 7	485 CE Conc. 4.5394 10.683 11.094 10.683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.2586 10.171 4.4285 0 485 CE Conc. 7.3247 5.0979	Cisi P.A. Cisi P.A. Cisi P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc. SA2 ( sDCE Conc.	p.A.	VC Conc. VC Conc. VC Conc.	Eth P.A. Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A. Ett P.A.	nane Conc.	0 1.5 Mett P.A. 4.7 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0 1.5 Mett P.A. 7 1.5 Mett P.A. 7 1.5 Mett P.A. 7 1.5 Mett P.A. 7 1.5 Mett P.A. 7 1.5 Mett P.A. 7 1.5 Mett P.A. 7 1.5 Mett P.A. 7 1.5 Mett P.A. 7 1.5 Mett P.A. 7 1.5 Mett P.A. 7 1.5 Mett P.A. 7 1.5 Mett P.A. 7 1.5 3.1 297.45 6.3.8 0 1.5 Mett P.A. 7 1.5 6.3.8 0 1.5 Mett P.A. 7 1.5 6.3.8 0 1.5 Mett P.A. 7 1.5 6.3.8 0 1.5 Mett P.A. 7 1.5 6.3.8 0 1.5 Mett P.A. 7 1.5 6.3.8 0 1.5 Mett P.A. 7 1.5 6.3.8 0 1.5 Mett P.A. 7 1.5 6.3.8 0 1.5 Mett P.A. 7 1.5 6.3.8 0 1.5 Mett P.A. 7 1.5 6.3.8 0 1.5 Mett P.A. 7 1.5 Mett P.A. 1.5 1.5 1.5 1.5 1.5 1.5 1.5 1.	0.00           333           hane           Conc.           1.17           105.25           107.64           119.01           89.82           49.18           7.19           0.00           333           hane           Conc.           1.75           4.57           37.08           119.07           125.58           74.24           15.92           0.00           333           hane           Conc.           1.18           6.56
RT: ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 IFF Ports 7 6 5 4 3 2 1 I I I I I I I I I I I I I	7.565         PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           707.88         10.9           1008.4         15.6           2148         32.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.70           335.59         5.20           635.03         9.84           890.59         13.8           932.17         14.4           172.69         26.7           3097         48.0           7.565         PCE           P.A.         Cond           90.89         1.41           36.4         0.56           167.71         2.66           167.71         2.67	5. 7 7 7 7 7 7 7 7 7 7 7 7 7	485 CE Conc. 4.5394 10.683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.2586 10.171 4.4285 0 485 CE Conc. 7.3247 5.0979 11.367	Cisl P.A. Cisl P.A. Cisl P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc. SA2 ( sDCE Conc.	column P.A. column P.A.	VC Conc. VC Conc. VC Conc.	Eth P.A. P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A. Ett	nane Conc.	0 1.5 Mett P.A. 4.7 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0 1.5 Mett P.A. 4.7 26.3 182.27 246.32 182.27 192.45 192.4	Conc.           1.17           105.25           107.64           119.01           89.82           49.18           7.19           0.00           333           nane           Conc.           1.75           37.08           119.07           125.58           74.24           15.92           0.00           333           nane           Conc.           1.75           37.08           119.07           125.58           74.24           15.92           0.00           333           nane           Conc.           1.18           6.56           45.49
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 8 7 6 5 4 3 2 1 IFF RT: 8 7 6 5 4 3 2 7 6 5 4 3 7 6 5 4 3 7 6 5 4 3 7 6 5 4 3 7 6 5 4 3 7 8 7 7 8 8 8 8 8 8 8 8 8 8 8 8 8	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           707.88         10.9           1008.4         15.6           2148         33.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.70           335.59         5.20           635.03         9.84           890.59         13.8           932.17         14.4           172.69         26.7           3097         48.0           7.565         PCE           P.A.         Cond           3097         48.0           7.565         PCE           P.A.         Cond           90.89         1.41           36.4         0.56           167.71         2.66           166.2, 7.27         7.27           766.73         11.8	5. 7. 7. 7. 7. 7. 7. 7. 7. 7. 7	485 CE Conc. 4.5394 10.683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.2586 10.171 4.4285 0 485 CE Conc. 7.3247 5.0979 11.367 5.0979 11.314 9.4297	Cis P.A. Cis P.A.	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc. SA2 ( sDCE Conc.	p.A.	VC Conc. VC Conc. VC Conc.	Eth P.A. P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A. Ett	nane Conc.	0 1.5 Mett P.A. 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0 1.5 Mett P.A. 7 28.8 0 1.5 18.3 148.55 477.06 503.13 297.45 63.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0 1.5 1.5 1.5 1.5 1.5 1.5 1.5 1.5	Conc. 1.17 105.25 107.64 119.01 89.82 49.18 7.19 0.00 333 hane Conc. 1.75 37.08 119.07 125.58 74.24 15.92 0.00 333 hane Conc. 1.75 37.08 119.07 125.58 74.24 15.92 0.00 333 hane Conc. 1.75 8.57 37.08 119.07 125.58 74.24 15.92 0.00 333 hane Conc. 1.75 8.57 37.08 119.07 125.58 74.24 15.92 0.00 333 hane Conc. 1.75 8.57 37.08 119.07 125.58 74.24 15.92 0.00 333 hane Conc. 1.75 8.57 37.08 119.07 125.58 74.24 15.92 0.00 333 hane Conc. 1.78 8.57 37.08 119.07 125.58 74.24 15.92 0.00 333 hane Conc. 1.75 8.57 37.08 119.07 125.58 74.24 15.92 0.00 333 hane Conc. 1.78 8.57 37.08 125.58 74.24 15.92 0.00 333 hane Conc. 1.78 8.57 37.08 125.58 74.24 15.92 0.00 333 hane Conc. 1.78 8.56 4.549 8.249 8.249 8.249 1.78
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           707.88         10.9           1008.4         15.6           2148         332.2           3413.9         52.9           7.565         PCE           P.A.         Cond           333.37         5.17           303.13         4.70           335.59         5.22           635.03         9.84           890.59         13.8           932.17         14.4           1726.9         26.7           3097         48.0           7.565         PCE           P.A.         Cond           3097         48.0           7.565         PCE           P.A.         Cond           90.89         1.41           36.4         0.50           167.71         2.60           469.2         7.27           766.73         11.8           1069.6         16.5	5. 7. 7. 7. 7. 7. 7. 7. 7. 7. 7	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.9288 9.9288 9.2586 10.171 4.4285 0 9.9288 9.2586 10.171 4.4285 0 485 CE Conc. 7.3247 7.3247 Conc. 7.3247 7.3247 Conc. 7.3247 7.3247 Conc. 7.3247 7.3247 Conc. 7.3247 7.3247 Conc. 7.3247 7.3247 Conc. 7.3247 7.3247 Conc. 7.3247 7.3247 Conc.	Cis P.A. Cis P.A. Cis	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran P.A.	EE3 ( sDCE Conc. SA3 ( sDCE Conc. SA2 ( sDCE Conc.	p.A.	VC Conc. VC Conc. VC Conc.	Eth P.A. Eth P.A. Eth P.A.	ylene Conc. ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	hane Conc.	0 1.5 Mett P.A. 4.7 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0 1.5 Mett P.A. 7 26.3 182.27 26.3 182.27 209.14	Conc.           1.17           105.25           107.64           119.01           89.82           49.18           7.19           0.00           333           nane           Conc.           1.75           45.18           7.19           0.00           333           nane           Conc.           1.75           4.57           37.08           74.24           15.92           0.00           333           nane           Conc.           1.75           4.592           0.00           333           nane           Conc.           1.18           6.56           45.49           61.58           82.42           52.20
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: 1 PORTS 7 6 5 4 3 2 1 IFF RT: 2 1 1 IFF RT: 2 1 1 IFF RT: 2 1 1 IFF RT: 2 1 1 IFF RT: 2 1 1 1 1 1 1 1 1 1 1 1 1 1	7.565           PCE           P.A.         Cond           36.04         0.56           30.59         0.47           101.2         1.57           487.75         7.56           707.88         10.9           1008.4         15.6           2148         332.           3413.9         52.9           7.565         PCE           P.A.         Cond           333.59         5.22           635.03         9.84           890.59         13.8           932.17         14.4           1726.9         26.7           3097         48.0           7.565         PCE           P.A.         Cond           309.59         13.8           932.17         14.4           1726.9         26.7           90.89         1.44           136.4         0.56           PCE         P.A.           08.9         1.41           36.4         0.56           167.71         2.60           469.2         7.27           766.73         113           1069.6	5. 7. 7. 7. 7. 7. 7. 7. 7. 7. 7	485 CE Conc. 4.5394 8.7409 11.094 10.683 11.234 7.9777 3.9401 0 485 CE Conc. 7.6361 13.157 14.689 9.9288 9.9288 9.9288 9.9288 9.9288 9.2586 10.171 4.4285 0 485 CE Conc. 7.3247 5.0979 11.367 13.114 9.4297 7.0326 0	Cisl P.A. Cisl P.A. Cisl	DCE Conc. DCE Conc.	Col 4 Tran P.A. Col 5 Tran P.A. Col 6 Tran P.A.	EE3 of sDCE Conc. SA3 of sDCE Conc. SA2 of sDCE Conc.	p.A.	VC Conc. VC Conc. VC Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Ett P.A. Ett P.A.	ane Conc.	1.5 Mett P.A. 421.69 431.26 476.82 359.87 197.05 28.8 0 1.5 Mett P.A. 7 18.3 148.55 477.06 503.13 297.45 63.8 0 1.5 Mett P.A. 4.73 26.3 182.27 246.72 330.2 209.14 1.03	Conc.           1.17           105.25           107.64           119.01           89.82           49.18           7.19           0.00           333           nane           Conc.           1.75           4.57           37.08           74.24           15.92           0.00           333           nane           Conc.           1.75           4.5.92           0.00           333           nane           Conc.           1.18           6.56           45.49           61.58           82.42           52.20           0.26



					Col 7	7 EE2 C	olumn us	ed 10/8	stds						
RT:	7.565 PCF	5.485 TCF		2.5 CisE	88 )CE	3.1 Tran	171 sDCE	\	/C	Eth	vlene	Eth	ane	1.5 Meth	i33 Jane
ports	P.A. Conc.	P.A. Cor	IC.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
7	290.54 4.50	105.23 7.53	45											33.95	8.47
6	549.18 8.51 205.04 3.18	157.44 11.2	38											282.55	70.52
4	334.64 5.19	124.06 8.88	27											662.02	165.24
3	497.81 7.72	151.23 10.8	28											270.35	67.48
2	1094.1 16.96	113.81 8.14	88											80.52	20.10
1 IFF	1931 29.93 3002 46.53	0 0												0	0.00
	40.00	<u> </u>	-			Col 8	Control 2	2 Colum	n					Ŭ	0.00
RT:	7.565	5.485				_								1.5	33
porte	PCE P A Conc		20	CisL	Conc	I ran	SDCE		/C	Eth	ylene	Eth PA	Conc	Meth	Conc
7	31.68 0.49	132.72 9.50	28	т. <del>.</del> .	00110.	1.4.	Conc.	т.д.	Conc.	1	Conc.	1.4.	Conc.	513.4	128.14
6	41.86 0.65	162.15 11.6	61											527.67	131.71
5	579.69 8.99	177.76 12.7	28											383.56	95.74
4	802.63 12.44 719.9 11.16	279.14 19.9	58											90.62	39.28 22.62
2	1835 28.44	143.62 10.2	83											7.48	1.87
1	1953.3 30.28	0 0												0	0.00
IFF	2985 46.27	0 0	L											0	0.00
RT:	7.565	5.485				Col 9	SA1 d	olumn						1.5	33
	PCE	TCE		CisE	CE	Tran	sDCE	١	/C	Eth	ylene	Eth	ane	Meth	nane
ports	P.A. Conc.	P.A. Cor	IC.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.	P.A.	Conc.
6	57.68 0.89 119.31 1.85	63.67 4.55 124.56 8.91	85											16.52 42.7	4.12
5	464.81 7.20	162.87 11.6	61											172	42.93
4	702.14 10.88	203.08 14.5	41											666.3	166.31
3	646.85 10.03	121.83 8.72	23											331.46	82.73
2	1213.1 18.80 2499.2 38.74	101.61 7.27	53											196.12	48.95
IFF	3284 50.90	0 0												0	0.00
						Cal 10									
DT	7.505	5 405					EEI	column				-			
RT:	7.565 PCE	5.485 TCE		CisE	CE	Tran	SDCE	column \	/C	Eth	ylene	Eth	iane	1.5 Meth	i33 nane
RT: ports	7.565 PCE P.A. Conc.	5.485 TCE P.A. Cor	nc.	CisE P.A.	OCE Conc.	Tran P.A.	sDCE Conc.	column ۱ P.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	iane Conc.	1.5 Meth P.A.	i33 nane Conc.
RT: ports 7	7.565 PCE P.A. Conc. 259.36 4.02 517.2 8.02	5.485 TCE P.A. Cor 564.9 40.4	nc. 47	CisE P.A.	OCE Conc.	Tran P.A.	sDCE Conc.	column ۱ P.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	iane Conc.	1.5 Meth P.A. 0	33 nane Conc. 0.00 61.83
RT: ports 7 6 5	7.565 PCE P.A. Conc. 259.36 4.02 517.2 8.02 704.2 10.92	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4	nc. 47 106	Cist P.A.	OCE Conc.	Tran P.A.	sDCE Conc.	<u>column</u> ۲.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	ane Conc.	1.5 Meth P.A. 0 247.73 429.98	33 nane Conc. 0.00 61.83 107.32
RT: ports 7 6 5 4	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10	nc. 47 06 44 86	CisE P.A.	OCE Conc.	Tran P.A.	sDCE Conc.	Column \ P.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	iane Conc.	1.5 Meth P.A. 0 247.73 429.98 401.01	i33 nane Conc. 0.00 61.83 107.32 100.09
RT: ports 7 6 5 4 3	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2	nc. 47 006 44 866 868	Cist P.A.	OCE Conc.	P.A.	SDCE Conc.	<u>column</u> ۲.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	iane Conc.	1.5 Meth P.A. 0 247.73 429.98 401.01 332.68	33 nane Conc. 0.00 61.83 107.32 100.09 83.04
RT: ports 7 6 5 4 3 2 1	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 231.26 16.5 18.5 1.32	nc. 47 006 444 868 58 58	CisE P.A.	OCE Conc.	Tran P.A.	SDCE Conc.	Column N.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	iane Conc.	1.5 Meth P.A. 0 247.73 429.98 401.01 332.68 135.89 1 27	33 nane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32
RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 231.26 16.5 18.5 1.32 0 0	nc. 447 006 444 986 988 958 958	CisE P.A.	OCE Conc.	Tran P.A.	SDCE Conc.	Column N P.A.	/C Conc.	Eth P.A.	ylene Conc.	Eth P.A.	ane Conc.	1.5 Meth P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0	33 nane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.00
RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 231.26 16.5 18.5 1.32 0 0	nc. 47 006 44 886 868 58 846	CisE P.A.	DCE Conc.	Tran P.A.	SDCE Conc.	P.A.	/C Conc. n	Eth P.A.	ylene Conc.	Eth P.A.	nane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT:	7.565 PCE P.A. Conc. 259.36 4.02 517.2 8.02 704.2 10.92 743.02 11.52 802.96 12.45 1168.2 18.11 1651.1 25.59 1063.6 16.49 7.565 PCE	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 231.26 16.5 18.5 1.32 0 0	nc. 447 006 444 868 558 558 246	CisE P.A. CisF	DCE Conc.	Col 10 Tran P.A. Col 11 Tran	SDCE Conc.	Column V P.A.	/C Conc. n	Eth P.A. Eth	ylene Conc.	Eth P.A. Eth	Conc.	1.5 Meth P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.00 33 hane
RT: ports 7 6 5 4 3 2 1 IFF RT: ports	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49           7.565         PCE           P.A.         Conc.	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 231.26 16.5 18.5 1.32 0 0 5.485 TCE P.A. Cor	nc. 447 006 444 986 558 558 446	CisE P.A. CisE P.A.	DCE Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	Control sDCE Conc.	Column V P.A. 1 colum V P.A.	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 1.27 0	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.00 333 hane Conc.
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 2	7.565 PCE P.A. Conc. 259.36 4.02 517.2 8.02 704.2 10.92 743.02 11.52 802.96 12.45 1168.2 18.11 1651.1 25.59 1063.6 16.49 7.565 PCE P.A. Conc. 170.63 2.64	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 231.26 16.5 18.5 1.32 0 0 5.485 TCE P.A. Cor 262.77 18.8	nc. 447 1006 444 1886 288 246 246 246	CisE P.A. CisE P.A.	DCE Conc.	Col 10 Tran. P.A. Col 11 Tran. P.A.	Control sDCE Conc.	P.A.	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc. nane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 1.27 0	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.32 0.00 333 hane Conc. 45.41
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565 PCE P.A. Conc. 259.36 4.02 517.2 8.02 704.2 10.92 743.02 11.52 802.96 12.45 1168.2 18.11 1651.1 25.59 1063.6 16.49 7.565 PCE P.A. Conc. 170.63 2.64 88.94 1.38	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 231.26 16.5 18.5 1.32 0 0 5.485 TCE P.A. Cor 262.77 18.8 140.25 10.0	nc. 447 006 444 886 858 446 .588 446 .58 .446	CisE P.A. CisE P.A.	DCE Conc.	Col 10 Tran. P.A. Col 11 Tran. P.A.	Control sDCE Conc.	P.A.	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	ane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 1.5 Mett P.A. 181.93 313.07 623.67	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.32 0.00 33 hane Conc. 45.41 78.14 155.67
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4	7.565 PCE P.A. Conc. 259.36 4.02 517.2 8.02 704.2 10.92 743.02 11.52 802.96 12.45 1168.2 18.11 1651.1 25.59 1063.6 16.49 7.565 PCE P.A. Conc. 170.63 2.64 88.94 1.38 109.24 1.69 206.62 3.20	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 231.26 16.5 18.5 1.32 0 0 5.485 TCE P.A. Cor 262.77 18.8 140.25 10.0 116.54 8.34	nc. 447 006 444 886 688 558 646 144 1442 443 77	CisE P.A. CisE P.A.	DCE Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	Control SDCE Conc. SDCE Conc.	Lolumn	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	ane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 1.5 Mett P.A. 181.93 313.07 623.67 14.53	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.32 0.00 33 hane Conc. 45.41 78.14 155.67 3.63
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49           PCE         P.A.         Conc.           170.63         2.64           88.94         1.38           109.24         1.69           206.62         3.20           405.43         6.28	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 231.26 16.5 18.5 1.32 0 0 5.485 TCE P.A. Cor 262.77 18.8 140.25 10.0 116.54 8.34 54.05 3.8 121.9 8.72	nc. 447 006 444 886 688 558 446 .58 446 .58 144 442 443 77 28	CisE P.A. CisE P.A.	DCE Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	Control SDCE Conc. SDCE Conc.	Lolumn	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 1.5 Mett P.A. 181.93 313.07 623.67 14.53 96.3	33 aane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.32 0.00 33 aane Conc. 45.41 78.14 155.67 3.63 24.04
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 2 1 IFF	7.565 PCE P.A. Conc. 259.36 4.02 517.2 8.02 704.2 10.92 743.02 11.52 802.96 12.45 1168.2 18.11 1651.1 25.59 1063.6 16.49 7.565 PCE P.A. Conc. 170.63 2.64 88.94 1.38 109.24 1.69 206.62 3.20 405.43 6.28 168.65 2.61	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 231.26 16.5 18.5 1.32 0 0 5.485 TCE P.A. Cor 262.77 18.8 140.25 10.0 116.54 8.34 54.05 3.8 121.9 8.77 165.37 11.4	nc. 447 006 444 886 688 558 646 558 646 100 114 442 443 77 28 84	CisE P.A. CisE P.A.	DCE Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	Control sDCE Control sDCE Conc.	<u>Column</u> P.A. <u>1 colum</u> P.A.	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 1.5 Mett P.A. 181.93 313.07 623.67 14.53 96.3 359.49	33 aane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.32 0.00 33 aane Conc. 45.41 78.14 155.67 3.63 24.04 89.73
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49           7.565         PCE           P.A.         Conc.           170.63         2.64           88.94         1.38           109.24         1.69           206.62         3.20           405.43         6.28           168.5         2.61           2138.6         33.15           3345         51.85	5.485           TCE           P.A.         Cor           564.9         40.4           107.41         7.69           257.6         18.4           29.45         2.10           157.37         11.2           231.26         16.5           18.5         1.32           0         0           5.485           TCE         P.A.           Cor         262.77           262.77         18.8           140.25         10.0           116.54         8.34           54.05         3.8           121.9         8.72           165.37         11.4           0         0           2         0.14	nc. 447 006 444 886 688 558 446 558 446 558 446 114 442 443 77 28 84 32	CisE P.A. CisE P.A.	DCE Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	Control sDCE Control sDCE Conc.	Lolumn	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 1.5 Mett P.A. 181.93 313.07 623.67 14.53 96.3 359.49 0 0	33 aane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.32 0.00 33 anane Conc. 45.41 78.14 155.67 3.63 24.04 89.73 0.00 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF IFF	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49           7.565         PCE           P.A.         Conc.           170.63         2.64           88.94         1.38           109.24         1.69           206.62         3.20           405.43         6.28           168.5         2.61           2138.6         33.15           3345         51.85	5.485           TCE           P.A.         Cor           564.9         40.4           107.41         7.69           257.6         18.4           29.45         2.10           157.37         11.2           231.26         16.5           18.5         1.32           0         0           5.485         TCE           P.A.         Cor           262.77         18.8           140.25         10.0           116.54         8.34           54.05         3.8           121.9         8.7           165.37         11.4           0         0           2         0.14	nc. 447 006 648 668 558 548 548 548 548 548 548 548 548 54	CisE P.A. CisE P.A.	DCE Conc. DCE Conc.	Col 11 Col 11 Tran P.A. Col 11 Col 12	Conc.	1 column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	nane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.00 333 hane Conc. 45.41 78.14 155.67 3.63 24.04 89.73 0.00 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 7 8 8 8 8 8 8 8 8 8 8 8 8 8	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49           7.565         PCE           P.A.         Conc.           170.63         2.64           88.94         1.38           109.24         1.69           206.62         3.20           405.43         6.28           1238.6         33.15           3345         51.85	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 231.26 16.5 18.5 1.32 0 0 5.485 TCE P.A. Cor 262.77 18.8 140.25 10.0 116.54 8.34 54.05 3.8 121.9 8.77 165.37 11.1 0 0 2 0.14	nc. 447 006 444 886 658 558 446  114 442 43 77 28 84 32	CisE P.A. CisE P.A.	DCE Conc.	Col 11 <u>Tran</u> P.A. <u>Col 11</u> <u>Tran</u> P.A.	Control Conc. Control SDCE Conc. SA0	1 column P.A. P.A. Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. Eth P.A.	ane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 1.5 Mett P.A. 181.93 313.07 623.67 14.53 96.3 359.49 0 0	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.32 0.32 0.00 33 hane Conc. 45.41 155.67 3.63 24.04 89.73 0.00 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 7 8 8 8 8 8 8 8 8 8 8 8 8 8	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49           7.565         PCE           P.A.         Conc.           170.63         2.64           88.94         1.38           109.24         1.69           206.62         3.20           405.43         6.28           168.65         2.61           2138.6         33.15           3345         51.85           PCE         P.A.           Conc.         7.565           PCE         P.A.	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 31.26 16.5 18.5 1.32 0 0 5.485 TCE P.A. Cor 262.77 18.8 140.25 10.0 116.54 8.34 54.05 3.8 121.9 8.77 165.37 11.4 0 0 2 0.14	nc. 447 006 444 886 668 558 646 100 114 442 443 77 28 84 32	CisE P.A. CisE P.A.	DCE Conc.	Col 10 Tran P.A. Col 11 Tran P.A.	Control sDCE Conc. sDCE Conc. SA0 sDCE Conc.	1 column N.A. P.A. P.A. column	/C Conc. n //C Conc.	Eth P.A. Eth P.A.	ylene ylene Conc. ylene ylene Conc.	Eth P.A. Eth P.A.	ane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 1.5 Mett P.A. 181.93 313.07 623.67 14.53 96.3 359.49 0 0	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.32 0.32 0.00 33 hane Conc. 45.41 155.67 3.63 24.04 89.73 0.00 0.00
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 7 8 7 7 8 8 8 8 8 8 8 8 8 8 8 8 8	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49           7.565         PCE           P.A.         Conc.           170.63         2.64           88.94         1.38           109.24         1.69           206.62         3.20           405.43         6.28           133.65         51.85           7.565         PCE           P.A.         Conc.           13345         51.85           7.565         PCE           P.A.         Conc.           7.565         PCE           P.A.         Conc.           7.565         PCE           P.A.         Conc.           7.565         PCE           P.A.         Conc.           7.506         2.73	5.485           TCE           P.A.         Cor           564.9         40.4           107.41         7.69           257.6         18.4           29.45         2.10           157.37         11.2           231.26         16.5           18.5         1.32           0         0           5.485         TCE           P.A.         Cor           262.77         18.8           140.25         10.0           116.54         8.34           54.05         3.8           121.9         8.77           165.37         11.3           0         0           2         0.14           5.485         TCE           P.A.         Cor           9         0.4	nc. 47 006 44 886 68 58 68 58 68 58 68 58 646 114 43 77 28 84 32	CisE P.A. CisE P.A. CisE P.A.	DCE Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 12 Tran P.A.	Control sDCE Conc. sDCE Conc. SA0 sDCE Conc.	1 column N.A. P.A. P.A. column	/C Conc. n //C Conc. //C Conc.	Eth P.A. P.A. P.A.	ylene Vlene Conc. ylene Conc.	Eth P.A. P.A. P.A.	ane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 1.5 Mett P.A. 181.93 313.07 623.67 14.53 96.3 359.49 0 0	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.32 0.32 0.32 0.32 0.32 0.32 0.3
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49           7.565         PCE           P.A.         Conc.           170.63         2.64           88.94         1.38           109.24         1.69           206.62         3.20           405.43         6.28           168.65         2.61           2138.6         33.15           3345         51.85           PCE         P.A.           P.A.         Conc.           7.565         PCE           P.A.         Conc.           3345         51.85	5.485 TCE P.A. Cor 564.9 40.4 107.41 7.69 257.6 18.4 29.45 2.10 157.37 11.2 231.26 16.5 18.5 1.32 0 0 5.485 TCE P.A. Cor 262.77 18.8 140.25 10.0 116.54 8.34 54.05 3.8 121.9 8.77 165.37 11.1 0 0 2 0.14 5.485 TCE P.A. Cor 91.08 6.52 118.79 8.50	nc. 47 006 44 886 68 58 68 58 68 58 68 58 68 58 67 84 6 84 6 84 32 6 77 28 84 32 6 77 28 84 32 6 77 28 84 32 6 77 77 28 84 32 6 77 77 77 77 77 77 77 77 77 77 77 77 7	CisE P.A. CisE P.A. CisE P.A.	DCE Conc. DCE Conc. DCE Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 12 Tran P.A.	Control SDCE Conc. SA0 SDCE Conc. SA0 Conc.	Column	/C Conc. n /C Conc. /C Conc.	Eth P.A. P.A. P.A.	ylene Vlene Conc. ylene Conc.	Eth P.A. P.A. Eth P.A.	ane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 1.5 Mett P.A. 181.93 313.07 623.67 14.53 96.3 359.49 0 0 0	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.32 0.32 0.32 0.32 0.32 0.32 0.32 0.3
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 7 6 5 4 8 7 7 6 5 4 8 7 7 6 5 4 8 7 7 6 5 4 8 7 7 6 5 4 8 7 7 6 5 4 8 7 7 6 5 4 8 7 7 8 7 8 7 8 8 8 8 8 8 8 8 8 8 8 8 8	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49           7.565         PCE           P.A.         Conc.           170.63         2.64           88.94         1.38           109.24         1.69           206.62         3.20           405.43         6.28           168.65         2.61           2138.6         33.15           3345         51.85           P.A.         Conc.           7.565         PCE           P.A.         Conc.           7.565         PCE           P.A.         Conc.           7.565         PCE           P.A.         Conc.           7.565         PCE           P.A.         Conc.           7.566         Conc.           7.566         Conc.           7.566	5.485           TCE           P.A.         Cor           564.9         40.4           107.41         7.69           257.6         18.4           29.45         2.10           157.37         11.2           231.26         16.5           18.5         1.32           0         0           5.485         TCE           P.A.         Cor           262.77         18.8           140.25         10.0           116.54         8.34           54.05         3.8           121.9         8.71           165.37         11.2           0         0           2         0.14           5.485         TCE           P.A.         Cor           9.08         6.52           118.79         8.50           141.09         10.1           15.485         TCE           P.A.         Cor           91.08         6.52           118.79         8.50           141.09         10.1           141.09         10.1	nc. 447 006 444 886 688 558 558 558 558 558 558 558 558	CisE P.A. CisE P.A. CisE P.A.	DCE Conc. DCE Conc. DCE Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 12 Tran P.A.	Control SDCE Conc. SA0 SDCE Conc. SA0 SDCE Conc.	1 column 1 colum P.A. N P.A.	/C Conc. n /C Conc. /C Conc.	Eth P.A. P.A. P.A.	ylene Vlene Conc. ylene Conc.	Eth P.A. P.A. Eth P.A.	ane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 14.53 96.3 353.07 623.67 14.53 96.3 359.49 0 0 0	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.00 33 hane Conc. 45.41 78.14 155.67 3.63 24.04 89.73 0.00 0.00 33 hane Conc. 9.10 11.61 33.40 26.06 26.06 26.06 26.06 26.06 26.06 26.06 26.07 26.06 26.
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 1 IFF RT: 7 6 5 4 3 2 7 6 5 4 3 7 6 5 4 3 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 6 5 7 8 7 8 7 8 7 8 7 8 7 8 7 8 7 8 7 8 7 8 8 8 8 8 8 8 8 8 8 8 8 8	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49           7.565           PCE         P.A.           7.06.3         2.64           88.94         1.38           109.24         1.69           206.62         3.20           405.43         6.28           168.65         2.61           2138.6         33.15           3345         51.85           7.565           PCE           P.A.         Conc.           176.06         2.73           183.77         2.85           440.58         6.83           336.36         5.21           778.75         12.07	5.485           TCE           P.A.         Cor           564.9         40.4           107.41         7.69           257.6         18.4           29.45         2.10           157.37         11.2           231.26         16.5           18.5         1.32           0         0           5.485         TCE           P.A.         Cor           262.77         18.8           140.25         10.0           116.54         8.34           54.05         3.8           121.9         8.7           165.37         11.2           0         0           2         0.14           5.485         TCE           P.A.         Cor           9.108         6.52           118.79         8.50           141.09         10.1           135.39         9.69	nc. 447 006 444 886 688 558 558 558 558 558 558 558 558	CisE P.A. CisE P.A. CisE P.A.	DCE Conc. DCE Conc.	Col 10 Tran P.A. Col 11 Tran P.A. Col 12 Tran P.A.	Conc. Control SDCE Conc. SA0 SDCE Conc.	1 column 1 colum P.A. P.A.	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Vlene Conc. ylene Conc.	Eth P.A. P.A. Eth P.A.	ane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 14.53 96.3 353.07 623.67 14.53 96.3 359.49 0 0 0 1.5 Mett P.A. 36.47 46.52 133.82 104.52 133.82	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.00 33 hane Conc. 45.41 78.14 155.67 3.63 24.04 89.73 0.00 0.00 33 hane Conc. 9.10 11.61 33.40 26.09 28.95
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF RT: 2 2 1 IFF RT: 2 2 1 IFF RT: 2 2 1 IFF RT: 2 2 1 IFF RT: 2 2 1 IFF RT: 2 2 1 IFF RT: 2 2 2 1 2 2 2 1 2 2 2 2 2 2 2 2 2 2 2 2 2	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49           7.565         PCE           P.A.         Conc.           170.63         2.64           88.94         1.38           109.24         1.69           206.62         3.20           405.43         6.28           168.65         2.61           2138.6         33.15           3345         51.85           PCE         P.A.           Conc.         77.565           PCE         P.A.           7.565         PCE           P.A.         Conc.           7.565         PCE           P.A.         Conc.           77.565         2.61           2138.6         33.15           336.36         5.21           778.75         12.07           9	5.485           TCE           P.A.         Cor           564.9         40.4           107.41         7.69           257.6         18.4           29.45         2.10           157.37         11.2           231.26         16.5           18.5         1.32           0         0           5.485         TCE           P.A.         Cor           262.77         18.8           140.25         10.0           116.54         8.34           54.05         3.8           121.9         8.7           165.37         11.2           0         0           2         0.14           5.485         TCE           P.A.         Cor           9.108         6.52           118.79         8.50           141.09         10.1           116.11         8.31           135.39         9.69           142.01         10.1	nc. 447 006 444 868 868 846 846 100 114 443 37 728 84 32 100 113 154 02 35 339 68	CisE P.A. CisE P.A. CisE P.A.	DCE Conc. DCE Conc.	Col 11 P.A. Col 11 P.A. Col 12 Col 12 Tran P.A.	Control SDCE Conc. SA0 SA0 SDCE Conc.	Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. P.A. Eth P.A.	ane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 14.53 96.3 359.49 0 0 0 0 1.5 Mett P.A. 14.53 96.3 359.49 0 0 0 0 1.5 Mett P.A. 15.89 11.53 96.3 359.49 0 0 0 0	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.00 33 hane Conc. 45.41 78.14 155.63 24.04 89.73 0.00 0.00 33 hane Conc. 9.10 11.61 33.40 26.09 28.95 16.69
RT: ports 7 6 5 4 3 2 1 IFF RT: ports 7 6 5 4 3 2 1 IFF RT: Ports 7 6 5 4 3 2 1 IFF	7.565           PCE           P.A.         Conc.           259.36         4.02           517.2         8.02           704.2         10.92           743.02         11.52           802.96         12.45           1168.2         18.11           1651.1         25.59           1063.6         16.49           7.565         PCE           P.A.         Conc.           170.63         2.64           88.94         1.38           109.24         1.69           206.62         3.20           405.43         6.28           168.65         2.61           2138.6         33.15           3345         51.85           7.565         PCE           P.A.         Conc.           176.06         2.73           183.77         2.85           440.58         6.83           336.36         5.21           778.75         12.07           969.88         15.03           1548.5         24.00	5.485           TCE           P.A.         Cor           564.9         40.4           107.41         7.69           257.6         18.4           29.45         2.10           157.37         11.2           231.26         16.5           18.5         1.32           0         0           5.485         TCE           P.A.         Cor           262.77         18.8           140.25         10.0           116.54         8.34           54.05         3.8           121.9         8.7           165.37         11.3           0         0           2         0.14           5.485         TCE           P.A.         Cor           9.10.8         6.52           118.79         8.50           141.09         10.1           135.39         9.69           142.01         10.1           31.55         2.2           0         0	nc. 447 006 444 868 868 846 846 100 114 443 37 728 84 32 100 113 154 02 35 339 68 59	CisE P.A. CisE P.A. CisE P.A.	DCE Conc. DCE Conc.	Col 11 P.A. Col 11 Tran P.A. Col 12 Tran P.A.	Control SDCE Conc. SA0 SA0 SDCE Conc.	1 column 1 colum P.A. Column	/C Conc. n /C Conc.	Eth P.A. Eth P.A.	ylene Conc. ylene Conc.	Eth P.A. P.A. Eth P.A.	ane Conc.	1.5 Mett P.A. 0 247.73 429.98 401.01 332.68 135.89 1.27 0 14.53 96.3 359.49 0 0 0 1.5 Mett P.A. 14.53 96.3 359.49 0 0 0 1.5 Mett P.A. 13.82 13.82 13.82 13.82 13.82 104.52 13.82 104.52 13.82 104.52 13.82 104.52 13.82 104.52 13.82 104.52 13.82 104.52 13.82 104.52 13.82 104.52 13.82 104.52 104.53 105.89 10.5 10.5 10.5 10.5 10.5 10.5 10.5 10.5	33 hane Conc. 0.00 61.83 107.32 100.09 83.04 33.92 0.00 33 hane Conc. 45.41 78.14 155.63 24.04 89.73 0.00 0.00 33 hane Conc. 9.10 11.61 33.40 26.09 18.89 16.69 1.69

Week 6 column 10 data was an outlier due to the low PCE readings within the columns. column 10 data was not used during average calculations.



### Appendix H nitrate and sulfate raw data

11/18/2005 Col 7 EE2 ALL CONC IN ppm(mg /L) RT: 12.250 12.830 14

Chloride

Nitrite

15.090

Sulfate

14.760

Nitrate

12.25 12.83 14.76	15.09
Chloride Nitrite Nitrate	Sulfate
Conc. Conc. Conc.	Conc.
119.714 8.175	18.989
107.655 6.795	13.810
105.232 5.663	14.549
104.911 5.408	18.546
104.551 5.307	23.010
100.840 5.052	29.373
97 348 4 879	28.843
011010	2010/10
CC2 ALL CONC IN ppm(mg /L)	
12.25 12.83 14.76	15.09
Chloride Nitrite Nitrate	Sulfate
Conc. Conc. Conc.	Conc.
121.285 3.918	16.718
100.194 7.988	13.392
101.802 0.800	14.721
100.676 5.235	20.802
100.789 5.204	24.981
98.448 5.085	28.493
95.327 4.951	28.134
LON         ALL CONC IN ppm(mg /L)           12.25         12.83         14.76	15.09
Chloride Nitrite Nitrate	Sulfate
Conc. Conc. Conc.	Conc.
99.690 5.228	21.462
99.373 5.348	19.919
97.200 6.055	15.531
97.416 5.328	17.789
98.406 5.129	21.368
97.950 5.015	22.683
	~~ ~
98.179 4.812	28.641
98.179 94.936 4.641	28.641 27.416
98.179 4.812 94.936 4.641 EE3 ALL CONC IN ppm(mg /L) 12.25 12.83 14.76	28.641 27.416
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)           12.25         12.83           Chloride         Nitrite	28.641 27.416 15.09 Sulfate
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)           12.25         12.83           Chloride         Nitrite           Nitrate         Conc.	28.641 27.416 15.09 Sulfate Conc.
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)           12.25         12.83           14.76           Chloride         Nitrite           Conc.         Conc.           151.81756         7.462	28.641 27.416 15.09 Sulfate Conc. 10.012
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)           12.25         12.83           14.76           Chloride         Nitrite           Nitrite         Nitrate           Conc.         Conc.           151.81756         7.462           135.0578         6.21	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)           12.25         12.83           14.76           Chloride         Nitrite           Nitrate           Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)           12.25         12.83           Chloride         Nitrite           Nitrate           Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)           12.25         12.83           Chloride         Nitrite           Nitrate           Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)           12.25         12.83           14.76           Chloride         Nitrite           Nitrite         Nitrate           Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)           12.25         12.83           14.76           Chloride         Nitrite           Nitrate           Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           1844.97         5.13           151.77573         4.065	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.440
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)           12.25         12.83           14.76           Chloride         Nitrite           Nitrite         Nitrate           Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           188.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25           12.83         14.76           Chloride         Nitrite	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462         135.0578           137.45724         5.63         140.01008           144.02705         5.14         148.4497           151.77573         4.065         167.96252           167.96252         4.08         SA3 ALL CONC IN ppm(mg /L)           12.25         12.83         14.76           Chloride         Nitrite         Nitrate	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc.
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)           12.25         12.83           14.76           Chloride         Nitrite           Nitrite         Nitrate           Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg /L)           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           132.898         8.243         8.243	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrate         Nitrate           Conc.         Conc.           Conc.         Conc.           132.898         8.243           109.440         8.417	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           188.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrate         Conc.           Conc.         Conc.           132.898         8.243           109.440         8.417           99.705         6.623	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           188.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrate         Conc.           Conc.         Conc.           12.25         12.83           109.440         8.417           99.705         6.623           98.710         5.755	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)         12.25           12.25         12.83         14.76           Choride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg /L)         12.25           132.898         8.243           109.440         8.417           99.705         6.623           98.710         5.755           99.650         5.093	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg /L)         12.25           12.89         14.76           Chloride         Nitrite           Nitrate         Nitrate           0.9440         8.243           109.440         8.417           99.705         6.623           98.710         5.755           99.650         5.093           99.605         5.002	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrate         0.623           132.898         8.243           109.440         8.417           99.705         6.623           98.600         5.093           99.605         5.002           98.263         4.817           99.605         5.002           98.263         4.817	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888 28.948 25.65
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg /L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrate         0.623           SA3 ALL CONC IN ppm(mg /L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrate         0.065           99.05         5.093           99.605         5.093           99.605         5.002           98.263         4.817           94.535         4.641	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888 28.948 27.534
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg /L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrate         0.623           98.710         5.755           99.650         5.093           99.605         5.093           99.605         5.092           98.263         4.817           94.535         4.641	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888 28.948 22.422 22.888
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrate         Nitrate           Conc.         Conc.           Conc.         Conc.           132.898         8.243           109.440         8.417           99.705         6.623           98.605         5.093           99.605         5.002           98.263         4.817           94.535         4.641           SA2 ALL CONC IN ppm(mg/L)         12.25	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888 28.948 27.534
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462         135.0578           137.45724         5.63           140.01008         5.39           144.02705         5.14           184.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25           132.898         8.243           109.440         8.417           99.705         6.623           98.605         5.002           98.263         4.817           94.535         4.641           SA2 ALL CONC IN ppm(mg/L)           12.25         12.83           14.76           Chloride         Nitrite           05.002         98.263           4.817         94.535           12.83         14.76           Chloride         Nitrite           SA2 ALL CONC IN ppm(mg/L)	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888 28.948 27.534
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)           12.25         12.83           14.76           Chloride         Nitrite           Nitrite         Nitrate           Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         12.83           SA3 ALL CONC IN ppm(mg/L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrate         Conc.           Conc.         Conc.           Conc.         Conc.           132.898         8.243           109.440         8.417           99.705         6.623           98.710         5.755           99.650         5.002           98.263         4.817           94.535         4.641           SA2 ALL CONC IN ppm(mg/L)           12.25         12.83 </td <td>28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888 28.948 27.534</td>	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888 28.948 27.534
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25           12.89         8.243           109.440         8.417           99.705         6.623           98.710         5.755           99.650         5.093           99.605         5.002           98.263         4.817           94.535         4.641           SA2 ALL CONC IN ppm(mg/L)           12.25         12.83           14.76           Chloride         Nitrite           14.817           94.535         4.641           SA2 ALL CONC IN ppm(mg/L)	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 15.321 14.689 18.028 22.422 22.888 28.948 27.534 15.09 Sulfate Conc. 13.848 4.700
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrite         Nitrate           Conc.         Conc.           Conc.         Conc.           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           132.898         8.243         109.440           98.710         5.755         5.993           99.605         5.002         98.263           94.535         4.641           SA2 ALL CONC IN ppm(mg/L)         12.25           12.25	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888 28.948 27.534 15.09 Sulfate Conc. 13.648 17.594
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrate         Nitrate           Conc.         Conc.           Conc.         Conc.           132.898         8.243           109.440         8.417           99.705         6.623           98.605         5.002           98.263         4.817           94.535         4.641           SA2 ALL CONC IN ppm(mg/L)           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc. <td>28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888 28.948 27.534 15.09 Sulfate Conc. 13.848 1.739 11.342 16.721</td>	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888 28.948 27.534 15.09 Sulfate Conc. 13.848 1.739 11.342 16.721
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrate         Conc.           Conc.         Conc.           12.25         12.83           14.76         S.093           99.605         5.002           98.710         5.755           99.650         5.002           98.263         4.817           94.535         4.641      SA2 ALL CONC IN ppm(mg/L)         12.25           12.83         14.76           Chloride         Nitrite	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888 28.948 27.534 15.09 Sulfate Conc. 13.848 1.739 11.342 16.731 22.310
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg /L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           151.81756         7.462           135.0578         6.21           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg /L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrate         Conc.           Conc.         Conc.           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           132.898         8.243         109.400           98.710         5.755         99.650           98.605         5.002         98.263           98.605         5.002         98.263           98.7	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 18.028 22.422 22.888 28.948 27.534 15.09 Sulfate Conc. 13.848 1.739 11.342 16.731 22.310 24.697
98.179         4.812           94.936         4.641           EE3 ALL CONC IN ppm(mg/L)         12.25           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           137.45724         5.63           140.01008         5.39           144.02705         5.14           148.4497         5.13           151.77573         4.065           167.96252         4.08           SA3 ALL CONC IN ppm(mg/L)         12.25           12.83         14.76           Chloride         Nitrite           Nitrite         Nitrate           Conc.         Conc.           Conc.         Conc.           12.25         12.83         14.76           Chloride         Nitrite         Nitrate           Conc.         Conc.         Conc.           132.898         8.243         109.440           98.710         5.755         5.093           98.650         5.093         99.605           5.093         99.605         5.002           98.263         4.817           94.535	28.641 27.416 15.09 Sulfate Conc. 10.012 8.346 12.036 22.198 25.857 31.281 43.984 50.110 15.09 Sulfate Conc. 15.949 15.321 14.689 15.321 14.689 15.321 14.689 18.028 22.422 22.888 28.948 27.534 15.09 Sulfate Conc. 13.848 1.739 11.342 16.731 22.310 24.697 28.174

	Onionae	TAILING	TNILLALO	Oullate
ports	Conc.	Conc.	Conc.	Conc.
7	124 023		7 4 1 7	9.812
, C	111.020		5.914	10.070
0	111.069		5.614	10.979
5	112.377		5.692	12.726
4	114.207		5.487	18.922
3	116 872		5 282	21 312
5	110.072		5.202	21.512
2	118.283		5.120	26.564
1	121.167		4.954	35.540
IFF	132.301		4.268	41.211
Col 8 CON	I 2 ALL CON	IC IN ppm(m	ıg /L)	
RT <sup>.</sup>	12 250	12 830	14 760	15 090
	Chlorido	Nitrito	Nitrato	Sulfato
	Onionae	Nittle	Nillale	Ounate
ports	Conc.	Conc.	Conc.	Conc.
7	107.495	n.a.	5.985	11.287
6	106.149	n.a.	6.206	8.018
5	106 878	na	5 7/3	16 454
	100.070	n.a.	5.745	10.454
4	106.731	n.a.	5.590	20.150
3	106.236	n.a.	5.501	23.361
2	105.538	n.a.	5.221	29.735
1	104,392	n.a.	5.204	30,672
IEE	104 250		E 160	20 604
0-10 00	104.200	11.d.	5.100	30.004
LOIY SA		וו ppm(mg	/L)	
Col 9 SA1	ALL CONC	IN ppm(mg /	′L)	
RT:	12.250	12.830	14.760	15.090
	Chloride	Nitrito	Nitrato	Sulfate
	Onionae	Orma	Orre	Ounate
ports	Conc.	Conc.	Conc.	Conc.
54	221.897	n.a.	11.417	19.089
45	112.613	n.a.	7.950	15.979
36	106.619	n.a.	6.157	18.623
27	106 668	n.a.	5 710	20.478
21	100.000	n.a.	5.715	20.470
21	107.410	n.a.	5.655	21.508
15	106.553	n.a.	5.387	26.127
9	104.510	n.a.	5.130	30.748
0	104 705	na	5 1 2 8	30 649
			a /L)	
			ig /L)	15 000
RI:	12.250	12.830	14.760	15.090
	Chloride	Nitrite	Nitrate	Sulfate
ports	Conc.	Conc.	Conc.	Conc.
7	106 872	0.000	7 275	10 128
6	100.072	0.000	1.213	10.120
ю	100.698	0.012	5.523	15.550
5	100.891	0.011	5.409	16.056
4	101.098	0.000	5.518	19.196
3	100 570			21.067
0	102.576	0.000	5.207	21.307
2	102.576	0.000	5.207 5.357	20 181
2	102.576	0.000	5.207 5.357	20.181
2	102.576 103.731 105.538	0.000 0.000 0.000	5.207 5.357 4.901	20.181 31.258
2 1 IFF	102.576 103.731 105.538 112.629	0.000 0.000 0.000 0.000	5.207 5.357 4.901 4.966	20.181 31.258 33.057
2 1 IFF Col 11 C0	102.576 103.731 105.538 112.629	0.000 0.000 0.000 0.000 NC IN ppm(n	5.207 5.357 4.901 4.966	20.181 31.258 33.057
2 1 IFF Col 11 CC	102.576 103.731 105.538 112.629 DN ALL COI	0.000 0.000 0.000 0.000 NC IN ppm(n	5.207 5.357 4.901 4.966 ng /L)	20.181 31.258 33.057
2 1 IFF Col 11 CC	102.576 103.731 105.538 112.629 DN ALL COI 12.250	0.000 0.000 0.000 0.000 NC IN ppm(n 12.830	5.207 5.357 4.901 4.966 ng /L) 14.760	20.181 31.258 33.057 15.090
2 1 IFF Col 11 CC RT:	102.576 103.731 105.538 112.629 ON ALL COI 12.250 Chloride	0.000 0.000 0.000 0.000 NC IN ppm(n 12.830 Nitrite	5.207 5.357 4.901 4.966 ng /L) 14.760 Nitrate	20.181 31.258 33.057 15.090 Sulfate
2 1 IFF Col 11 CC RT: ports	102.576 103.731 105.538 112.629 ON ALL COI 12.250 Chloride Conc.	0.000 0.000 0.000 NC IN ppm(n 12.830 Nitrite Conc.	5.207 5.357 4.901 4.966 ng /L) 14.760 Nitrate Conc.	20.181 31.258 33.057 15.090 Sulfate Conc.
2 1 IFF Col 11 CC RT: ports 7	102.576 103.731 105.538 112.629 ON ALL COI 12.250 Chloride Conc. 119.511	0.000 0.000 0.000 NC IN ppm(n 12.830 Nitrite Conc. n.a.	5.207 5.357 4.901 4.966 ng /L) 14.760 Nitrate Conc. 6.709	20.181 31.258 33.057 15.090 Sulfate Conc. 18.880
2 1 IFF Col 11 CC RT: ports 7 6	102.576 103.731 105.538 112.629 ON ALL COI 12.250 Chloride Conc. 119.511 106.757	0.000 0.000 0.000 NC IN ppm(n 12.830 Nitrite Conc. n.a. n.a	5.207 5.357 4.901 4.966 ng /L) 14.760 Nitrate Conc. 6.709 6.241	20.181 31.258 33.057 15.090 Sulfate Conc. 18.880 9.485
2 1 IFF Col 11 CC RT: ports 7 6 5	102.576 103.731 105.538 112.629 ON ALL COI 12.250 Chloride Conc. 119.511 106.757 107.020	0.000 0.000 0.000 NC IN ppm(n 12.830 Nitrite Conc. n.a. n.a.	5.207 5.357 4.901 4.966 ng /L) 14.760 Nitrate Conc. 6.709 6.241 5.846	21.307 20.181 31.258 33.057 15.090 Sulfate Conc. 18.880 9.485 15.072
2 1 IFF Col 11 CC RT: ports 7 6 5	102:576 103:731 105:538 112:629 DN ALL COI 12:250 Chloride Conc. 119:511 106:757 107:090	0.000 0.000 0.000 NC IN ppm(n 12.830 Nitrite Conc. n.a. n.a. n.a.	5.207 5.357 4.901 4.966 ng /L) 14.760 Nitrate Conc. 6.709 6.241 5.846	20.181 31.258 33.057 15.090 Sulfate Conc. 18.880 9.485 15.072
2 1 IFF Col 11 CC RT: ports 7 6 5 4	102.576 103.731 105.538 112.629 ON ALL COI 12.250 Chloride Conc. 119.511 106.757 107.090 106.126	0.000 0.000 0.000 NC IN ppm(n 12.830 Nitrite Conc. n.a. n.a. n.a. n.a.	5.207 5.357 4.901 4.966 ng /L) 14.760 Nitrate Conc. 6.709 6.241 5.846 5.437	20.181 31.258 33.057 15.090 Sulfate Conc. 18.880 9.485 15.072 20.379
2 1 IFF Col 11 CC RT: ports 7 6 5 4 3	102.576 103.731 105.538 112.629 ON ALL COI 12.250 Chloride Conc. 119.511 106.757 107.090 106.126 105.628	0.000 0.000 0.000 NC IN ppm(n 12.830 Nitrite Conc. n.a. n.a. n.a. n.a. n.a. n.a.	5.207 5.357 4.901 4.966 mg /L) 14.760 Nitrate Conc. 6.709 6.241 5.846 5.437 5.313	20.181 31.258 33.057 15.090 Sulfate Conc. 18.880 9.485 15.072 20.379 23.769
2 1 IFF Col 11 CC RT: ports 7 6 5 4 3 2	102.576 103.731 105.538 112.629 ON ALL COI 12.250 Chloride Conc. 119.511 106.757 107.090 106.126 105.628 105.958	0.000 0.000 0.000 NC IN ppm(n 12.830 Nitrite Conc. n.a. n.a. n.a. n.a. n.a. n.a. n.a.	5.207 5.357 4.901 14.760 Nitrate Conc. 6.709 6.241 5.846 5.437 5.313 5.211	20.181 31.258 33.057 15.090 Sulfate Conc. 18.880 9.485 15.072 20.379 23.769 28.912
2 1 IFF Col 11 CC RT: 7 6 5 4 3 2 1	102.576 103.731 105.538 112.629 DN ALL COI 12.250 Chloride Conc. 119.511 106.757 107.090 106.126 105.628 105.958 104.220	0.000 0.000 0.000 NC IN ppm(n 12.830 Nitrite Conc. n.a. n.a. n.a. n.a. n.a. n.a. n.a. n	5.207 5.357 4.901 4.966 ng /L) 14.760 Nitrate Conc. 6.709 6.241 5.846 5.437 5.313 5.211 5.136	20.181 31.258 33.057 15.090 Sulfate Conc. 18.880 9.485 15.072 20.379 23.769 28.912 30.600
2 1 IFF Col 11 CC RT: ports 7 6 5 4 3 2 1 IFF	102.576 103.731 105.538 112.629 ON ALL COI 12.250 Chloride Conc. 119.511 106.757 107.090 106.126 105.628 105.958 104.220	0.000 0.000 0.000 NC IN ppm(n 12.830 Nitrite Conc. n.a. n.a. n.a. n.a. n.a. n.a. n.a.	5.207 5.357 4.901 4.966 ng /L) 14.760 Nitrate Conc. 6.709 6.241 5.846 5.437 5.313 5.211 5.136 5.966	20.181 31.258 33.057 15.090 Sulfate Conc. 18.880 9.485 15.072 20.379 23.769 23.769 23.769 23.769 23.769
2 1 IFF Col 11 CC RT: 7 6 5 4 3 2 1 IFF	102.576 103.731 105.538 112.629 N ALL COI 12.250 Chloride Conc. 119.511 106.757 107.090 106.126 105.628 105.958 104.220 102.658	0.000 0.000 0.000 <u>0.000</u> <u>Nitrite</u> Conc. n.a. n.a. n.a. n.a. n.a. n.a. n.a. n	5.207 5.357 4.901 4.966 ng /L) 14.760 Nitrate Conc. 6.709 6.241 5.846 5.437 5.313 5.211 5.136 5.066	20.181 31.258 33.057 15.090 Sulfate Conc. 18.880 9.485 15.072 20.379 23.769 28.912 30.600 30.067
2 1 IFF Col 11 CC RT: 7 6 5 4 3 2 1 IFF	102.576 103.731 105.538 112.629 DN ALL COI 12.250 Chloride Conc. 119.511 106.757 107.090 106.126 105.628 105.958 104.220 102.658	0.000 0.000 0.000 NC IN ppm(n 12.830 Nitrite Conc. n.a. n.a. n.a. n.a. n.a. n.a. n.a. n	5.207 5.357 4.901 4.966 ng /L) 14.760 Nitrate Conc. 6.709 6.241 5.846 5.437 5.313 5.211 5.136 5.066	20.181 31.258 33.057 15.090 Sulfate Conc. 18.880 9.485 15.072 20.379 23.769 23.769 23.769 23.769 23.912 30.600 30.067

Col 12 SA 0 ALL CONC IN ppm(mg /L)										
RT:	12.250	12.830	14.760	15.090						
	Chloride	Nitrite	Nitrate	Sulfate						
ports	Conc.	Conc.	Conc.	Conc.						
7	106.187	n.a.	6.351	6.534						
6	120.845	n.a.	8.138	8.966						
5	106.669	n.a.	5.841	18.491						
4	105.077	n.a.	5.648	17.860						
3	105.188	n.a.	5.300	22.768						
2	104.148	n.a.	5.167	25.907						
1	103.528	n.a.	5.106	31.898						
IFF	104.067	n.a.	5.035	30.473						



### Bibliography

### (1) Literature Cited

- Amon, J. P.; Shelly, M. L.; Agrawal, A., Development of a wetland constructed for treatment of groundwater contaminated by chlorinated ethenes. Hydrology, soil, and vegetation. *Submitted to environmental engineering*, 2006.
- (2) Anderson, T. A.; Coats, J. R. (Eds). *Bioremediation through rhizosphere technology;* American Chemical Society: Washington, DC: 1994; p 249.
- (3) Armstrong, W.; Cousins, D.; Armstrong, J.; Turner, D. W.; Beckett, P. M. Oxygen Distribution in Wetland Plant Roots and Permeability Barriers to Gas-exchange with the Rhizosphere: a Microelectrode and Modelling Study with Phragmites australis. *Annals of Botany* 2000, *86*, 687-703.
- (4) Astin, J. and Sanders, D.A. The Role of Cost in Remedy Selection *J. of Env. Cleanup Costs, Technologies, and Techniques* 1996, *6* (3), 55-67.
- (5) Bankston, J. L.; Sola, D. L.; Komor, A. T.; Dwyer, D. F. Degradation of trichloroethylene in wetland microcosms containing broad-leaved cattail and eastern cottonwood. *Water Res.* 2002, *36* (6), 1539-1546.
- (6) Bouwer, E. J. *Bioremediation of Organic Contaminates in the subsurface* In Mitchell, R. (Ed.); Environmental Microbiology; John Wiley & Sons: New York: 1992; p 287-318.
- (7) Bouwer, E. J.; McCARTY, P. L. Transformations of 1- and 2-Carbon Halogenated Aliphatic Organic Compounds Under Methanogenic Conditions. *Applied And Environmental Microbiology* 1983, *Vol. 45* (No. 4), 1286-1294.
- (8) Bouwer, E. J.; Rittmann, B. E.; McCarty, P. L. Anaerobic Degradation of Halogenated 1 and 2 Carbon Organic Compounds. *Environmental Science & Technology* 1981, *15* (5), 596-599.
- (9) Chapelle, F. H. *Groundwater Microbiology and Geochemistr;* John Wiley and Sons Inc.: New York: 1993; p 588.



- (10) Clemmer, N. D. Characterization of Chlorinated Solvent Degradation in a Constructed Wetland.Graduate School ofEngineering and Management. *MS thesis* 2003, *AFIT/GEE/ENV/03-03*.
- (11) Cunningham, S. D.; shann, J. R.; Crowley, D. E. *Phytoremediation of contaminated water and soil;* In Kruger, E. L.; Anderson, T. A.; Coats, J. R. (Eds.);
  Phytoremediation of soil and water contaminates; American Chemical Society: Washington, D.C.: 1997; p 2-9.
- (12) Dolan; E, M.; McCarty; L, P. Small-column microcosm for assessing methanestimulated vinyl chloride transformation in aquifer samples. *Environmental Science* & *Technology* 1995, 29 (8), 1892.
- (13) Eguchi, Masahiro et al. A Field Evaluation Of In Situ Biodegradation Of Trichloroethylene Through Methane Injection. *Water. Resource* 2001, *35* (9), 2145-2152.
- (14) Environmental Protection Agency (EPA). *Treatment Technology for site cleanup: Annual Status Report (11th edition).*; EPA: Washington, D.C.: Office of solid waste and Emergency Response, 2004; EPA-542-R-03-001.
- (15) Environmental Protection Agency (EPA). Drinking Water Standards. Adapted from EPA 816-F-02-013 <u>http://www.epa.gov/safewater/mcl.html.</u> 2002, 2005 (Feb/01), 1.
- (16) Environmental Protection Agency (EPA). Innovative Treatment Technologies: Annual Status Report (8th Edition).; EPA: Washington, D.C.: Office of solid waste and Emergency response, 1996; EPA-542-R-96-010.
- (17) Fan, A. M. Trichloroethylene: water contamination and health risk assessment. . *Rev. Environ. Contam toxicol* 1988, *101*, 55-92.
- (18) Fogel, M.; Tadde, A.; Fogel, S. Biodegradation of Chlorinated Ethenes by a Methane-Utilizing Mixed Culture. *Applied And Environmental Microbiology* 1986, 4, 720-724.
- (19) Freemann, D. L.; Gossett, J. M. Biological Reductive Dechlorination of Tetrachloroethylene and Trichloroethylene to Ethylene under Methanogenic Conditions. *Applied and Environmental Microbiology* 1989, *55* (9), 2144-2151.



- (20) Isalou; Mansour; Sleep; E, B.; Liss; N, S. Biodegradation of high concentrations of tetrachloroethene in a continuous flow column system. *Environmental Science & Technology* 1998, *32* (22), 3579.
- (21) Johnson, R. L.; Pankow, J. F. Dissolution of Dense Chlorinated Solvents into Groundwater. 2. Source Functions for Pools of Solvent. *Environ. Sci. Technol.* 1992, 26, 896-901.
- (22) Lee, M. D.; Odom, J. M.; Buchanan, R. J. New Perspectives On Microbial DehalogenationOf Chlorinated Solvents:Insights from the Field. *Annu. Rev. Microbiol.* 1998, *52*, 423-452.
- (23) Lelie, Dan van der; Schwitzguebel, J. P.; Glass, D. J.; Vangronsveld, J.; Baker, A. Assessing Phytoremediation's Progress. *Environmental science and technology* 2001, *35* (21), A446-A452.
- (24) Little, C. Deane et al. Trichloroethylene Biodegradation by a Methane-Oxidizing Bacterium *Applied And Environmental Microbiology* 1988, *54* (4), 951-956.
- (25) Lorah, M. M.; Voytek, M. A.; Spencer, T. A. Preliminary Assessment of Microbial Communities and Biodegradation of Chlorinated Volatile Organic Compounds in Wetland at Cluster 13, Lauderick Creek Area, Aberdeen Proving Ground, Maryland. USGS 2003, 03-4119.
- (26) Lorah, M.; Olsen, L. Degradation of 1,1,2,2-Tetrachloroethane in a Freshwater Tidal Wetland: Field and Laboratory Evidence. *Environ. Sci. Technol.* 1999, *33*, 227-234.
- (27) Lunney, A. I.; Zeeb, B. A.; Reimer, K. J. Uptake of weathered DDT in vascular plants: Potential for phytoremediation. *Environmental Science & Technology* 2004, 38 (22), 6147.
- (28) Lynge, E.; Anttila, A.; Hemminki, K. Organic solvents and cancer. *Cancer Causes & Control* 1997, 8 (3), 406-419.
- (29) Masters, G. M. *Introduction to environmental engineering and science;* Prentice Hall: Upper Saddle River, N.J: 1997; p 651.
- (30) Mitsch, W. J.; Gosselink, J. G. Wetland; Van Nostrand Reinhold: New york: 1993; p 721.



105

- (31) Pankow, James F. and Cherry, John A. *Dense Chlorinated Solvents and other DNAPLs in Groundwater;* Waterloo Press: Portland, Oregon: 1996; p 515.
- (32) Pankow, J. F.; Cherry, J. A. *Dense Chlorinated Solvents and other DNAPLs in Groundwater;* Waterloo Press: Portland: 1996; p 511.
- (33) Schnoor, J. L.; Louis, L. A.; Mccutcheon, S. C.; Wolfe, N. L.; Carreira, L. H. Phytoremediation of Organic and Nutrient contamination. *Environmental science* and technology 1995, 29 (7), 318A-323A.
- (34) Smidt, Hauke and de Vos, Willem M. Anaerobic Microbial Degradation. *Annu. Rev. Microbiol.* 2004, *58*, 43-73.
- (35) Sobolewski, T. A. Characterization of solvent degradation profile due to microbial and chemical processes in a constructed wetland. 2003, *AFIT/GEM/ENV/04M-17*.
- (36) United States General Accounting Office. Environmental Protection: Challenges in Defense Environmental Program Management; US Government Printing Office: Washington D.C., 1995; GAO/T-NSIAD-95-121.
- (37) USDA and NRCS. The Plant Database, Version 3.5 <u>http://plants.usda.gov</u> National Plant Data Center, Baton Rouge, LA 70874-4490 USA. 2004, .
- (38) Van Der Lelie, Daniel; Schwitzguebel, J.; Glass, D. J.; Vangronsveld, J.; Baker, A. Assessing Phytoremediation's Progress. *Environmental Science and Technology* 2001, *35* (21), A446.
- (39) Vogel, T.; Criddle, C.; McCarty, P. Transformations of halogenated aliphatic compounds. *Environmental Science & Technology* 1987, *21* (8), 722-736.
- (40) Vogel, T. M.; Roudier, P. Considering innovative groundwater technologies. *Pollution Engineering* 1998, , 18.
- (41) White, J. C.; Parrish, Z. D.; Isleyen, M.; Gent, M. P. N.; Iannucci-Berger, W.; Eitzer, B. D.; Mattina, M. I. Influence of nutrient amendments on the phytoextraction of weathered 2,2-bis(p-chlorophenyl)-1,1-dichloroethylene by cucurbits. *Environ. Toxicol. Chem.* 2005, *24* (4), 987-994.



www.manaraa.com

- (42) Wiedemeier, T. H.; Rifai, H. S.; Newell, C. J.; Wilson, J. T. Natural Attenuation of Fuels and Chlorinated Solvents in the Subsurface; John Wiley & Sons, Inc: New York: 1999; p 611.
- (43) Wilson, J. T.; Wilson, B. H. Biotransformation of Trichloroethylene in Soil . *Applied And Environmental Microbiology* 1985, *49* (1), 242-243.
- (44) Yang, Lei et al. Feasibility of bioremediation of trichloroethylene contaminated sites by nitrifying bacteria through cometabolism with ammonia. *Journal of Hazardous Materials* 1999, *B69*, 111-126.
- (45) Zumdahl, s. s. Chemistry; Houghton Mifflin Company: Boston: 1997; p 1118.



#### Vita

2<sup>nd</sup> Lt Jun Yan graduated from Niles North High School in Skokie, Illinois. He attended the University of Illinois at Champaign-Urbana and completed his undergraduate studies in Civil Engineering in 2004. Upon graduation he was commissioned through the Air Force ROTC and entered AFIT's Graduate School of Engineering and Management. His next assignment will be at Misawa AB, Japan.



www.manaraa.com

	F	Form Approved OMB No. 074-0188							
The public reporting burden for this collection of information is estimated to average 1 hour per response, including the time for reviewing instructions, searching existing data sources, gathering and maintaining the data needed, and completing and reviewing the collection of information. Send comments regarding this burden estimate or any other aspect of the collection of information, including suggestions for reducing this burden to Department of Defense, Washington Headquarters Services, Directorate for Information Operations and Reports (0704-0188), 1215 Jefferson Davis Highway, Suite 1204, Arlington, VA 22202-4302. Respondents should be aware that notwithstanding any other provision of law, no person shall be subject to an penalty for failing to comply with a collection of information if it does not display a currently valid OMB control number. <b>PLEASE DO NOT RETURN YOUR FORM TO THE ABOVE ADDRESS.</b>									
1. REPOR 23-03-2000	RT DATE (DD-1 6	MM-YYYY)	2. REPORT TYPE Master's Thesis			<b>3. DATES COVERED</b> ( <i>From</i> – <i>To</i> ) Jan 2005 – Mar 2006			
4. TITL	E AND SUBTI	TLE			5a.	5a. CONTRACT NUMBER			
Evalu	ation Of Chlo	prinated Solv	ents Removal Efficiency Among Three Wetlan		Wetland <b>5b</b> .	d 5b. GRANT NUMBER			
		T lant Spc	cies. A mesocom stud	y	5c.	PROGRAM ELEMENT NUMBER			
6. AUT	HOR(S)				5d.	PROJECT NUMBER			
Jun Yan,	, $2^{nd}$ Lt, USA	F			5e.	TASK NUMBER			
					5f.	WORK UNIT NUMBER			
<ul> <li>7. PERFORMING ORGANIZATION NAMES(S) AND ADDRESS(S)         <ul> <li>Air Force Institute of Technology</li> <li>Graduate School of Engineering and Management (AFIT/ENV)</li> <li>2950 Hobson Way</li> <li>WPAFB OH 45433-7765</li> </ul> </li> <li>9. SPONSORING/MONITORING AGENCY NAME(S) AND ADDRESS(ES)</li> </ul>						8. PERFORMING ORGANIZATION REPORT NUMBER AFIT/GEM/ENV/06M-18 10. SPONSOR/MONITOR'S ACRONYM(S)			
						11. SPONSOR/MONITOR'S REPORT NUMBER(S)			
12. DISTRIBUTION/AVAILABILITY STATEMENT APPROVED FOR PUBLIC RELEASE; DISTRIBUTION UNLIMITED.									
13. SUPP	LEMENTARY	NOTES							
<b>14. ABSTRACT</b> Different species of plants need to be studied individually to compare the remediation efficiency of each species. This research will study three different wetland plants species and an unplanted control, under a laboratory setting. Each plant has a different characteristic favorable for chlorinated solvent degradation. <i>Eleocharis erythropoda</i> (Spike Rush) are plants with thin tube like leaves and large root mass. <i>Carex comosa</i> (Bearded Sedge) has broad leaves and <i>Scirpus atrovirens</i> (Green Bulrush) are broad leafed wetland plants with a long flowering stem during reproduction. PCE will be injected into the plant mesocosm and any possible PCE degradation will be observed. It is my hypothesis that PCE will be degraded into daughter products in all the mesocosms. However, the question will be which plant is the most efficient at chlorinated solvent degradation and is there difference between the planted reactors and the control reactors?									
15. SUBJECT TERMS Phytoremediation, bioremediation, PCE, TCE, chlorinated solvents									
16. SECURITY CLASSIFICATION OF:			17. LIMITATION OF ABSTRACT	18. NUMBER OF	<b>19a. NAME OF RESPONSIBLE PERSON</b> Bleckmann, Charles A., Civilian (ENV)				
REPORT U	ABSTRACT U	c. THIS PAGE U	UU	122	<b>19b. TELEPHO</b> (937) 255-3636, ex	<b>NE NUMBER</b> (Include area code) (t 4721; e-mail: Charles.Bleckmann@afit.edu			

