DL-methionine and _{DL}-methionyl-_{DL}-methionine increase intestinal development and activate Wnt/β-catenin signaling activity in domestic pigeons (*Columba livia*)

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ABSTRACT This experiment was undertaken to investigate the effects of parental dietary _{DL}-methionine (**DL-Met**) and **DL-methionyl-DL-methionine** (**DL-Met-**Met) supplementation on the intestinal development of young squabs. A total of 108 pairs of breeding pigeons and 432 one-day-old squabs were randomly divided into 3 groups: the control group (CON) was fed a basal diet (CP = 15%) and the experimental groups were fed a basal diet supplemented with 0.3% _{DL}-Met or _{DL}-Met-Met. Each pair of breeding pigeons nourished 4 young squabs, and 8 squabs from each treatment were randomly sampled at the end of the experiment. The results indicated that $_{DL}$ -Met and $_{DL}$ -Met-Met supplementation improved the intestinal morphology and structure in the squabs, as reflected by the increased relative intestinal weight of each small intestinal segment, villus height, and villus to crypt ratio. In addition, _{DL}-Met and _{DL}-Met-Met supplementation significantly increased the protein expression of cell proliferation markers (Ki67 and PCNA) and tight junction proteins (ZO-1 and

Claudin-1) in the jejunum and strengthened the fluorescence signal intensity of Ki67, PCNA and Villin. Moreover, the expression of Wnt/β -catenin signaling pathway-related proteins (Frizzled 7 [FZD7], p-GSK- 3β , Active β -catenin, β -catenin, TCF4, c-Myc, and Cyclin D1), and intestinal peptide transporter 1 (**PepT1**) in the jejunum was considerably higher in the treatment group than in the CON group (P < 0.05), with the _{DL}-Met-Met group having the highest expression. Consistently, the molecular docking results predicted the possibility that _{DL}-Met or _{DL}-Met-Met binds to the membrane receptor FZD7, which mediates Wnt/β -catenin signaling. Collectively, the improvement of the intestinal development in squabs after parental dietary 0.3%_{DL}-Met and _{DL}-Met-Met supplementation could be through activation of Wnt/ β -catenin signaling pathway, and _{DL}-Met-Met is superior to _{DL}-Met. Our findings may provide basic data for further optimizing the feeding formula of breeding pigeons and improving the growth and development of squabs.

Key words: pigeon, _{DL}-Met-Met, intestinal development, Wnt/β -catenin signaling, PepT1

INTRODUCTION

The mortality of young squabs is as high as 15%, which is critically affected by intestinal development retardation, low utilization of nutrients and a high prevalence of intestinal diseases. Particularly, young pigeons are initially nourished by parents with crop milk containing mainly proteins and lipids (Dong et al., 2013; Chen et al., 2020). As pigeons grow, the crop milk is

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gradually replaced by grains, a transition characterized by abrupt removal from nurturing by parents and instantaneous exposure to solid feed (Sales and Janssens, 2003; Xie et al., 2019).

As a result, the gut architecture and permeability of the epithelial barrier are altered to adapt to these changes, resulting in excessive multiplication of potential pathogens in the intestinal tract and inducing stress (Spreeuwenberg et al., 2001; Dong et al., 2012). Therefore, it is widely believed that the dietary transition is a stressful event associated with intestinal impairment, most often manifested by villus atrophy and barrier dysfunction, especially in the small intestine (Lalles et al., 2004; Pieper et al., 2008; Wijtten et al., 2011). Nevertheless, the small intestine, especially the jejunum, is a primary site for the

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absorption of nutrients and it plays a vital role in determining the developmental potential of hatching birds (Uni et al., 1998; Liu et al., 2014).

Dietary amino acids are crucial for the maintenance of gut integrity and function. (Wu et al., 2014). As the first-limiting amino acid in diets for poultry, substantial studies have demonstrated that increased methionine (Met) provision above basal requirements can enhance growth performance and meat quality (Wen et al., 2014; Li et al., 2017; Park et al., 2018; Jiang et al., 2019). In addition, supplementation with _{DL}-Met-Met is able to mitigate pathogenic infections and consequent intestinal cellular damage in broilers infected with *Eimeria spp*. (Khatlab et al., 2019). However, studies of the effects of _{DL}-methionine (_{DL}-Met) and _{DL}-methionyl-_{DL}-methionine (_{DL}-Met) on intestinal development in pigeons are still scarce.

Extensive evidence indicates the crucial role of Wnt/ β -catenin signaling in maintaining the gut architecture and homeostasis. More importantly, it facilitates intestinal renewal by controlling cell proliferation and differentiation (Ootani et al., 2009; Clevers and Nusse, 2012; Schuijers et al., 2015). Interestingly, Met deprivation suppressed intestinal stem cell (ISC) proliferation and marker expression in the Drosophila midgut (Saito et al., 2017; Obata et al., 2018). Consistently, Metdepleted medium inhibits canonical Wnt/ β -catenin signaling, while the addition of S-adenosylmethionine (SAM) is sufficient to rescue Wnt signaling (Albrecht et al., 2019). Importantly, Wnt signaling is mediated by the membrane receptor Frizzled7 (FZD7). Nevertheless, further in-depth research is required to reveal the exact mechanism by which _{DL}-Met or _{DL}-Met-Met promotes the intestinal development of pigeons and how Wnt/β -catenin signaling exerts its function in response to _{DL}-Met or _{DL}-Met-Met.

Given the peculiar feeding pattern of breeding pigeons, precise nutrition provision to the parents seems to be an effective method to improve the health status of squabs. Therefore, we explored whether appropriate $_{\rm DL}$ -Met and $_{\rm DL}$ -Met-Met supplementation could improve the intestinal structural morphology and barrier function of squabs and evaluated the effects of $_{\rm DL}$ -Met and $_{\rm DL}$ -Met on the intestinal development of squabs.

MATERIALS AND METHODS

Ethical Statement

All methods and management procedures used in this study were complied with the guidelines established by South China Agricultural University (Guangzhou, China), and the protocol for this experiment was approved by the Animal Ethics Committee of South China Agricultural University (Guangzhou, China).

Animals and Experimental Treatments

 $_{\rm DL}$ -Met (purity >99%) and $_{\rm DL}$ -Met-Met (purity >95%) purchased from Evonik Degussa (Frankfurt,

Germany) were added to the basal diet at a dose of 0.3% (CP = 15%). The methionine level is the optimal doses based on our previous study (Chen et al., 2020).

White king pigeons and experimental sites were provided by WENS Foodstuff Group Co., Ltd. (Yingde, China). A total of 108 pairs of white king breeding pigeons with similar production performance, reproduction performance, and laying eggs on the same day were selected and randomly allocated to 3 treatments, which consisted of 6 replications of 6 pairs of pigeons each. The three experimental diets were formulated as follows: control group (basal diet), 0.3% _{DL}-Met supplementation group and 0.3% _{DL}-Met-Met supplementation group (Table 1). Each pair of breeding pigeons was tending 4 young squabs. The parent pigeons were fed the experimental diets continually for the entire experimental period (46 d), including during the egg incubation period and the feeding period. The squabs were fed crop milk secreted by the parent pigeons. On the 46th d, eight squabs per treatment were euthanized to collect intestine samples and to measure the unit intestinal weight of each section of the small intestine.

Hematoxylin and Eosin Staining

After fixing with 4% paraformaldehyde, the jejunum samples were washed with water, dehydrated using

Table 1. Compositions and nutrient levels of the experimental diets (air-dried basis, %).

Ingredient $(\%)^1$	Basal diet	0.30% DL-Met	0.30% DL- Met-Met
Corn	51.00	51.00	51.00
Soybean meal	17.50	17.50	17.50
Wheat	24.00	24.00	24.00
Sorghum	3.00	3.00	3.00
Dicalcium phosphate	1.20	1.20	1.20
Shell powder	1.00	1.00	1.00
Salt	0.30	0.30	0.30
Vitamin and mineral premix ²	1.00	1.00	1.00
Lysine (98.5%)	0.40	0.40	0.40
DL-Methionine (DL-Met)	0.00	0.30	0.00
DI-Methionyl-DI-	0.00	0.00	0.30
Methionine			
$(_{DL}-Met-Met)$			
Zeolite powder	0.60	0.30	0.30
Total	100.00	100.00	100.00
Formulated nutrient con	nposition $(\%)$		
$\begin{array}{c} {\rm Metabolic\ energy}\\ {\rm (MJ/kg)} \end{array}$	12.13	12.13	12.13
Crude protein	15.00	15.00	15.00
Calcium	0.76	0.76	0.76
Total phosphorus	0.56	0.56	0.56
Nonphytate phosphorus	0.32	0.32	0.32
DL-Met	0.25	0.55	0.25
_{DL} -Met-Met	-	-	0.30
Lysine	0.96	0.96	0.96

¹Provided per kilogram of diet: vitamin A, 4,000.00 IU; vitamin D₃, 1,725.00 IU; vitamin E, 24.00 mg; vitamin K₃, 1.00 mg; vitamin B₁, 3.00 mg; vitamin B₂, 13.00 mg; vitamin B₆, 2.00 mg; vitamin B₁₂, 25.00 μ g; niacin, 15.00 mg; folic acid, 0.55 mg; pantothenic acid, 7.50 mg; biotin, 0.12 mg; choline chloride, 200.00 mg; copper, 10.00 mg; iron, 35.00 mg; manganese, 55.00 mg; zinc, 35.00 mg; iodine, 0.20 mg; selenium, 0.25 mg.

graded ethanol, immersed in wax and made into paraffin-embedded tissues for subsequent histological analysis. Then, 5- μ m thick serial sections of the paraffinembedded tissues were cut by a microtome (Microm-HM340E, Thermo Fisher Scientific), attached to glass slides and stained with hematoxylin and eosin (**H & E**). Finally, the jejunum tissues were observed and photographed under a microscope (Eclipse Ti-S, Nikon, Melville, NY), and the villus height and crypt depth were measured with Image-Pro Plus software.

Western Blotting

Proteins were isolated from jejunum samples and analyzed as described previously (Xie et al., 2020). The expression levels were quantified using ImageJ software (version 1.8.0 112, National Institute of Health, Bethesda, MD). The antibodies used were: anti-ZO-1 (#339100, Thermo Fisher, Waltham, MA), anti-Claudin-1 (#374900, Thermo Fisher), anti-Frizzled7 (FZD7, #bs-5125R, Bioss ANTIBODIES, Beijing, China), antip-GSK-3 β (#5558, Cell Signaling Technology, Danvers, MA), anti-GSK-3 β (#340449, Zen BioScience, Chengdu, Sichuan, China), anti-Active β -Catenin (#19807, CST), anti- β -catenin (#201328, Zen BioScience), anti-T cell factor (TCF4, ab130014, Abcam, Cambridge, MA), anti-CyclinD1 (#SC-450, Santa Cruz Biotechnology, Dallas, TX), anti-c-Mvc (#SC-41, Santa Cruz), antiproliferating cell nuclear antigen (PCNA, #200947, Zen BioScience), anti-PepT1(#SC-373742, Santa Cruz), anti- β -actin (#250132, Zen BioScience), and anti-rabbit IgG (#511203, Zen BioScience) and anti-mouse IgG (#511103, Zen BioScience) secondary antibodies.

Immunofluorescence

The jejunum sections were placed on glass slices as described previously (Chen et al., 2020). Briefly, the samples were blocked in 5% BSA for 2 h and incubated with primary antibodies against ZO-1 (#bs-1329R, Bioss), Ki67 (#NB500-170, Novus Biologicals, Littleton, CO), PCNA (#2586, CST, Danvers, MA), Villin (#SC-58897, Santa Cruz), FZD7 (#bs-5125R, Bioss) and PepT1 (#SC-373742, Santa Cruz) overnight at 4° C and with FITC-(#115-545-003, Jackson Laboratory, Jackson, MS) or Cy3-conjugated secondary antibodies (#111-165-045, Jackson Laboratory) for 2 h at room temperature. The nuclei were stained with DAPI (Sigma-Aldrich, Burlington, MA) for 20 min at room temperature. Fluorescence images were acquired with a microscope (NIS-Elements, Nikon, Tokyo, Japan).

Molecular Docking

The three-position format of $_{DL}$ -Met (Zinc1532529) or $_{DL}$ -Met-Met (ZINC1605259) and the crystal structure of the FZD7 cystine-rich region (10.2210/pdb5T44/pdb) were obtained from ZINC (http://zinc.docking.org/)

and Protein Data Bank (https://www.rcsb.org/). Autodock serials software was used for molecular structure optimization, conformation search, and the molecular docking process. Meanwhile, PyMOL was used for the molecular graphic display to screen the binding sites of $_{\rm DL}$ -Met or $_{\rm DL}$ -Met-Met with FZD7 that could be involved in flexible docking.

Statistics

The data were analyzed using one-way ANOVA with SPSS software version 20 (SPSS Inc., Chicago, IL). The t test and LSD multiple range test were used to evaluate the differences between 2 or multiple groups. The results are expressed as the mean \pm standard error, and *P*-values < 0.05 were considered significant.

RESULTS

DL-Met and *DL*-Met-Met Improve Growth Performance and Epithelial Structural Integrity

To elucidate the effect of _{DL}-Met and _{DL}-Met-Met on the growth performance and intestinal development of squabs, we recorded related indicators of squab growth performance, measured the length and weight of the duodenum, jejunum, and ileum, and examined the morphology of the villi by H&E staining. The results showed a significant increase in average daily gain and a decrease in the mortality of the squabs after $_{\rm DL}$ -Met or $_{\rm DL}$ -Met-Met supplementation (P < 0.05, Figures 1A and 1B). Moreover, compared with the CON group, _{DL}-Met-Met supplementation significantly increased the small intestinal weight of each section (P < 0.05, Figure 1C), while the _{DL}-Met group only had a significant increase in the duodenum. H&E staining indicated that compared with the CON group, parental supplementation with $_{DL}$ -Met-Met significantly increased the height of the villi and the villous crypt ratio (VCR) (P < 0.05, Figures 1D -1G), although the crypt depth in the jejunum was not significantly different among all groups.

_{DL}-Met and _{DL}-Met-Met Increase Abundance of Intestinal Tight Junction Proteins

To clarify whether _{DL}-Met and _{DL}-Met-Met supplementation enhances the intestinal development in squabs, we detected the expression of the tight junction proteins ZO-1 and Claudin1. As shown in Figure 2, compared with the CON group, the expression of ZO-1 and Claudin-1 in the jejunum were significantly increased after _{DL}-Met-Met supplementation (P < 0.05, Figures 2A and 2B), as was the fluorescence signal intensity of ZO-1 (Figures 2C and 2D).



Figure 1. _{DL}-methionine and _{DL}-methionyl-_{DL}-methionine improve growth performance and epithelial structural integrity. (A, B) Growth performance of squabs. (A) Average daily gain; (B) mortality; (C) unit intestinal weight; (D) representative images of H&E staining of the jejunums are shown (40 ×). Scale bar = 500 μ m. (E–G) Statistical analysis of villus height (E), crypt depth (F), and villous crypt ratio (G). "a-c" indicates significant differences (P < 0.05). The values are mean ±S.E.M. with n = 8.



Figure 2. _{DL}-methionine and _{DL}-methionyl-_{DL}-methionine increase abundance of intestinal tight junction proteins. (A) The expression of ZO-1 and Claudin-1 in the jejunum of squabs raised by pigeons supplemented _{DL}-Met or _{DL}-Met-Met. (B) Statistical analysis of the expression of ZO-1 and Claudin-1 based on the images shown in (A). (C) Representative images of IF staining with ZO-1 antibodies of the pigeon jejunum (100 ×). Scale bar = 100 μ m. (D) Statistical analysis of the expression of ZO-1 based on the images shown in (C). "a-b" indicates significant differences (P < 0.05). The values are mean ±S.E.M. with n = 3.

DL-Met and *DL*-Met-Met Promote the Proliferation and Differentiation of Intestinal Cells

Subsequently, we tested the effects of _{DL}-Met and _{DL}-Met-Met on the proliferation and differentiation of intestinal epithelial cells. A stronger fluorescence signal intensity of Ki67 (a marker of proliferating cells) (P < 0.05, Figures 3A and 3D), PCNA (P < 0.05, Figures 3B and 3D) and Villin (marker of absorptive cells) (P < 0.05, Figures 3C-3D), coupled with increased expression of PCNA (a marker of proliferating cells) (P < 0.05, Figures 3E and 3F), was observed in squab jejunum after parental dietary _{DL}-Met and _{DL}-Met-Met supplementation, suggesting that _{DL}-Met promotes the differentiation of ISCs into epithelial absorptive cells.

DL-Met and *DL*-Met-Met Potentiate PepT1 Expression

In addition, immunofluorescence (Figures 4A and 4B), together with western blotting (Figures 4C and 4D), showed that the protein expression of intestinal

peptide transporter 1 (**PepT1**) in the treatment group was significantly increased (P < 0.05) in comparison with that in the CON group, and the _{DL}-Met-Met group exhibited the highest expression of PepT1.

_{DL}-Met and _{DL}-Met-Met Intensify Wnt/ β-Catenin Activity by Upregulating the Expression of Membrane Receptor FZD7

Next, we explored the probable mechanism of $_{\rm DL}$ -Met or $_{\rm DL}$ -Met-Met in promoting intestinal development and barrier function. We found that the Wnt/ β -catenin pathway was significantly upregulated after $_{\rm DL}$ -Met or $_{\rm DL}$ -Met-Met supplementation. Specifically, the GSK-3 β level was decreased, and the levels of FZD7, p-GSK-3 β , Active β -catenin, β -catenin, TCF4, c-Myc, and Cyclin D1 were increased (Figures 5A and 5B). Consistently, the fluorescence signal intensity of FZD7 was significantly increased (P < 0.05, Figures 3B and 3C). The docking results predicted that there were 5 or 7 hydrogen bonds between $_{\rm DL}$ -Met or $_{\rm DL}$ -Met and the cysteine-rich region of FZD7, suggesting that $_{\rm DL}$ -Met or $_{\rm DL}$ -Met-Met might bind to the subdomain of FZD7. The ZHONG ET AL.



Figure 3. _{DL}-methionine and _{DL}-methionyl-_{DL}-methionine promote the proliferation and differentiation of intestinal cells. (A–C) Representative images of IF staining with Ki67 (A), PCNA (B), and Villin (C) antibodies of the pigeon jejunum (100 ×). Scale bar = 100 μ m. (D) Statistical analysis of the expression of Ki67, PCNA and Villin based on the images shown in (A–C). (E) The expression of PCNA in the jejunum of squabs raised by pigeons supplemented with _{DL}-Met or _{DL}-Met-Met. (F) Statistical analysis of the expression of PCNA based on the images shown in (E) "a-c" indicates significant differences (P < 0.05). The values are mean ±S.E.M. with n = 3.

docking calculations showed the lowest binding energy conformer, with binding energies of -2.86 and -1.72 kcal/mol, respectively (Figures 5E and 5F).

DISCUSSION

The intestines are the most important organs wherein most nutrients are absorbed (Feng et al., 2019). Desirable dietary interventions targeting parental pigeons are proposed to be a good strategy to facilitate the maintenance of intestinal health status and homeostasis in squabs (Xu et al., 2020a,b). It is now generally accepted that intestinal functions are reflected by the intestinal size and structure, especially the villus height and crypt depth. Our present study demonstrates that parental supplementation of _{DL}-Met and _{DL}-Met resulted in a heavier intestinal weight, especially in the duodenum, compared with that of the CON group. In addition, a significant increase in villus height was observed in the DL-Met and DL-Met-Met groups. Increased intestinal weight and villus height are usually associated with an enhancement of digestive and absorptive capacity by the small intestine, which in turn improves the health status of the animals (Upadhaya et al., 2020). Furthermore, the VCR in the jejunum was significantly increased in squabs by parental dietary _{DL}-Met and _{DL}-Met-Met supplementation. A higher VCR generally equates to a more absorptive area, and fewer intestinal epithelial cells (IECs) are recruited for cell renewal; therefore, more abundant enterocytes are available for the digestion and absorption of nutrients (Qin et al., 2019). Collectively, _{DL}-Met and _{DL}-Met-Met enhanced intestinal development by improving the small intestinal



Figure 4. _{DL}-methionine and _{DL}-methionyl-_{DL}-methionine potentiate PepT1 expression. (A) Representative images of IF staining with PepT1 antibody in the pigeon jejunum (100 ×). Scale bar = 100 μ m. (B) Statistical analysis of the fluorescence signal intensity based on the images shown in (A). (C) The expression of PepT1 in the jejunum of squabs raised by pigeons supplemented with _{DL}-Met or _{DL}-Met-Met. (D) Statistical analysis of the expression of PepT1 based on the images shown in (C). "a-c" indicates significant differences (P < 0.05). The values are mean ±S.E.M. with n =3.

morphology, and significant improvements were observed in the $_{\rm DL}$ -Met-Met group compared with the $_{\rm DL}$ -Met group.

The intestinal barrier is a semipermeable structure that is comprised of tightly attached continuous monolayer IECs and tight junctions (**TJs**) between enterocytes. The TJs serve as a protective barrier that allows for the uptake of necessary nutrients while restricting pathogenic molecules and bacteria from the external environment from escaping from the intestinal lumen (Chelakkot et al., 2018). It has been confirmed that compromised intestinal barrier function contributes to the risk of a variety of intestinal and systemic diseases (Odenwald and Turner, 2017). Importantly, transmembrane protein complexes (e.g., claudins and occludin) and peripheral membrane proteins (e.g., ZO-1 and ZO-2) together constitute TJs. The assembly and combination of diverse TJs are of premier importance to seal the intercellular space and further regulate the paracellular and transcellular pathways. Interestingly, Claudin-ZO protein interactions, which link claudins to the actin cytoskeleton through scaffold proteins such as ZO-1, are essential for tight junction assembly (Findley and Koval, 2009). Our study demonstrated that parental dietary _{DL}-Met-Met improved the intestinal development of squabs, as shown by significantly increased expression of tight junction proteins in the jejunum (ZO-1 and Claudin-1). Based on the size-selective sieving of molecules by TJs, different claudin isoforms may be differentially expressed to perform their corresponding functions (Peltonen et al., 2007). This might explain the inconsistent expression of claudin-1 and ZO-1 in the _{DL}-Met group.

Enterocytes express tight junction proteins that modulate the permeability of the epithelium and thereby maintain the intestinal integrity (Peterson and Artis, 2014; Ahmad et al., 2017). Moreover, constant self-renewal of IECs depends on vigorous proliferation of ISCs in crypts. Consequently, the proliferation of ISCs is indispensable for sustaining the integrity of the intestinal structure and barrier function. Both Ki67 and PCNA are markers of cell proliferation. In the present study, _{DL}-Met and _{DL}-Met-Met supplementation significantly increased the expression of Ki67 and PCNA in the jejunum, indicating that _{DL}-Met and _{DL}-Met-Met stimulate the prompt renewal of the intestinal epithelium by accelerating cell proliferation. During development, IECs differentiate into firmly cohesive polarized cells, wherein many microvilli could increase the cell surface contact area and favor absorptive and secretory functions (Lhocine et al., 2015). In particular, as a plentiful actin-modifying protein in the epithelial brush border, Villin participates in the formation of the epithelial structure and the function of the microvilli (Delacour et al., 2016). Our study showed that Villin was highly expressed after _{DL}-Met or _{DL}-Met-Met supplementation (especially _{DL}-Met-Met), suggesting that _{DL}-Met and _{DL}-Met-Met could maintain the architecture of the IECs and enhance their absorptive capacity.

The maintenance of stem cells in the intestine involves complex interactions of multiple signal transduction pathways, especially Wnt/ β -catenin (van der Flier and Clevers, 2009; Merenda et al., 2020). More importantly, attenuating Wnt/ β -catenin leads to decreased cell proliferation capacity and intestinal dysfunction (Hu et al., 2018; Li et al., 2019). Similarly, our previous study demonstrated that Met or HMB ameliorates deoxynivalenolinduced ISC injury through Wnt/ β -catenin signaling reactivation, which promotes ISC expansion and further drives intestinal epithelial regeneration (Zhou et al.,



Figure 5. _{DL}-methionine and _{DL}-methionyl-_{DL}-methionine increase abundance of FZD7 and activate Wnt/ β -catenin signaling pathway in pigeon jejunal tissue. (A) The expression of Wnt/ β -catenin signaling pathway-associated proteins in the jejunum of pigeons with _{DL}-Met or _{DL}-Metore. (B) Statistical analysis of the expression of target proteins based on the images shown in (A). (C) Representative images of IF staining with FZD7 antibodies of the pigeon jejunum (100 ×). Scale bar = 100 μ m. (D) Statistical analysis of the expression of FZD7 based on the images shown in (C). (E, F) Binding mode of _{DL}-Met to FZD7 (E) and _{DL}-Met-Met to FZD7 (F). The protein is shown in cartoon form. Surrounding residues are represented in stick form. The dashed lines indicate the formation of hydrogen bonds. The numbers beside the dash show the distance (Å) of hydrogen bonds. "a-c" indicates significant differences (P < 0.05). The values are mean ±S.E.M. with n = 3.

2019). Previous study have shown that the methionine metabolite SAM might be activating wnt signaling (Albrecht et al., 2019), However, whether DL-Met can act on the Wnt/ β -catenin pathway in other fashion is poorly understood. In the current study, we confirmed that parental dietary _{DL}-Met or _{DL}-Met-Met resulted in a higher level of FZD7 in the squab jejunum. Similarly, molecular docking analysis predicted the possible sites

where $_{DL}$ -Met or $_{DL}$ -Met-Met might bind to the membrane receptor FZD7.

Based on our results, we speculate that the direct binding between _{DL}-Met or _{DL}-Met-Met and FZD7 leads to enhanced activity of the canonical Wnt pathway, which promotes the phosphorylation of GSK-3 β and inhibits GSK-3 β (Figure 6). As a consequence, the protein complex (including CK1 α , GSK3, Axin1, and APC)



Figure 6. Integrated model depicts the mechanism by which $_{DL}$ -methionine and $_{DL}$ -methionyl- $_{DL}$ -methionine intensify FZD7-mediated Wnt/ β -catenin signaling activity and the pathway involved in the cellular uptake of $_{DL}$ -methionyl- $_{DL}$ -methionine mediated by PepT1 in the brush border membrane of enterocytes.

falls apart, and unphosphorylated β -catenin is stabilized and accumulates, subsequently translocating into the nucleus and binding to TCF to induce the expression of its target genes (Nusse and Clevers, 2017). Interestingly, _{DL}-Met-Met seems to be more efficient than _{DL}-Met, which may be explained by the fact that the high transport capacity of PepT1 activated by _{DL}-Met-Met allows intestinal $_{\rm DL}$ -Met (Daniel, uptake of 2004:Flanagan et al., 2019). Furthermore, _{DL}-Met-Met might bind to FZD7 directly or be transported into the cytoplasm by PepT1 to be hydrolyzed to the corresponding _{DL}-Met, which engages with FZD7, which in turn enhances Wnt/ β -catenin signaling activity. Nonetheless, further research should be performed to reveal the exact mechanism by which _{DL}-Met or _{DL}-Met-Met regulates Wnt/β -catenin during intestinal development.

In the jejunum, PepT1 is abundantly expressed specifically in well-differentiated absorptive epithelial cells in the villi, shedding light on the pivotal role of PepT1 in the absorption process (Nguyen, et al., 2007; Dranse et al., 2018). Through the PepT1 KO mouse

model, Zhang et al. (2016) demonstrated that PepT1 critically modulates miRNA and protein expression directly or indirectly along the crypt/villus axis in the jejunum and thus facilitates the maintenance of intestinal homeostasis and proper functions. Consistent with previous results, the immunofluorescence of PepT1 showed that the highest expression was detectable at the villus tip, while little expression was detectable at lower crypt cells (Spanier and Rohm, 2018). It is plausible that the expression gradient of PepT1 along the crypt-villus axis is quite important in intestinal function and the maintenance of normal intestinal epithelial integrity. Notably, the _{DL}-Met-Met group exhibited the highest expression of PepT1, which may be attributed to the fact that Met-Met serves as a PepT1 substrate to activate the PepT1 promoter and enhance the transcriptional activity of PepT1 (Wang et al., 2017). Furthermore, PepT1 activity may be directly regulated by amino acids (Shiraga et al., 1999). However, the regulatory mechanisms by which _{DL}-Met acts on PepT1 still require further investigation.

CONCLUSIONS

Our study found that parental dietary supplementation with _{DL}-Met and _{DL}-Met-Met effectively increases intestinal development in squabs, we speculate that the reason for this might be the reinforcement of PepT1 expression and Wnt/ β -catenin signaling activity (Figure 6). Most importantly, _{DL}-Met-Met appears to be more efficient than _{DL}-Met. Our findings support a potential nutritional manipulation strategy to improve the growth and intestinal development of squabs.

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DISCLOSURES

We declare that we have no financial and personal relationships with other people or organizations that can inappropriately influence our work, there is no professional or other personal interest of any nature or kind in any product, service and/or company that could be construed as influencing the content of this paper.

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