

Corrections

APPLIED BIOLOGICAL SCIENCES

Correction for “In vivo fluorescence imaging of exogenous enzyme activity in the gastrointestinal tract,” by Gregor Fuhrmann and Jean-Christophe Leroux, which appeared in issue 22, May 31, 2011, of *Proc Natl Acad Sci USA* (108:9032–9037; first published online May 16, 2011; 10.1073/pnas.1100285108).

The authors note that, on page 9036, right column, line 5, “(0.1 µg)” should instead appear as “(1 µg).”

This error does not affect the conclusions of the article.

www.pnas.org/cgi/doi/10.1073/pnas.1216389109

DEVELOPMENTAL BIOLOGY

Correction for “Transcriptional network controlled by the trithorax-group gene *ash2* in *Drosophila melanogaster*,” by Sergi Beltran, Enrique Blanco, Florenci Serras, Beatriz Pérez-Villamil, Roderic Guigó, Spyros Artavanis-Tsakonas, and Montserrat Corominas, which appeared in issue 6, March 18, 2003, of *Proc Natl Acad Sci USA* (100:3293–3298; first published March 7, 2003; 10.1073/pnas.0538075100).

The authors note that there is a broken link in the legend of Fig. 4, at the end of *Materials and Methods*, in *Results and Discussion*, and in *SI Materials and Methods*. The link www.ub.es/epidd/arrays/index.htm should instead appear as <http://compfly.bio.ub.es/Ash2Microarrays>.

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PHARMACOLOGY

Correction for “Chemical and genetic evidence for the involvement of Wnt antagonist Dickkopf2 in regulation of glucose metabolism,” by Xiaofeng Li, Jufang Shan, Wochul Chang, Ingyu Kim, Ju Bao, Ho-Jin Lee, Xinxin Zhang, Varman T. Samuel, Gerald I. Shulman, Dakai Liu, Jie J. Zheng, and Dianqing Wu, which appeared in issue 28, July 10, 2012, of *Proc Natl Acad Sci USA* (109:11402–11407; first published June 25, 2012; 10.1073/pnas.1205015109).

The authors note that the affiliation for Jufang Shan, Ju Bao, Ho-Jin Lee, Xinxin Zhang, and Jie J. Zheng should instead appear as Department of Structural Biology, St. Jude Children’s Research Hospital, Memphis, TN 38105. The corrected affiliation line appears below. The online version has been corrected.

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In vivo fluorescence imaging of exogenous enzyme activity in the gastrointestinal tract

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Exogenous enzymes are administered orally to treat several diseases, such as pancreatic insufficiency and lactose intolerance. Due to the proteinaceous nature of enzymes, they are subject to inactivation and/or digestion in the gastrointestinal (GI) tract. Here we describe a convenient fluorescence-based assay to monitor the activity of therapeutic enzymes in real time in the GI tract. To establish the proof of principle, the assay was applied to proline-specific endopeptidases (PEPs), a group of enzymes recently proposed as adjuvant therapy for celiac disease (a highly prevalent immunogenetic enteropathy). A short PEP-specific peptide sequence which is part of larger immunotoxic sequences of gluten was labeled with a fluorescent dye and a corresponding quencher. Upon enzymatic cleavage, the fluorescence emission was dequenched and detected with an in vivo imaging system. PEPs originating from *Flavobacterium meningosepticum* (FM) and *Myxococcus xanthus* (MX) were evaluated after oral administration in rats. While MX PEP could not cleave the peptide in the stomach, FM PEP showed significant gastric activity reaching 40–60% of the maximal in vivo signal intensity. However, both enzymes produced comparable fluorescence signals in the small intestine. Coadministration of an antacid drug significantly enhanced MX PEP's gastric activity due to increased pH and/or inhibition of stomach proteases. With this simple procedure, differences in the in vivo performance of PEPs, which could not be identified under in vitro conditions, were detected. This imaging assay could be used to study other oral enzymes in vivo and therefore be instrumental in improving their therapeutic efficiency.

celiac sprue | prolyl oligopeptidase | autoimmune disease

Exogenous enzymes are being tested or already used orally to treat several diseases, including pancreatic insufficiency, lactose intolerance, and phenylketonuria (1, 2). However, harsh pH changes and the presence of proteases in the gastrointestinal (GI) environment can strongly alter the activity of orally-administered enzymes. Strategies to prevent GI degradation of enzymes include enteric-coated formulations (3), coadministration of H₂-receptor antagonists (4), and polymer-based matrices (5, 6). However, to the best of our knowledge, convenient methods to evaluate oral enzyme activity in vivo are still lacking. Here, we present a method for the real time measurement of exogenous enzyme activity in the GI tract by fluorescence dequenching using celiac disease (CD) as model application. CD is a highly prevalent (~1%), genetically-based illness of the gut, triggered by the ingestion of cereal proteins, such as wheat gluten (7).

Generally, the digestive process of proteins in mammals is initiated in the stomach by low pH and proteases (mainly pepsin). Digestion is pursued in the small intestine (pH 5–7) by pancreatic proteases (e.g., trypsin and chymotrypsin) and brush-border membrane enzymes, leading to amino acids and oligopeptides which are then absorbed (8, 9). Owing to their exceptionally high proline and glutamine content, immunogenic peptides from gluten resist enzymatic degradation by GI enzymes (10).

The current and only treatment for CD is life-long elimination of gluten from the diet (11). This dietary restriction is a difficult experience for many patients and is often associated with decreased quality of life. Poor compliance, whether inadvertent

or voluntary, to a strict gluten-free diet is frequent and predisposes patients to CD complications (e.g., nutritional deficiencies, osteoporosis, secondary autoimmune disorders, malignancies) (12). Hence, there is an urgent need for complementary nondietary therapies to help treat this common disorder because it is associated with increased morbidity and mortality (13). Various therapeutic avenues are currently being explored to manage this pathology (14–17). Among these, oral administration of prolyl endopeptidases (PEPs) is one of the most studied approaches (18). Unlike human GI enzymes, PEPs can efficiently hydrolyze proline-rich gluten peptides (9) and they have been shown to alleviate gluten immunotoxicity under simulated intestinal conditions (19, 20). Earlier studies conducted with PEPs from *Flavobacterium meningosepticum* (FM) and *Myxococcus xanthus* (MX) revealed moderate enzyme stability in vitro under artificial GI conditions (21). *Aspergillus niger* (AN) PEP was recently investigated in an artificial compartmental system mimicking the human GI tract (TIM model). This PEP efficiently accelerated gluten breakdown under gastric conditions in vitro (22). Another PEP from *Sphingomonas capsulate* (SC) was evaluated in a clinical trial in a combinatory approach (23). The enzyme reduced the gluten-specific T cell response but did not significantly diminish CD-associated GI symptoms (24). Thus, a better understanding of the behavior of PEPs in vivo after oral administration may pave the way for a more efficient optimization of CD treatment (25). In this contribution, we report an in vivo fluorescence-based assay that allows for monitoring enzymatic PEP activity in real time in the GI tract. This assay could, in principle, be adapted to other oral enzymes.

Results and Discussion

In Vitro Activity of PEPs. The activity of FM, MX, and SC PEPs was first characterized in vitro and their low stability in artificial GI fluids (Fig. S1A) was confirmed (21). Moreover, PEPs could not reduce the abundance of the peptide QLQPFQPOLPYQPQPF from whole wheat gluten under conditions simulating in vitro GI digestion (Fig. S1B). This model peptide comprises the major immunogenic peptide QLQPFQPOLPY (26, 27). Since the in vivo predictability of in vitro studies is often limited, we sought to analyze PEP activity in rats.

Design of In Vivo Fluorescence Assay. To develop the fluorescence assay, the short peptide sequence LPYQPF was selected as substrate as it constitutes the core region of both the CD-toxic peptide p62-75 from α 2-gliadin (21) and from QLQPFQPOLPYQPQPF (Fig. 1A). To establish proof of concept, LPYQPF was first labeled with the fluorescent dye EDANS ($\lambda_{em} = 490$ nm) and corresponding quencher Dabcyl [E(EDANS)-

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The authors declare no conflict of interest.

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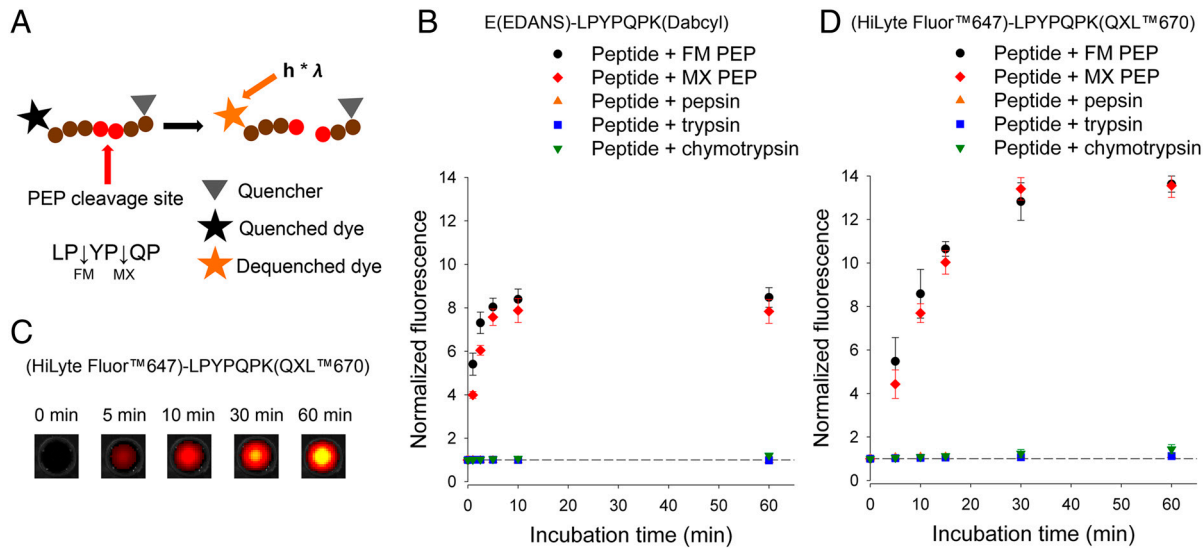


Fig. 1. In vitro cleavage of E(EDANS)-LPYPQPK(Dabcyl) and (HiLyte Fluor™647)-LPYPQPK(QXL™670). (A) Schematic representation of the dequenching assay. Arrows within the peptide sequence indicate preferential cleavage sites for FM and MX PEP. (C) In vitro cleavage of (HiLyte Fluor™647)-LPYPQPK(QXL™670) by FM PEP in phosphate buffer (pH 6.8) measured with the in vivo imaging system. (B, D) The peptides (5 and 3 μ M, respectively) were incubated in the presence of FM and MX PEPs (1 μ g/mL), pepsin (0.20 mg/mL), trypsin (0.22 mg/mL), or chymotrypsin (0.22 mg/mL). Mean \pm SD, $n = 3$, fluorescence intensities in B and D were measured using plate reader and in vivo imaging system, respectively.

LPYPQPK(Dabcyl)]. Fluorescence of the intact peptide was low while its proteolysis by PEPs in vitro led to its complete cleavage [as verified by liquid chromatography-mass spectrometry (LC-MS)] concurrent with an 8-fold increase in fluorescence intensity after 15 min incubation. Native enzymes of the GI tract barely hydrolyzed this peptide (Fig. 1B). For the in vivo experiments, in order to increase signal depth penetration and to reduce auto fluorescence within tissue (28), LPYPQPK was labeled with the fluorescent dye HiLyte Fluor™647 ($\lambda_{em} = 700$ nm) and corresponding quencher QXL™670 [(HiLyte Fluor™647)-LPYPQPK (QXL™670)]. In vitro proteolysis of this modified peptide resulted in a >13-fold increase of fluorescence after complete cleavage, while luminal proteases could not digest the peptide (Fig. 1C and D). A control peptide without quencher [(HiLyte Fluor™647)-LPYPQPK] produced comparable fluorescence intensities in vitro (Fig. S2). LC-MS analysis additionally revealed that FM PEP mainly cleaved at LPYP \downarrow QP, while MX PEP preferred the LP \downarrow YPQP position (Fig. 1A).

In Vivo Dequenching Assay. With this tool in hand, the enzymatic activity of PEPs was examined in vivo in female rats (see Movie S1 for a live illustration of the in vivo assay). FM and MX PEP were selected for the in vivo experiments because of their comparable in vitro cleavage pattern against gliadin peptides with similar chain length (21). The animals were gavaged with PEPs, followed by the labeled peptide 5 min later. Whole wheat gliadin was coadministered to all rats to create a more realistic environment. To quantify the extent of peptide cleavage in vivo, the fully cleaved peptide (precleaved peptide) was also given to rats in a separate experiment, and its fluorescence intensity taken as 100%. After gavage, the animals were either quickly anesthetized continuously for 4 h or at selected time points for 15 min. For imaging, the rats were placed in an in vivo imaging system (Fig. S3). Images were analyzed by selecting regions of interest (ROIs) over the stomach and small intestine based on the rat's anatomy (Fig. S4). In terms of size, form, and position on the animal's abdominal area, ROIs were kept constant, for a given animal, throughout the duration of experiments. The fluorescence signal was not tracked in the lower GI tract because inflammation in CD occurs in the early small intestine.

In Vivo Imaging under Continuous Anesthesia. For stomach analysis experiments, rats were anesthetized 5 min after gavage. By doing so, gastric motility decreased, and the effect of prolonged exposure to gastric conditions was examined. Administration of the peptide or PEPs alone did not result in significant fluorescence in the stomach (Fig. 2, Fig. S5), while the precleaved peptide produced a strong fluorescent signal. Rats receiving FM PEP prior to the peptide displayed a rapid increase in fluorescence intensity in the stomach region (Fig. S5), reaching 60% of the signal of the precleaved peptide after 20 min (Fig. 2B). Conversely, MX PEP

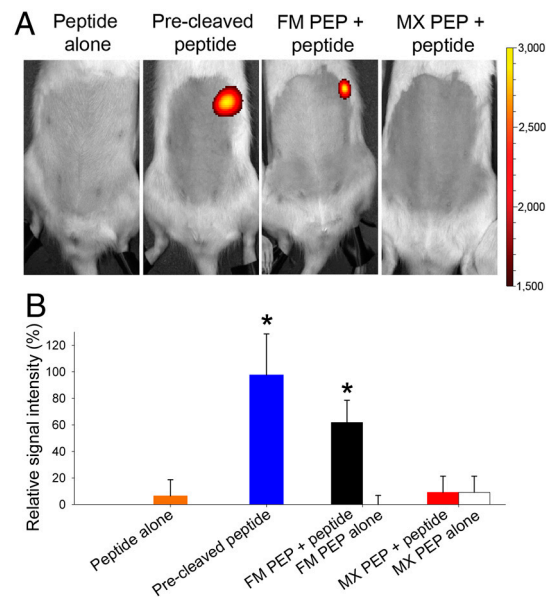


Fig. 2. Real time measurement of PEP activity in vivo under continuous anesthesia. (A) Fluorescence imaging and (B) relative signal intensity in the stomach region 20 min after oral administration of the labeled peptide alone, the precleaved peptide, FM, or MX PEP with or without peptide. Representative rats from each set are shown in A. Color scales are identical for all pictures. In B, signals were plotted by setting the average in vivo maximum signal (precleaved peptide after 18 min) to 100%. Mean \pm SD, $n = 6$. *Precleaved peptide and FM PEP + peptide vs. all other groups ($p < 0.05$).

The presented assay could also serve to monitor the degradation of peptide drugs by GI enzymes *in vivo* and devise new delivery strategies. The work highlighted significant discrepancies between the *in vitro* and *in vivo* properties of oral enzymes. In particular, this imaging method (potentially in combination with the TIM model) could prove useful in the optimization of PEPs and their delivery strategies for the treatment of CD.

Materials and Methods

Materials. FM PEP (25 U/mg) was purchased from LuBioScience, MX PEP (25 U/mg) was obtained from Zedira, and SC PEP (19 U/mg) was custom-made by Prozomix Ltd., as described elsewhere (34). The labeled peptides (EDANS)-LPYPQPK(DabcyI), (HiLyte Fluor™647)-LPYPQPK(QXL™670) and (HiLyte Fluor™647)-LPYPQPK were custom-synthesized by Peptide 2.0 and Anaspec, respectively. Whole wheat gluten, gliadin, pepsin (2,188 U/mg protein), α -chymotrypsin (96 U/mg protein), and pancreatin (USP grade) were purchased from Sigma-Aldrich. Z-Gly-Pro-pNA and trypsin (3,480 U/mg) were procured from Bachem and AppliChem, respectively. Magnesium aluminate monohydrate (magaldrate, Riopan®) was purchased in a retail pharmacy.

In Vitro Stability and Activity of PEPs. FM, MX, and SC PEPs were incubated at 50 mU/mL in simulated gastric (pH 1.2 with pepsin) and intestinal (pH 6.8 with pancreatin) fluids United States Pharmacopeia (USP), and in USP acetate buffer (pH 4.5 with pepsin). pH-dependent activity profiles of FM and MX PEPs were determined at 50 mU/mL in HCl/KCl solution USP (pH 1.2), acetate buffer USP (pH 4.5), phosphate buffer (100 mM, pH 7.0), and trometamol/sodium chloride buffer (50/100 mM, pH 9). Activity was assessed by measuring the cleavage of Z-Gly-Pro-pNA using an Infinite M200 microplate reader (Tecan Ltd.) (20).

In vitro activity of PEPs was analyzed by incubating the enzymes directly with whole wheat gluten in 50 mM acetate buffer USP (pH 4.5). Pepsin (0.8 mg/mL) was added, and the mixture was incubated for 1 h at 37 °C. Subsequently, the pH was raised to 6.8 (using 0.2 M Na₂HPO₄ and 0.2 M NaOH) and trypsin and chymotrypsin (0.4 mg/mL) were added. The mixture was again incubated for 2 h at 37 °C. After heat inactivation (10 min at 95 °C), the mixture was centrifuged (14,000 × *g* for 10 min) and the supernatant analyzed by LC-MS.

Samples were analyzed using an LC system composed of a Rheos Allegro quaternary pump, column oven (Hot Dog 5090), C18-column (Hypersil Gold, 100 × 1 mm, 1.9 μm) and XCalibur control software (all from Thermo Fisher Scientific Inc.). Samples were injected (10 μL) at 35 °C, and a flow rate of 50 μL/min using water (solvent A) and acetonitrile (solvent B) (both with 0.1% formic acid). The gradient consisted of: 95% solvent A as initial value, 95–50% A in 1–19 min, 50–5% A in 19–23 min, 5% A in 23–27 min, 5–95% A in 27–28 min, and 95% A in 28–31 min. The LC was directly connected to an LTQ XL linear quadrupole Ion Trap (Thermo Fisher Scientific Inc.) equipped with an electrospray ion source. Data were acquired by full MS, followed by MS²-fragmentation of the five most intense signals in an automated data-dependent scan.

Raw data were subjected to the Sequest search algorithm using Proteome Discoverer Software (Thermo Fisher Scientific Inc.). Detected peptides were searched against all sequences from the Expasy Proteomics database (Swiss Institute of Bioinformatics) that matched the following search terms: triticum, aestivum, wheat, and gliadin. Abundance of QLQFPQQLPYPQ PPF was analyzed for the incubation of PEPs with whole wheat gluten. Each value was normalized to a non-PEP-treated sample. The linearity of MS quantification was determined by spiking a synthetic gluten peptide (PQPQLPYPQQLP) into gluten-digest samples in a dose-dependent manner ($R^2 = 0.9945$).

Dequenching Assay. *In vitro* cleavage of the peptides E(EDANS)-LPYPQPK (DabcyI) ($\lambda_{ex} = 340$ nm, $\lambda_{em} = 490$ nm) and (HiLyte Fluor™647)-LPYPQPK (QXL™670) was monitored by spectrofluorometry ($\lambda_{ex} = 640$ nm, $\lambda_{em} = 700$ nm) at 37 °C on an Infinite M200 microplate reader and an IVIS® Spectrum *in vivo* imaging system (Caliper Life Science), respectively. Complete cleavage was confirmed by LC-MS.

In Vivo Imaging. All animal experiments were approved by the Cantonal Veterinary Office Zurich. *In vivo* experiments were conducted with (HiLyte Fluor™647)-LPYPQPK(QXL™670). Female Sprague-Dawley rats (4–8 weeks old, 125–200 g) on gluten-free food were fasted for 4 h, unless stated otherwise. The abdominal region of rats was shaved and a pre-gavage picture ta-

ken. The rats were gavaged orally with peptide alone (negative control), precleaved peptide (positive control), FM or MX PEP alone (negative control), FM or MX PEP, followed by peptide solution (peptide given 5 min after PEP; study group) (Fig. S3). Oral solutions were prepared as follows: FM and MX PEPs (0.1 μg) were diluted in 150 μL 10 mM phosphate buffer (pH 7.0). 5 μL of peptide stock solution (250 μM in 35% DMSO) was diluted in 200 μL sodium cholate [0.5%, w/volume (v)] to ensure proper solubilization. For the precleaved peptide solution, peptide and PEP were mixed and shaken at 37 °C for at least 4 h prior to each experiment. For all experiments, PEPs were admixed with gliadin, the dominant CD-toxic fraction in gluten (0.5 or 2.5 mg corresponding to a PEP:gliadin ratio of 1:500 or 1:2,500, w/w), 30 s prior to oral gavage. To estimate whether modified peptide was absorbed, rats were administered FM PEP with peptide and the fluorescence intensity in the saphenous vein area was imaged during 4 h of discontinuous anesthesia. In experiments aimed at assessing the impact of stomach acidity on MX PEP activity, the rats were given peptide diluted in magaldrate (Riopan®).

After oral administration, the rats were anesthetized (2% isoflurane, 2 mL/min oxygen) for immediate imaging (continuous anesthesia) or returned to their cages for imaging at later time points (discontinuous anesthesia) (Fig. S3). Rats were placed in an IVIS® Spectrum *in vivo* imaging system equipped with a cone nose for anesthesia (1.5–2% isoflurane, 0.5 mL/min oxygen) and a heated platform (37 °C) to maintain body temperature during analysis. Pictures were recorded in the following settings: $\lambda_{ex} = 640$ nm, $\lambda_{em} = 700$ nm, binning 8, f/stop 2, exposure time = 2 s. The rats were assigned to groups in such a way that mean weights were equal in all groups.

For quantifying residual peptide leaving the stomach and intestine, rats were administered FM PEP with peptide and they were killed after 1 h or the feces were collected 24 h after oral gavage, respectively. Feces or stomach content were suspended in a mixture (3:2, v/v) of phosphate buffer (10 mM, pH 7.0) and sodium cholate (0.5%) and centrifuged (3,000 × *g*, 5 min). 5 μL of FM PEP (0.25 mg/mL) was added to the supernatant and the extent of fluorescence dequenching (IVIS® Spectrum *in vivo* imaging system) was used to calculate the residual proportion of intact peptide. Then, the feces or stomach samples were extracted three times using sodium cholate dissolved in acetonitrile (0.25%, w/v). After centrifugation (20,000 × *g*, 10 min) supernatants were collected, combined and freeze-dried. Lyophilizates were dissolved in water and acetonitrile (1:1, v/v) and the concentration of cleaved peptide was quantified by spectrofluorometry.

Analysis of Recorded Images. Images were analyzed with Living Image software (Caliper Life Science). In each experiment, ROIs were defined for the stomach and small intestine (Fig. S4). At each time point, the average fluorescence intensity in each ROI was normalized to the pre-gavage status. The relative signal intensity was expressed in percentage of mean maximal *in vivo* value for each region (stomach and small intestine) obtained by administering the precleaved peptide (100%). The following equation was employed (normalized fluorescence of 1 is set to 0%)

$$\text{Relative signal intensity (\%)} = ((x_t - 1) \times 100\%) / (x_{\max} - 1),$$

where x_t represents normalized fluorescence at a single time point and x_{\max} is the maximal *in vivo* signal in the specific ROI.

Autofluorescence not related to the peptide and located outside the abdominal area was erased in all displayed pictures. All pictures were smoothed by 7 × 7 pixels to reduce background noise.

The video was created by editing the still pictures using Nero StartSmart software.

Statistical Analysis. Mean and standard deviation (SD) are reported for all data (unless stated otherwise); *n* values in the legends indicate the number of independent experiments. Group values were analyzed by nonparametric Kruskal-Wallis analysis of variance, followed by Dunn's multiple comparisons test to ascertain the significance of all paired combinations. Differences between groups were considered significant for $p < 0.05$.

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