

Common and heritable components of white matter microstructure predict cognitive function at 1 and 2 y

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Previous studies indicate that the microstructure of individual white matter (WM) tracts is related to cognitive function. More recent studies indicate that the microstructure of individual tracts is highly correlated and that a property common across WM is related to overall cognitive function in adults. However, little is known about whether these common WM properties exist in early childhood development or how they are related to cognitive development. In this study, we used diffusion tensor imaging (DTI) to investigate common underlying factors in 12 fiber tracts, their relationship with cognitive function, and their heritability in a longitudinal sample of healthy children at birth ($n = 535$), 1 y ($n = 322$), and 2 y ($n = 244$) of age. Our data show that, in neonates, there is a highly significant correlation between major WM tracts that decreases from birth to 2 y of age. Over the same period, the factor structure increases in complexity, from one factor at birth to three factors at age 2 y, which explain 50% of variance. The identified common factors of DTI metrics in each age group are significantly correlated with general cognitive scores and predict cognitive ability in later childhood. These factors are moderately heritable. These findings illustrate the anatomical differentiation of WM fiber from birth to 2 y of age that correlate with cognitive development. Our results also suggest that the common factor approach is an informative way to study WM development and its relationship with cognition and is a useful approach for future imaging genetic studies.

factor analysis | white matter | DTI | cognition | heritability

Individual differences in white matter (WM) microstructure have been shown to be related to cognitive functions in adults (1, 2) and children (3–5). Diffusion tensor imaging (DTI) parameters in different WM regions or tracts are variably associated with processing speed, general intelligence, and higher-order cognitive abilities (1–5).

To date, the majority of DTI studies of WM microstructure have analyzed each tract or WM region independently. However, more recent studies clearly demonstrate that fractional anisotropy (FA) and other DTI metrics of WM microstructure correlate highly across WM tracts and across a wide age range, including neonates, older children, and adults (6–8). Furthermore, studies indicate that a general factor derived by principal component analyses (PCA) describing common variance in FA across WM tracts is associated with information processing speed (9) as well as general intelligence (10), at least in older adults. In children ages 8 to 16 y, three principal components of FA have been identified; the one corresponding to the corpus callosum was significantly related to IQ (11).

The identification of one or more common WM factors that influence variation in multiple WM tracts or regions is important for several reasons. Such factors could prove to be a good target for imaging genetic studies, as one would expect genetic variants influencing WM development and function to act on WM throughout the brain. Describing WM tracts in terms of their common underlying dimensions may also provide invaluable knowledge on how they are functionally linked. In addition, there is a practical value to combining and presenting complex

microstructural data of every different fiber as a small number of summary scores, because they could be readily applied in clinical settings and easy to analyze in research studies. Finally, an understanding of how these common factors develop will be critical for understanding how WM tracts develop in relation to genetic, environmental, and functional influences.

Early childhood is a time of rapid WM development, with a significant increase of myelination (12) and of FA and a decrease of radial diffusivity (RD) and axial diffusivity (AD) during the first 2 y of life (13, 14). Two recent studies found that myelination, as measured by myelin water fraction (MWF), is predictive of general cognitive ability in young children (15, 16). However, little is known about how intertract microstructural covariance develops in this period of rapid growth, and how it is related to cognitive development. Furthermore, although studies have shown that regional or tract-based WM microstructure is heritable in children and adults (17), there have been no previous studies of the genetic and environmental contributions to the common factors that underlie WM integrity and development. Therefore, this study reports the covariance of tract-averaged DTI values across WM tracts, their relationship with cognitive functions, and their heritability in healthy children with longitudinal data collected at birth and ages 1 and 2 y.

Results

Changes in Intertract Correlations from Neonates to 2-y-Olds. We studied the correlation of tract-averaged DTI metrics between each WM tract across all subjects at each age. For this, tracts of interest were identified in an atlas, each subject's images were registered to the atlas, and average AD, RD, and FA in each

Significance

The integrity or microstructure of white matter as determined by diffusion tensor imaging (DTI) is related to cognitive function. Most studies focus on individual tracts, even though the microstructure of white matter tracts throughout the brain is highly correlated. In older adults, a common property of white matter predicts cognitive function, though it is not known if common factors are present in early childhood development or how they relate to cognitive function. Here, we found that DTI-based common underlying factors that emerge at this age are significantly related to cognitive abilities from birth to age 2 y. These findings indicate that the functional specialization of cognition and the anatomical differentiation of fibers cooccur in the neonatal and infant brain.

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tract in the atlas space was determined. There was a general decrease in intertract correlations for all DTI-derived metrics from birth to 2 y of age, represented by the cooler colors in correlation matrices of 2-y-olds (Fig. 1). At birth, RD had the highest mean correlation among tracts, followed by FA and AD. This pattern was also observed at ages 1 and 2, but the correlations were not as strong at these older ages. Interestingly, each DTI parameter had a different developmental pattern; correlational coefficients of AD dropped extensively from birth to 1 y of age to the level observed at age 2, and those of RD and FA showed a more gradual reduction from birth to 2 y. Homologous pathways of the left and right cerebral hemispheres had higher correlation coefficients for all DTI parameters than did non-homologous ones. Even though correlations of other nonhomologous tracts generally decreased, homologous tracts maintained fairly high correlations over time.

In neonates, the splenium showed relatively low correlations with all other tracts across all DTI parameters (Table S1). In addition, AD and FA in the cingulate cingulum (CGC) and AD in the inferior longitudinal fasciculus (ILF) also had relatively poor correlations with the other tracts. At age 1 y, the genu and the splenium had high correlations with the corticothalamic tract (CT) and the ILF, respectively, across DTI parameters (Table S2). At age 2 y, some of these nonhomologous correlation coefficients were moderately high (Table S3): right CT–genu ($\rho = 0.59$) for AD; right CT–genu ($\rho = 0.62$), right uncinate fasciculus (UNC)–genu ($\rho = 0.63$), and both ILF–splenium ($\rho = 0.66$) for RD; and right CT–genu ($\rho = 0.63$) for FA. Note that correlation patterns are very similar when one twin from each twin pair is excluded from the analysis across age groups (Tables S1–S3).

To help ensure that the different numbers of gradient directions used for scans in this analysis did not contribute to the observed correlations, we analyzed intertract DTI correlations in a subgroup ($n = 34$) with all three time points using the same gradient directions (Table S4). In this subgroup, intertract correlation from birth to 2 y decreased in the same pattern with higher correlations of RD compared with AD and FA in a given age group, although many correlations were not significant due to the small sample size and increased variability of DTI parameters at ages of 1 and 2 y.

Factor Analysis. Fig. 2 depicts extracted factors and their factor loading across 12 tracts using the principal axis factoring (also see Fig. S1 and Table S5). In neonates, factor analyses demonstrated a one-factor solution, with the first factor explaining 72%, 69%, and 80% of the total variance of FA, AD, and RD, respectively. Factor loadings for individual tracts were evenly high ranging from 0.66 to 0.95, with the splenium having relatively low loadings across DTI metrics.

At age 1 y, a single factor best describes FA and RD, explaining 47% and 56% of the total variance, respectively. Although the percentage of explained common variance and factor loading decreased somewhat, general patterns of factor loading were similar to that of neonates. In contrast, two factors were identified for AD, with the first two factors explaining 46%: F1 with high loading for genu, ILF, and prefrontal CT, and F2 with high loading for CGC, arcuate (ARC), and UNC.

At age 2 y, three factors were extracted for each DTI parameter. Three factors together explained 54% of total common variance of FA, 52% of AD, and 62% of RD. The overall pattern of factor loading was consistent for FA, AD, and RD. The ILF,

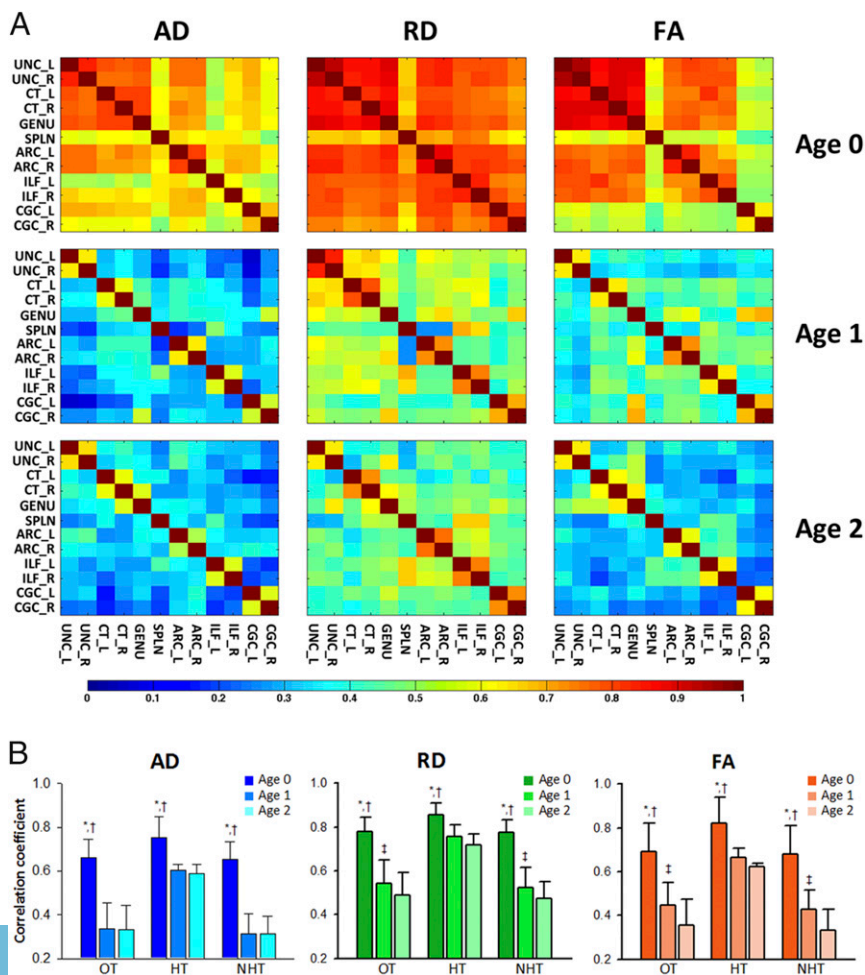


Fig. 1. (A) Heat maps of the intertract correlation matrices obtained from tract-level AD, RD, and FA and (B) mean correlation coefficients of overall (OT), homologous (HT), and nonhomologous tracts (NHT). * $P < 0.05$ compared with a group of 1-y-olds; †, $P < 0.05$ compared with a group of 2-y-olds; ‡, $P < 0.05$ compared with a group of 2-y-olds. Bilateral cingulate cinguli, CGC; callosal genu, GENU; callosal splenium, SPLN; uncinate, UNC.

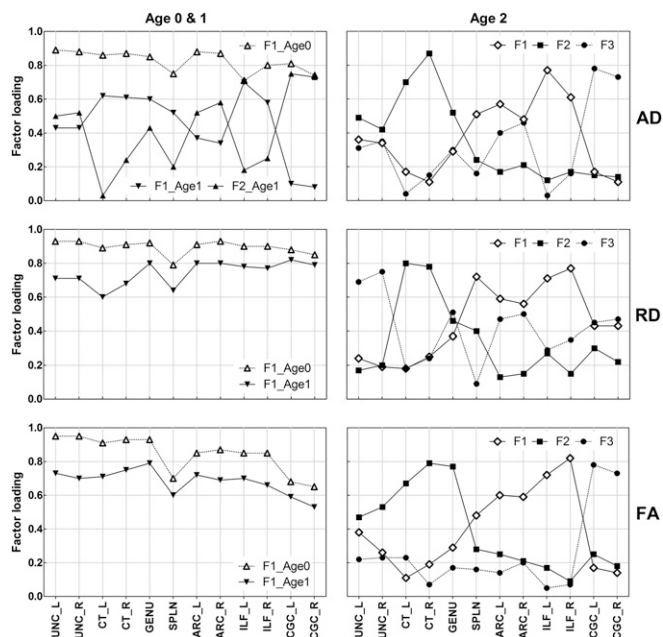


Fig. 2. Factor loadings of 12 tracts on extracted factors in three age groups. Refer to Fig. 1 for abbreviations.

the ARC, and the splenium had high factor loadings for factor 1, the genu and the prefrontal projection of CT had high loading for factor 2, and the CGC and the ARC had high loading for factor 3. Interestingly, AD factor 3 at age 2 y corresponded to AD factor 2 at age 1 y. All individual tracts had a positive weight on the factor across age groups, whereas the weight was near zero for some tracts. In general, the degree of factor loading was similar for homologous tracts in each hemisphere.

The similar factor structures were also found in two subgroups to which only one member of each of the twin pairs was randomly allotted across age groups (Fig. S2). Comparison of factor in the 6- and the 42-direction data found many similar factors with some differences; the 6- and 42-direction factors were related, suggesting differences were due to the small sample size of 6-direction data (Fig. S3 and Dataset S1).

Cognitive Relationships. There were several significant relationships between DTI parameter factors and Mullen early learning composite (ELC) scores (Table 1 and Fig. S4), including significant correlations between AD factor 1 in neonates and ECL at 1 y, AD factor 1 at 1 y and ECL at 2 y, and AD factor 2 at 1 y and ECL at 2 y, indicating that AD factors have a predictive ability for cognitive function at later ages. Interestingly, AD factors 1 or 2 at 1 y did not have any significant relationships with ECL at 1 y, whereas both FA factor 1 and RD factor 1 were significantly associated with ECL at 1 y, but not at other ages, indicating that DTI factor-cognitive function relationships are age-specific. Supplementary Spearman's correlation indicated that FA had a positive relationship whereas RD had a negative relationship with ELC at age 1 y. Neonatal AD factor 1 had negative relationship with ELC at 1 y, whereas AD factor 1 at ages 1 and 2 y showed positive relationships with ELC at age 2 y (Table S6).

Secondary analysis of distance correlations with verbal and nonverbal developmental quotients (VDQ and NVDQ, respectively) revealed several significant relationships of interest not captured by correlations with ECL (Table S7). Neonatal factor 1 for FA, AD, and RD each was significantly correlated with NVDQ at age 1 y, and neonatal AD factor 1 was significantly correlated with VDQ at age 2 y, providing more evidence of the predictive ability of DTI factors for later cognitive function. FA factor 3 at 2 y was also significantly correlated with VDQ at age 2 y. Analysis of the Mullen subscales revealed several significant

correlations consistent with correlations identified with ECL, VDQ, and NVDQ (Table S7).

These results were generally replicated in subgroups in which only one member of each of the twin pairs was assigned (Tables S8 and S9).

Heritability Analysis. Genetic contributions to DTI factors are presented in Table 2. In neonates, the single factor of AD, RD, and FA had relatively little genetic variance (0.25 to 0.29), whereas 1-y-olds had higher heritability estimates (0.38 to 0.74). Genetic contributions were somewhat lower and more variable at age 2 y (0.20 to 0.53) given the complex factor structure present at age 2 y. These differences are likely nonsignificant; statistical tests involve multivariate longitudinal analyses, which are beyond the scope of this article.

Discussion

We have shown that there are highly significant correlations in measures of microstructure between major WM tracts in neonates and that these correlations tend to decrease after birth while the factor structure develops increased complexity by age 2 y. The factors that describe common WM microstructure are significantly related to cognitive development in the first years of life, and these relationships are age-specific. We also found that these common WM microstructure factors are moderately heritable.

Factor Structure of Microstructural WM Parameters. This study investigates the factor structure of DTI parameters. We believe that explorative factor analysis (EFA) is more suitable than the PCA that previous studies have used because the main purpose of this study is not merely to reduce data but to identify unobservable latent variables that account for the shared variances (covariance) among variables that incorporate error of measurement (18).

In the current study, factors were extracted based on statistically significant correlations of WM microstructure between fibers. The biological underpinning of the factors is fundamentally the basis of diffusion properties in WM, effect of longitudinally oriented axonal membranes, the coherence of the parallel organization of axons within fibers, the packing density of the fibers, and their degree of myelination (19). Functional relatedness of fibers (7, 20) as well as technical causes such as intravoxel partial volume averaging (21) have also been proposed as possible explanations of correlation.

Table 1. Distance correlation coefficients (dCor) between Mullen ELC and factors of DTI parameters

Group	DTI factor	ELC at age 1 y		ELC at age 2 y	
		dCor	FDR-p	dCor	FDR-p
Age 0	FA_F1	0.10	0.45	0.12	0.24
	AD_F1	0.13	0.02	0.12	0.35
	RD_F1	0.11	0.24	0.12	0.24
Age 1 y	FA_F1	0.20	$<10^{-5}$	0.10	0.71
	AD_F1	0.11	0.446	0.20	$<10^{-4}$
	AD_F2	0.10	0.745	0.09	0.78
Age 2 y	RD_F1	0.20	$<10^{-4}$	0.12	0.45
	FA_F1	–	–	0.12	0.56
	FA_F2	–	–	0.12	0.52
	FA_F3	–	–	0.13	0.45
	AD_F1	–	–	0.19	0.001
	AD_F2	–	–	0.11	0.71
	AD_F3	–	–	0.16	0.17
	RD_F1	–	–	0.11	0.63
	RD_F2	–	–	0.10	0.39
RD_F3	–	–	0.13	0.78	

FDR-p, false discovery rate *P* value. Boldface *P*-values indicate significance level <0.05 .

Table 2. Variance component estimates of each factor for DTI parameters

Group	DTI factor	Variance components		
		a ²	c ²	e ²
Age 0	FA_F1	0.29****	0.00	0.06****
	AD_F1	0.25***	0.02	0.07****
	RD_F1	0.25****	0.03	0.04****
Age 1	FA_F1	0.55**	0.00	0.24***
	AD_F1	0.38**	0.06	0.12****
	AD_F2	0.55*	0.02	0.23****
Age 2	RD_F1	0.74****	0.00	0.12****
	FA_F1	0.49***	0.00	0.12****
	FA_F2	0.53*	0.00	0.21***
	FA_F3	0.46**	0.03	0.12****
	AD_F1	0.30	0.00	0.25
	AD_F2	0.51***	0.00	0.06****
	AD_F3	0.32	0.10	0.17****
	RD_F1	0.20	0.06	0.21*
	RD_F2	0.35***	0.10	0.07****
RD_F3	0.41**	0.00	0.16***	

*P < 0.05; **P < 0.01; ***P < 0.001; ****P < 0.0001. Values are unstandardized variance component estimates.

Factor analyses in this study indicated that the number of factors increases from one at birth to three at age 2. In neonates, one-factor structure with a single factor explained over 69% of the variance for AD, RD, and FA. At 1 y, FA and RD are best described by a one-factor structure, but the amount of the variance explained by the factor was somewhat decreased compared with this relationship at birth. At 1 y, AD is best described by two factors: one that includes high factor loading for the UNC, ARC, and CGC and much weaker loading for other tracts, and one that generally has moderate loading for all tracts except the CGC. The UNC–ARC–CGC factor persists at age 2 y; RD also “develops” this factor by age 2 y.

At age 2 y, three factors were extracted from all three DTI metrics. Fibers located in the posterior part of the cerebrum including the ILF, ARC, and splenium were dominant in factor 1 while the CGC and CT tracts were weakly loaded; fibers located in the anterior cerebrum including the CT and genu were dominant in factor 2. These findings are consistent with the notion that the global pattern of WM development follows a posterior–anterior gradient and thus the factors seem to reflect a shared maturation between fibers. Previous DTI studies have demonstrated that many of the same frontal fiber systems that make up factor 2 at age 2 y (e.g., uncinate, genu, corticospinal tract) mature later in development than more posterior fiber systems that make up factor 1 (e.g., ILF, splenium) (22–24). In addition, there are differences in other WM-related profiles such as diameter of axon and thickness of myelin sheaths between anterior and posterior regions of the cerebrum (25, 26). In parallel with this posterior–anterior theory of cortical maturation, the synchrony we observed in WM tracts is also reflective of the theory that cortical development is hierarchical, with sensorimotor regions maturing before regions that support higher-order processing (27). Factor loading was similar in each hemisphere for homologous tracts, suggesting that even though FA, RD, and AD tend to be somewhat asymmetric on a tract basis (14), the overall factor loading is similar across hemispheres. In terms of functional similarities, the genu connects left and right prefrontal cortex (7, 20) and explains prefrontal functions predominantly (28), whereas the splenium, ARC, and ILF are major tracts in which associations are consistently revealed between diffusion measures and language function (29, 30). Factor 3, the UNC–ARC–CGC dominant factor present at age 2 y, was already apparent at age 1 y for AD, suggesting that AD may be a more sensitive marker of differences in early maturation compared

with RD and FA. The three dominant fibers in this somewhat unexpected factor 3 are all curvilinear and later maturing association fibers connecting cortical regions within hemispheres. Because fibers with a higher degree of orientation dispersion reduce the FA due to partial volume effects (31), these three fibers may have relatively low FA in common (14) and cluster together. However, note that labeling a factor runs the risk of reifying a statistical finding, and further studies are needed to investigate the functional implication of these factors.

At this time, it is not clear how the factor structure of common WM microstructure develops between age 2 y and adulthood. A previous PCA study identified one FA component in older adults (9). Although it is difficult to compare these studies directly, it may be that WM tracts differentiate into several complex factors during childhood development but, in adulthood at the endpoint of development, converge to make a single universal factor again. This assumption may corroborate the canalization effect, in which cumulative genetic effects determine the endpoint of development (32).

Cognitive Relationships. This study found that the common factor for AD, RD, and FA at birth was significantly correlated with NVDO at age 1 y, and neonatal AD was also correlated with ECL at age 1 y and VDO at age 2 y. As such, neonatal WM microstructure at birth is somewhat predictive of cognitive function at later ages. AD factor 1 in 1-y-olds was predictive of ECL at age 2 y. It is interesting that the common FA and RD factor at age 1 y was significantly related to ELC at age 1 y whereas there was no relationship of any of the FA or RD factors with ECL at 2 y. In contrast, the AD factors at age 1 y are not related to cognitive function at age 1 y, whereas AD factor 1 (with high loading from ILF, ARC, and splenium) at ages 1 and 2 y is related to ECL at age 2 y. At ages 1 and 2 y, less maturity (higher AD) in these association fibers is associated with better cognitive development. In contrast, the negative correlation of neonatal AD factor 1 with ECL at 1 y indicates that relatively more mature (lower AD) fibers at birth are associated with better cognitive development at 1 y. These results indicate that the relationship between WM microstructure and cognitive development is age-specific in early childhood.

In the developing brain, AD is thought to reflect overall fiber density and complexity whereas RD is more reflective of myelination (12). These measures are more proximal to the neuro-anatomical processes (i.e., axonal pruning and change of intracellular cytoskeleton during early development) than FA (22), and FA and RD change more rapidly from 1 to 2 y compared with AD (14). AD may be a better representative of WM fibers as a whole because of these properties (12). AD at birth was predictive of ECL at age 1 y and of VDO at age 2 y, and AD factor 1 at age 1 y was predictive of ECL at age 2 y, suggesting that overall WM complexity and integrity is more highly correlated with later cognitive development. There is rapid development of myelin and a decrease in RD (13, 14) in the first year of life, and the relationship of the RD factor at 1 y with ELC at 1 y may reflect the role that myelination plays in cognitive development in the first year of life, although the role of myelination (at least as much as it is reflected in RD) is not as strong at age 2 y, perhaps due to a ceiling effect. This may be because the bulk of myelination has occurred by age 1 y and individual variation in cognitive development in the second year of life is less dependent on further maturation of myelin and more dependent on inherent structure reflected in AD. This notion is supported by a previous study that found only a very weak association of MWF, a myelin-related imaging parameter, with Mullen cognitive scores in young children (16). In addition, a similar study found MWF had more widespread correlations with ECL in 1- to 2-y-olds compared with 2- to 4.8-y-olds (15).

FA and cognitive function relationships have been studied mainly in adults and older children (3, 33), and some studies find lower RD in association with better cognition as an index of myelination (4, 5). However, few studies have reported relationships between AD and cognition (1). AD appears to be more related to cognitive development in early childhood than are RD or FA (12).

Overall, the correlations between the DTI factors and cognitive functions were weak and ranged from 0.13 to 0.23. These correlations are, in fact, very similar to those found by the previous studies of older adults using a similar approach (9, 10), suggesting that the magnitude of the relationship between WM microstructure and overall cognitive function may be consistent throughout life. The tract-averaged analysis performed in this study may not be the best way to uncover structure–function relationships in WM, as specific regions within tracts may have stronger relationships with behavior and cognitive function.

Heritability of Identified Factors. We observed that single factors of FA, AD, and RD in neonates had significant heritability ranging from 0.24 to 0.28. These findings are consistent with a mean heritability of around 0.30 in individual tracts in neonates using the same quantitative tractography (34). Factor heritability demonstrated a moderate increase at age 1 y—0.37 to 0.71—and then a mild decrease to 0.20 to 0.52 at age 2 y as the three factors were divided, with genetic estimates of factors at age 2 y still higher than those in neonates.

On the other hand, the present study found that the common environment component for DTI factors was around zero across this age range, whereas the unique environment component showed considerable increase at age 1 y. These findings suggest that environmental effects shared by twin pairs are minimal, whereas those not shared by twin pairs increase with age; thus environment may influence the relative decrease of heritability from birth to 2 y of age.

As imaging genetic studies of WM indicated that several genes are associated with WM microstructure (35, 36), WM-related genes likely have effects across all WM fibers. These possibilities, in turn, suggest that factor analysis or PCA may be more informative for imaging genetics studies.

Microstructural Intertract Correlation. When two tracts have a positive correlation, a subject with a large FA value in one tract is likely to have a large FA in the other tract. A previous study found that adults had higher intertract correlations than neonates (6), although the trajectory between birth and adulthood is not clear. We found that, after initial decline, means of between-tract correlations in 2-y-olds reached 0.35 ± 0.12 for FA, 0.33 ± 0.11 for AD, and 0.49 ± 0.10 for RD in this study, lower than neonates, and also supported for adolescents (0.35 ± 0.18 for FA, 0.78 ± 0.06 for AD, and 0.75 ± 0.08 for RD) (6) and adults (0.43 ± 0.18 for FA, 0.41 ± 0.16 for AD, and 0.51 ± 0.13 for RD) (7). Healthy neonates are born with mostly unmyelinated WM, and myelination occurs primarily over the first few years of life. Hence, the correlations between tracts would be expected to be high at birth given that tracts would tend to be homogenous, with little myelination and differentiation, and lower in the early postnatal period as tract maturation and myelination occurs in response to individual experience and heterogeneous genetic and environmental influences on individual tracts (6). Taken together, we propose U-shaped trajectory of microstructural intertract correlation, with a decrease from birth to 2 y and then a gradual increase throughout adolescence and into adulthood. Longitudinal studies in children are necessary to confirm this.

This study has several limitations that need to be considered when interpreting the results. First, the use of different scanners with different gradient directions may introduce differences in DTI microstructural measurements (21, 37), although correlations between tracts in individuals would likely be less affected given they were determined at a single time point with the same sequence on the same scanner. In general, subjects across age groups showed similar factor structures between the 6- and the 42-direction data, with some differences that were likely caused by the small sample size of the 6-direction data. In addition, the results for intertract DTI correlations over time using a subgroup with all three time points using the same gradient directions were similar to the larger group. Second, attrition across age groups is another caveat that may affect results, including reduced power

to detect significant correlations at ages 1 and 2 y. In addition to loss of subjects over time due to dropouts, a further loss of analyzable scans in 1- and 2-y-olds was caused by failure to get the child to sleep and enter the scanner, which is more likely in older children, as well as motion once in the scanner, even when cognitive data were available for the subject. Finally, our analysis study was focused on correlations of tract averages, mainly to be comparable to previous studies in adults that used this approach. An analysis of pointwise DTI metrics, namely tract profiles along each tract, may provide more precise spatial information about correlations and how they change over development (14).

In conclusion, our longitudinal study has demonstrated decreased intertract correlations, increased complexity of factor structure, and genetic heritability from birth to 2 y of age in terms of DTI parameters. Moreover, these common components of WM microstructure are associated with present and future cognitive ability and are age-specific. This suggests that studies of WM structure and cognitive ability need to be age-specific and that combining wide age ranges in development will miss age-specific relationships (30). Further studies at later ages in childhood are needed to understand how these relationships evolve. Finally, this study also suggests that the common factor approach is a fruitful way to study WM microstructure and the relationship to cognitive function in development and complements an individual-fiber-centered approach.

Materials and Methods

Participants. Children analyzed in this study are healthy singleton and twin subjects in the ongoing longitudinal University of North Carolina Early Brain Development Studies (EBDS). Pregnant women were recruited from the outpatient obstetrics–gynecology clinics at the University of North Carolina hospitals and Duke University Medical Center. Exclusion criteria for mothers included major maternal illness or infection during pregnancy, and maternal diagnosis of a major psychiatric disorder; for neonates, exclusion criteria included chromosomal abnormalities, severe congenital abnormalities, major medical illness or infection, and abnormalities on MRI other than small intracranial hemorrhages that are common in neonates (38). Zygosity was determined with PCR short tandem repeat analysis of 14 loci on DNA prepared from buccal swab cell collection (BRT Laboratories). Written informed consent was obtained from a parent of all infant participants. This study was approved by the Institutional Review Boards of the University of North Carolina School of Medicine and Duke University Medical Center.

DTI and cognitive data from a total of 685 subjects were collected across three age groups (Dataset S2) including 429 twin (173 monozygotic, 256 dizygotic) and 256 nontwin subjects (male/female: 366/319, Caucasian/African American/Asian: 525/146/14). There were 535 subjects with analyzable DTI scans at birth, 322 at 1 y, and 244 at 2 y; among 322 subjects with usable scans at age 1 y, 217 (67%) were included at birth; of the 244 with usable scans at age 2 y, 164 (67%) were included at birth and 134 (55%) were included at age 1 y. Demographic variables for each group are presented in Table S10 and Fig. S5. Data from these participants have been published partially in our previous studies (14, 34).

Image Acquisition. Most MRI data were acquired on a 3T Siemens Allegra head-only scanner (Siemens Medical System), comprising 86% of the neonates, 82% of the 1-y-olds, and 77% of the 2-y-olds in the present study. For earlier Allegra diffusion-weighted imaging (DWI) data, a single-shot echo-planar imaging spin-echo sequence was used with the following parameters: Repetition Time (TR)/Echo Time (TE) = 5,200/73 ms, slice thickness = 2 mm, and in-plane resolution = $2 \times 2 \text{ mm}^2$, with a total of 45 slices for six directions using b value of $1,000 \text{ s/mm}^2$ and 1 baseline image (b value = 0) per sequence, repeated five times total to improve signal-to-noise. For the remaining Allegra DWI data, 42 directions of diffusion sensitization were acquired with a b value of $1,000 \text{ s/mm}^2$ in addition to seven images with no diffusion weighting for reference. The parameters were as follows: TR/TE/Flip angle = 7,680/82/90°, slice thickness = 2 mm, and in-plane resolution = $2 \times 2 \text{ mm}^2$, with a total of 60 to 72 slices. The rest of the study subjects were scanned using an upgraded Siemens model, the 3T Tim Trio (Siemens Medical System), following the same sequencing parameters as the 42-direction Allegra sequence detailed above.

Quantitative Tractography. A study-specific quality control protocol was applied to all raw DWI data using DTIPrep (www.nitrc.org/projects/dtiprep), which included slice-wise and gradient-wise artifact detection, as well as eddy current and motion correction (39). Skull and nonbrain tissue was masked out using

Brain Extraction Tool (BET) (40), and tensors were estimated using a weighted least-squares algorithm (41). Using the University of North Carolina –Utah National Alliance for Medical Image Computing DTI framework (www.nitrc.org/projects/dtiatlasbuilder, 42), the University of North Carolina EBDS combined neonate and pediatric (1 to 2 y) atlas was created, on which a total of 12 fiber tract segments consistent with those of Penke et al. (12) were reconstructed using streamline tractography using 3D Slicer (<https://www.slicer.org>) and defined as follows: callosal genu and splenium; bilateral cingulate cinguli; ARC; uncinate; ILF; and prefrontal projections of the CT (see ref. 14 and appendix of ref. 34 for details). Each subject's image was registered to the atlas. For group analysis, statistical diffusion profiles were generated for AD, RD, and FA at equally spaced points along the length of each fiber tract in each subject's original DTI space. Note that the tractography can be considered to identify regions of interest along the tract, as crossing fibers from other tracts may be contributing to the results. Average AD, RD, and FA values were computed over a given fiber segment for each tract. These generated tract-averaged DTI parameters were used in the following analysis.

Cognitive Measure: Mullen Scales of Early Learning. The Mullen Scales of Early Learning (MSEL) is a standardized developmental test for children ages 3 to 60 mo; it consists of five scales: gross motor, fine motor, visual reception, expressive language, and receptive language (43). The MSEL was administered to all participants by trained researchers using standardized protocols at ages 1 and 2 y.

The primary cognitive variable for the present study was the conventional Mullen ELC, which is a T score representing an average of the four scales: visual reception, fine motor, expressive language, and receptive language.

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- Mullen VQ, NVDQ, and raw scores composed of the five Mullen scales were also used as secondary variables (5).

Statistical Analysis. For demographic variables, frequency distributions were calculated for categorical variables, and the means and SDs were calculated for continuous variables.

Spearman's correlation analyses were used to study the intercorrelations of the tract averages of DTI parameters between WM tracts. Then, EFA for each tract average DTI parameter was conducted in the whole sample using the package “psych” (44) and “MVN” (45) in R (46). The relationships between extracted factor scores of the tract average DTI metrics and the cognitive scores were studied using distance correlation performed by the package “energy” in R (47). The Benjamini and Hochberg (1995) false discovery rate (FDR) adjustment was used to adjust *P* values for multiple comparisons (48).

For genetic analysis of twins, we fitted ACE (additive genetics, common environment, and unique environment) model to study the importance of environmental and genetic influences. The variance was decomposed into three components of ACE. Parameters were estimated using Full Information maximum Likelihood in the R package “OpenMx” (44). A full description of statistical analysis is provided in [Supporting Information](#).

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