

Cite this: *RSC Adv.*, 2017, 7, 43345

Effects of wheat peptide supplementation on anti-fatigue and immunoregulation during incremental swimming exercise in rats

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This study elucidated the effects of wheat peptide administration on anti-fatigue and immunoregulation functions in rats. Wheat peptides were separated and the fraction with the highest radical scavenging activity *in vitro* was subjected to mass spectrometry to identify the peptide sequences. Sixty rats were randomized into 5 groups: no exercise control group (C), no exercise with low dose [20 mg kg⁻¹ d⁻¹] group (M), exercise control group (E), exercise with low dose group (Z), and exercise with high dose [100 mg kg⁻¹ d⁻¹] group (D). After training for 4 weeks with incremental swimming exercise, bodyweight and exhaustive time were tested and serum, small intestine, skeletal muscle and brain tissues of the rats were collected. A total of four peptide sequences from the highest active fraction were identified. The exhaustive time of group D was significantly longer than groups E and Z. The malondialdehyde content of group M was significantly lower than group C ($p < 0.01$), but secretory immunoglobulin A and 5-hydroxytryptamine (5-HT) were higher (both $p < 0.01$). Compared to group E, the activities of superoxide dismutase in skeletal muscle and acetylcholinesterase were significantly higher in groups Z and D ($p < 0.01$ or 0.05), but caspase-3 was lower ($p < 0.01$). Glutathione peroxidase and 5-HT in group D were both significantly higher than in group E ($p < 0.01$ or 0.05), but interleukin-6 and interleukin-8 were lower ($P < 0.05$). Supplementation of wheat peptide could effectively improve the ability of one-time exhaustive exercise of rats, remove free radicals from skeletal muscle in time, and alleviate the intestinal and blood inflammatory responses.

Received 18th July 2017
Accepted 1st September 2017

DOI: 10.1039/c7ra07860a

rsc.li/rsc-advances

1. Introduction

Wheat protein is a byproduct of wheat starch processing, and is mainly composed of gliadin and glutenin.¹ Wheat protein contains abundant hydrophobic amino acids and uncharged amino acids, and leads to a large area of intramolecular hydrophobicity and poor solubility in water,² which consequently hinders the utilization of wheat protein. Enzymatic modification increases the solubility of wheat protein, which in turn improves the processing performance and functional properties of wheat protein. Enzymatic hydrolysis of wheat produces a large amount of peptide material with different molecular weights. In recent years, plant-derived peptides, including soy peptides, corn peptides, *etc.*, have been widely used as raw food materials in the food industry. But wheat peptide has not been widely used as a new source of raw food

material. The development of deep processing of wheat protein showed good application prospects for wheat peptide in food industry.

Current research is focusing on the functional properties of wheat peptides. Wheat peptides have a variety of biological functions, such as antioxidant activities. Žilić *et al.*³ have reported that the free radical scavenging potential of wheat peptide was stronger than that of legume protein hydrolysate. Similarly, few other studies^{4,5} have also indicated that wheat peptides have stronger antioxidant activity. On the other hand, a large number of studies^{6–9} indicated that wheat peptides can significantly inhibit ACE activity and have anti-hypotensive effect. As early as 1979, Ziadrou *et al.*¹⁰ have obtained opioid-like active peptides from wheat protein hydrolysate. Fukudome *et al.*¹¹ have also shown that high concentrations of wheat enkephalin could be obtained in the wheat protein hydrolysates. In addition, few other studies have shown the tumor suppressor activity of wheat protein. Calzuola *et al.*¹² isolated acidic peptides from the chromatin of wheat germ by using infrared spectroscopy and mass spectrometer, and demonstrated its role in controlling cell proliferation. Jeong *et al.*¹³ also reported that the tumor suppressor peptide extracted from the wheat protein inhibited the acetylation of nucleoprotein.

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Wheat peptide has a variety of functional properties, but few studies have demonstrated its anti-fatigue and regulation of immune functions. In the studies that reported anti-fatigue function of plant-derived peptides, soybean and corn peptides tend to have a certain anti-fatigue effects.^{14,15} Yimit *et al.*,¹⁶ demonstrated that soybean peptide regulates immune function well. Wheat peptides, soybean peptides, as well as corn peptides were obtained in plant protein hydrolysates, and therefore, wheat peptide might be associated with the anti-fatigue and immune regulation functions. High intensity exercise causes a series of physiological and biochemical changes in muscle, intestine, blood, and nervous system, leading to the occurrence of fatigue and decreased immunity. These mainly reflect in the declination of glycogen amount, increase oxidative stress, descent intestinal mucosal permeability, muscle damage, inflammatory responses, *etc.*¹⁷ Animal testing is a good way to validate both anti-fatigue and immune regulation functions of wheat peptide, and is the essential part before human clinical trials. Therefore, animal experiments were conducted in this study to elucidate the effects of wheat peptide on the growth of rats, as well as the related indexes of serum, small intestine, skeletal muscle, and brain tissue in rats. Exhaustive time that reflects the direct impact of rat exercise capacity was tested, and the mechanism of anti-fatigue and immune regulation of wheat peptide was also discussed. Hence, the aim of this study was to provide a theoretical basis for the functional diversity of wheat peptides, and further develop a kind of functional food ingredient with anti-fatigue and immunoregulation functions for processing military food. This can alleviate the fatigue and disturbances of immune regulation in a large number of physical training of soldiers.

2. Materials and methods

2.1 Materials

2.1.1 Samples. Wheat protein was provided by Shangqiu Huayang Ecological Agriculture Development Co., Ltd., and the protein content was 81.91%.

Alkaline protease (2.2×10^5 U mL⁻¹) and flavor protease (3.1×10^4 U mL⁻¹) were provided by Novozymes Biotechnology Co., Ltd.

2.1.2 Experimental animals. Sixty specific pathogen free (SPF) Sprague-Dawley (SD) rats weighing 282.19 ± 7.95 g were provided by Beijing Veitong Lihua Experimental Animal Center. Animal license number: SCXK (Beijing) 2006-0009.

2.1.3 Animal feed. Conventional feeds consisting of 58.9% corn, 25% soybean meal, 10% sucrose, 1% lard, 0.4% salt, 3.5% mineral premix, 1% vitamin premix, 0.2% choline tartrate were provided by the PLA Academy of Military Medical Sciences. Preparation of mineral and vitamin premixes were conformed according to the AIN-93G feed standard.

2.1.4 Reagents. MDA kit, SOD kit, GSH-Px kit, TTE radioimmunoassay kit, COR radioimmunoassay kit, IL-6 kit, IL-8 kit, Caspase-3 kit, sIgA radioimmunoassay kit, ET kit, AchE kit were provided by the Beijing Huayang Biotechnology Research Institute. CK kit was supplied by Zhongsheng Beikong Biotechnology Co., Ltd. Glycogen kit was provided by Nanjing Institute

of Jiancheng Bioengineering. 5-HT kit was supplied by DIASOURCE.

2.2 Methods

2.2.1 Preparation of wheat peptide. Wheat protein was dispersed in deionized water at a concentration of 11.2% (w/v), incubated in a water bath at 55 °C, the pH of the solution was adjusted to 8.5, the alkaline protease of 2200 U g⁻¹ was added, and then treated by enzymatic hydrolysis for 4.3 h. After enzymolysis, the enzyme was inactivated by boiling water bath for 10 min. After adjusting the temperature to 50 °C and pH value to 6.5, the flavor protease of 1070 U g⁻¹ was added and the enzymatic hydrolysis lasted for 2.2 h. pH of the solution was kept constant with 1.0 mol L⁻¹ NaOH or HCl for every 20 min during the enzymatic hydrolysis. The enzyme was inactivated by boiling water bath for 10 min at the end of enzymatic reaction. Centrifugation was performed at 3000 rpm for 20 min after cooling down, and the supernatant was freeze-dried at -20 °C and allowed to stand-by.

2.2.2 Determination of molecular weight distribution of wheat peptide. Molecular weight distribution of wheat peptide was analyzed by high performance gel filtration chromatography (Shimadzu, Kyoto, Japan). The chromatographic conditions were as follows. Column: TSK gel G2000 SWXL (300 mm × 7.8 mm). Mobile phase: volume ratio of acetonitrile–water–TFA was 45 : 55 : 0.1. Detection wavelength: 220 nm. Flow rate: 0.5 mL min⁻¹. Column temperature: 30 °C. Injection volume: 10 μL. The standard calibration curve of relative molecular mass was prepared with bacitracin (M_w 1500 Da), cytochrome C (M_w 12 500 Da), aprotinin (M_w 6500 Da), glycine-glycine-tyrosine-arginine (M_w 451 Da) and glycine-glycine-glycine (M_w 189 Da). The regression equation between the logarithm of relative molecular mass and retention time is $y = -0.2095x + 6.6081$ ($R^2 = 0.9975$). The relative molecular mass and distribution range of the sample were obtained by substituting the chromatographic data of the sample into the calibration curve equation.

2.2.3 Amino acid composition analysis. The total and free amino acid composition of wheat peptide sample was determined by an automatic amino acid analyzer (Hitachi, Tokyo, Japan), according to the method of Gu *et al.*,¹⁸ with some modifications. In total amino acid composition analysis, all amino acids except tryptophan were analyzed by hydrolyzing samples with 6 mol L⁻¹ HCl in tubes sealed under nitrogen at 110 °C for 24 h, and the samples were treated with performic acid in cysteine and methionine analysis. Tryptophan was analyzed by dissolving the samples at 110 °C for 22 h in 5 mol L⁻¹ NaOH. After removal of impurities, the free amino acid composition test was directly performed. The bound amino acid composition was obtained by subtracting the content of free amino acid from total amino acid.

2.2.4 Separation of wheat peptide. The lyophilized wheat protein hydrolysate was dissolved in deionized water and passed through an ultrafiltration membrane with a molecular weight cutoff value of 1000 Da. The fraction with molecular weight less than 1000 Da was collected and lyophilized. Wheat



Table 1 Details of rats swimming training program^a

Weeks	Monday (min)	Tuesday (min)	Wednesday (min)	Thursday (min)	Friday (min)	Saturday (min)
1	20	30	40	50	60	60
2	70	75	80	85	90	90
3	95	100	105	110	115	120
4	120*1	120*1	120*1	120*2	120*2	120*2

^a Note: the number after each “*” was the ratio of weight-bearing to weight of each rat (%).

peptide with molecular weight less than 1000 Da was continuously separated by reversed-phase high performance liquid chromatography (RP-HPLC) (Waters, USA) using a XBridge BEH130 C18 column (4.6 × 250 mm, 5 μm, Waters, USA) as described by Gu *et al.*,¹⁸ with some modifications. Peptides were eluted with eluent A [deionized water containing 0.1% (v/v) trifluoroacetic acid (TFA)], and eluent B [acetonitrile with 0.1% (v/v) TFA] using the following gradient program: 0–80 min, 0–35% B; 80–100 min, 35–80% B; 100–105 min, 80% B; 105–113 min, 80–5% B; 113–120 min, 5% B. Wheat peptide was isolated at a flow rate of 5 mL min^{−1} and determined at 220 nm. Sample concentration was 50 mg mL^{−1} and the injection volume was 300 μL. The major fractions were collected and lyophilized for radical scavenging activity assay and sequence identification. 1,1-Diphenyl-2-picrylhydrazyl (DPPH) radical scavenging activity assay was performed according to the method described in Li *et al.* study.¹⁹

2.2.5 Identification of peptides. A Waters corporation matrix-assisted laser desorption/ionization time of flight mass spectrometry (MALDI-TOF-MS) was used for the identification of fractions. A binary mobile phase consisted of acetonitrile and deionized water with 0.1% (v/v) formic acid. The solution was injected with a flow rate of 0.3 mL min^{−1}. The injection volume was 5 μL. The capillary voltage was 3.5 kV in the positive ion mode. Instrumental control and data analysis were performed using Waters MassLynx software.

2.2.6 Feeding conditions. The experimental animals were fed in the animal room under SPF barrier environment of Beijing Sport University. The rats included were 4 per cage, maintained a room temperature of 20–24 °C, with a relative humidity of 50–70%, free intake of drinking water, good indoor ventilation conditions, and normal day and night changes. The animals were treated according to the ethical guidelines of Beijing Sport University. All animal test programs were approved by the Institutional Animal Ethics Committee of Beijing Sport University.

2.2.7 Grouping and training programs. Sixty rats were randomly divided into five groups: no exercise control group (C), no exercise with low dose group (M), exercise control group (E), exercise with low dose group (Z), and exercise with high dose group (D), with 12 rats in each group. Groups E, Z and D took incremental load swimming exercise for four weeks. Training program details were shown in Table 1.

2.2.8 Dose of gavage. The gavage administration with wheat peptide solution, prepared with fresh saline, was carried out at 15 : 00 to 16 : 00 every day from Monday to Saturday, and

stopped on Sunday. The mass concentration of wheat peptide solution was 25 mg mL^{−1}. The modest gavage dose of 20 mg kg^{−1} body weight per day was given to the rats of M and Z groups, and the gavage dose of 100 mg kg^{−1} body weight per day was given to the rats of group D.

2.2.9 Exhaustive experiment. An exhaustive experiment after the last training session was conducted in E, Z, and D groups. The exhaustive experiment conditions were as follows: use of tail-weight swimming, weight-bearing of 3% of weight, pool water depth of 50 cm, water temperature of 31 ± 1 °C, and 5 to 6 rats swam at the same time in the water per square meter. Rats swam to the bottom of the water for 3 consecutive times, and each turn lasted for more than 10 s. The rats were then considered as exhausted, and their exhausted time was recorded.

2.2.10 Sampling. After 24 hours of exhaustive experiment, all the rats were administered anesthesia and sacrificed. The abdominal aorta blood was taken and centrifuged to extract the serum. Brain tissue, proximal jejunum of 3 cm, and quadriceps were taken and stored at −80 °C.

2.2.11 Index determination. CK, TTE, COR, ET, IL-6 and IL-8 of serum sample, IL-6, IL-8, sIgA, caspase-3 after intestinal sample homogenate, and IL-6 in skeletal muscle samples were measured by radioimmunoassay. SOD, MDA, GSH-Px, muscle glycogen, and AchE and 5-HT of brain tissue were measured by colorimetry.

2.3 Statistical analysis

Data were processed using SPSS 13.0 statistical software. Results were expressed as mean ± standard deviation (means ± SD). Using single factor analysis of variance, *P* < 0.05 showed significant difference, *P* < 0.01 indicated that the difference was very significant.

3. Results and discussion

3.1 Characterization and identification of wheat peptide

As shown in Table 2, wheat protein was digested by alkaline protease and flavor protease to prepare wheat peptide samples, and components with molecular weight less than 1000 Da accounted for 82.15%. Molecular weight less than 180 Da is usually considered as a free amino acid component. Therefore, the ratio of wheat peptide with molecular weight less than 1000 Da was 71.94% and the major molecular weight was located at 180–500 Da (54.66%) in this experiment. These results indicated that wheat peptide was mainly composed of



Table 2 Molecular weight distribution of wheat peptide^a

Molecular weight range/Da	Content/%
>10 000	1.42 ± 0.24
5000–10 000	1.65 ± 0.36
3000–5000	2.19 ± 0.61
2000–3000	3.05 ± 0.12
1000–2000	9.54 ± 0.44
500–1000	17.28 ± 0.83
180–500	54.66 ± 1.89
<180	10.21 ± 0.58

^a Data are shown as mean ± SD.

small peptides. It was reported that small active peptides were more easily absorbed and demonstrated higher biological activities than longer peptides *in vivo* as they were less susceptible to undergo gastrointestinal hydrolysis.²⁰

The amino acid composition of wheat peptide sample was analyzed and the results were shown in Table 3. Results demonstrated that the most abundant amino acids were glutamic acid and glutamine (37.7 g/100 g in total amino acids). It was reported that glutamine is beneficial for improving the body's immune function,²¹ so wheat peptide may also have its role in immunoregulation. Saito *et al.* have shown that His, Pro and Tyr could contribute to the higher radical scavenging ability.²² In this study, the total content of His, Pro and Tyr was 10.5 g/100 g in total amino acids, which may provide the wheat peptide with strong antioxidant capacity. These results suggest that the biological activities of wheat peptide seemed to be due to these specific amino acids.

Table 3 Amino acid composition of wheat peptide sample (g/100 g peptide)

Amino acid	Total amino acid	Free amino acid	Bound amino acid
Asp ^a	2.3	0.1	2.2
Glu ^b	37.7	0.4	37.3
Ser	3.7	0.2	3.5
His	1.6	0.2	1.4
Gly	3.0	0.2	2.8
Thr	1.7	0.3	1.4
Arg	2.7	0.8	1.9
Ala	1.9	0.5	1.4
Tyr	2.6	0.5	2.1
Cys	0.7	n.d. ^c	0.7
Val	3.1	0.3	2.8
Met	1.5	0.4	1.1
Phe	4.6	0.6	4.0
Ile	2.7	0.2	2.5
Leu	5.0	1.2	3.8
Lys	1.1	0.4	0.7
Pro	6.3	0.1	6.2
Trp	0.6	n.d. ^c	0.6
Total	82.8	6.4	76.4

^a Aspartic acid + asparagine. ^b Glutamic acid + glutamine. ^c Not determined.

Wheat peptide with molecular weight less than 1000 Da was separated by RP-HPLC, and the chromatogram was shown in Fig. 1. Five fractions were selected and designated as F1–F5. Research suggested that fatigue was positively correlated with free radical increase.²³ Thus, DPPH radical scavenging activities of these five fractions were measured *in vitro* at a concentration of 3.0 mg mL^{−1}. As shown in Fig. 2, fraction F4 (86.61 ± 3.22%) showed the highest DPPH radical scavenging activity. Therefore, fraction F4 was selected for further identification of amino acid sequences.

MALDI-TOF-MS for peptide identification was performed using the highest active fraction F4. A total of four peptide sequences were identified (Table 4). The molecular masses of all these identified peptides are less than 500 Da. All of them were peptides with 2–4 amino acid residues. The biological activities of these peptides depend on their amino acid composition and structure. Suetsuna *et al.*²⁴ demonstrated that hydrophobic amino acid residues, such as Ala or Leu, were helpful for radical scavenging. In addition, the active peptides usually have specific amino acid sequences. One study demonstrated strong antioxidant activity of the peptides containing Leu residues at the N-terminus and Arg residues at the C-terminus (Leu-Asp-Arg).²⁵ The study of Saito *et al.* also proved higher antioxidant property of Leu-Trp-Arg.²² In this study, Leu-Ala-Arg was identified from fraction F4, and meant that this was associated with higher radical scavenging activity, which may be due to a tripeptide with specific amino acid sequence.

3.2 Effect of wheat peptides on exercise ability and fatigue resistance

Long-term high-intensity training may lead to endocrinal imbalance, movement system strain, energy metabolism disorder, and further may affect the ability to exercise and recovery ability after fatigue. It has been reported that excessive training would increase blood viscosity,²⁶ and inhibit brain absorption of complex ammonia, leading to emotional depression.²⁷

The indicators for determining the declination of body function includes TTE and COR. TTE is the main male androgen and it increases the protein synthesis metabolism and improves athletic ability. COR is mainly generated in stress conditions, which is related to the body's catabolism, leading to the declination of exercise capacity. Under normal circumstances, synthesis and decomposition of blood TTE and COR maintains balance. However, long-term high-intensity training would lead to hypothalamic dysfunction, and gradual decline of TTE in the blood. The reason could be due to the decrease in the cleavage enzyme activity of cholesterol side chain in the testicular interstitial cells.²⁸ At the same time, the concentration of COR was gradually increased, and the protein catabolism was increased, resulting in the decrease ratio of blood TTE to COR.^{29,30} In this study, the model of increasing load intensity swimming exercise in rats was used. Results showed the weight gain of the rats in the exercise group was slightly slower than that of the control group (Fig. 3), proving that the long-term and high-intensity exercise training hindered the normal growth of



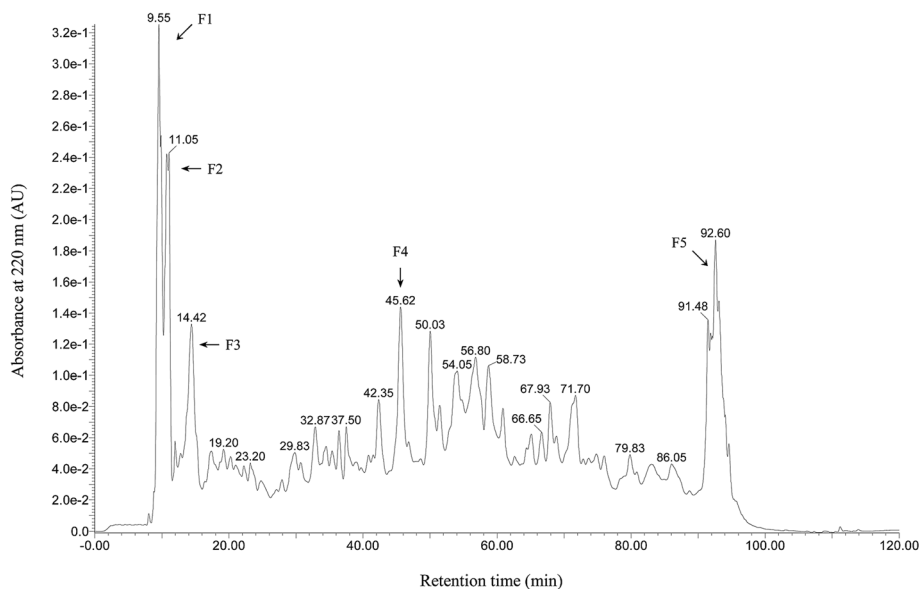


Fig. 1 RP-HPLC chromatogram of wheat peptide. Separation was programmed on a XBridge BEH130 C18 column (4.6 × 250 mm, 5 μm) using gradients: eluent A (0.1% TFA in deionized water, v/v) and eluent B (acetonitrile with 0.1% TFA, v/v) at a flow rate of 5 mL min⁻¹. Collected fractions are numbered from F1 to F5.

the body. However, the level of TTE and COR in this study showed no significant changes between each group (Table 6), which may be due to the loss of endocrine system dysfunction with training. Another reason for this may be that the serum was taken 24 h after the rat's exhaustive test, and the fatigue of the exercise stress in the rats was fully restored, resulting in a stable level of TTE and COR. Rahimi *et al.*³¹ have shown that during high-intensity endurance training, TTE/COR ratio of long-term resting was higher than that of short-term resting, which was helpful in the stable maintenance of TTE and COR levels. In addition, supplementation of wheat peptides in this study showed no significant effect on TTE and COR levels.

Although TTE and COR are associated with the synthesis and decomposition of proteins, additional protein supplementation may not have any effect on both the substances.

In Table 6, there was no significant difference in CK between each group. CK is a non-functional enzyme present in human serum that does not catalyze in the blood and reflects only the destruction of organ cells or cell permeability. CK is mainly present in skeletal muscle, myocardium and brain tissues. Changes in CK levels are often used as an important indicator for the assessment of skeletal muscle load, injury, adaptability and recovery.^{32,33} Results of this study indicated that the high-intensity training did not cause skeletal muscle injury, and that supplemented with wheat peptides had no significant effect on skeletal muscle injury. According to the previous studies, supplementation of protein or protein hydrolysates showed no significant preventive effect on skeletal muscle injury and recovery during exercise,^{34,35} and these results were consistent with our study findings.

Glycogen is mainly stored in the skeletal muscle and liver, and serves as a form of energy storage. Long-term exercise can promote the body's capacity of uptake and utilization of sugar, and increase muscle glycogen and liver glycogen storage.³⁶ In Table 7, muscle glycogen of group M was significantly lower than that in group E ($p < 0.05$), group Z ($p < 0.05$) and group D ($p < 0.01$). Results of our study showed that the muscle glycogen of exercise group was significantly higher than the quiet group, indicating that exercise can improve muscle glycogen storage. In the exercise groups, there was no significant effect with wheat peptide supplementation on muscle glycogen storage, indicating that supplementation of wheat peptide did not have a significant effect on muscle glycogen storage during exercise. Glycogen is mainly composed of excess glucose, and wheat peptide is the hydrolysate derived during the wheat protein

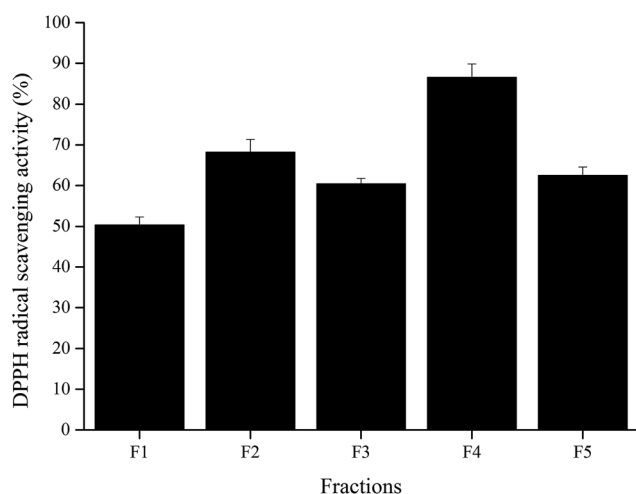
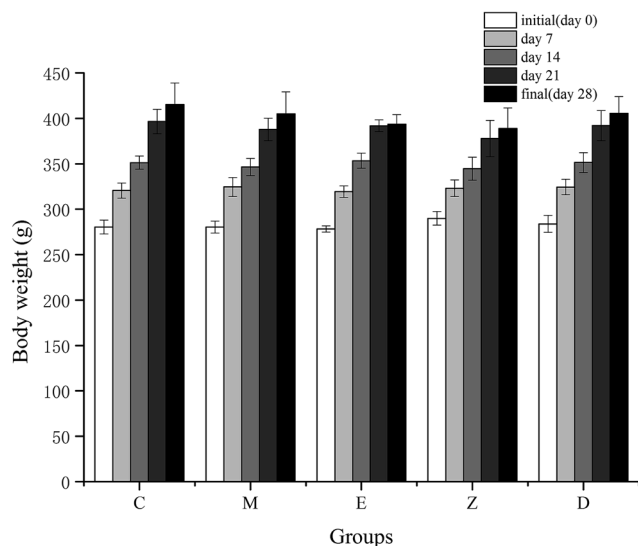


Fig. 2 DPPH radical scavenging activities of the collected fractions from RP-HPLC system at a concentration of 3 mg mL⁻¹. The data are shown as mean ± SD.



Table 4 The identification of peptides by MALDI-TOF-MS from selected RP-HPLC fraction F4^a

Amino acid sequence	Observed mass (Da)	Calculated mass ^b (Da)	Probability (%)
Leu-Ala-Arg (LAR)	360.3	358.5	98.3
Gln-Asp (QD)	263.2	261.3	99.1
Gln-Pro (QP)	244.2	243.3	99.5
His-Gln-Gly-Ile (HQGI)	453.4	453.6	90.2

^a Fractions are termed as in Fig. 1. ^b Average mass.**Fig. 3** Effect of wheat peptide on body weight changes in each group. The data are shown as mean \pm SD. C, no exercise control group; M, no exercise with low dose group; E, exercise control group; Z, exercise with low dose group; D, exercise with high dose group.

hydrolysis. The addition of protein may not have a significant effect on the synthesis of glycogen, but the excess protein and carbohydrate could be helpful in the maintenance of glycogen storage. Morifuji *et al.*³⁷ reported that the addition of glucose and whey protein hydrolysate significantly reduced the consumption of muscle glycogen compared with glucose supplementation alone during exercise. Ivy *et al.*³⁸ also showed

that the addition of carbohydrate and protein was more helpful in improving the ability of high-intensity exercise than carbohydrate supplementation alone. Supplementation of carbohydrates alone can maintain the level of muscle glycogen during exercise,³⁹ and supplementation of carbohydrates with additional protein may show better results. Bringing up the results of this study, the additional protein or protein hydrolysate alone has no significant influence on glycogen storage, but it may play a complementary role in promoting supplementation effects of carbohydrates. The reason may be the protein or protein hydrolysate is more capable of activating the key protein kinase in the skeletal muscle, which determines the absorption of glucose and the synthesis of glycogen during exercise.³⁷

In the long-term study of fatigue and injury mechanism, the relationship between generation/removal of free radical and fatigue-injury of body were paid more attention. Many studies^{23,40–44} have shown that exercise fatigue was associated with increased free radicals in the body. In the process of movement, generation and removal of free radicals play a decisive role in the levels of free radicals *in vivo*, thus affecting the occurrence and elimination of exercise-induced fatigue, and the extent of oxidative stress injury.^{23,41}

Because of the differences in the blood distribution of respective tissues during exercise, as well as different sensitivities to exercise stimulation, there were differences in free radical metabolism and oxidative damage of different tissues on the same exercise. Skeletal muscle involves strong metabolic organization in the body, which is the driving force of movement, and is the organ of maximal consumption of energy and oxygen. So, skeletal muscle represents to be the most severe lipid peroxidative tissue that is affected with acute strenuous exercise, and is the most vulnerable tissue to free radical damage. SOD is an important antioxidant enzyme in the body, which can remove oxygen free radicals. MDA is the main product of lipid peroxidation, reflecting the lipid peroxidation level of body tissues and cells. GSH-Px is an important peroxide decomposition enzyme in the body, which can protect the body from peroxide damage. In Table 7, the SOD activity of skeletal muscle in group D was significantly higher than that in groups C, M and E ($p < 0.01$), and group Z was significantly higher than that in group E ($p < 0.05$). The content of MDA in group M was significantly lower than groups C ($p < 0.05$) and E ($p < 0.01$), and group Z was significantly lower than that in group E ($p < 0.05$).

Table 5 Effects of wheat peptide on the correlation indexes of small intestine in rats^a

	C	M	E	Z	D
SOD (U mg ⁻¹)	5.31 \pm 0.45	5.96 \pm 2.08	5.46 \pm 2.57	5.59 \pm 1.33	6.43 \pm 1.34
MDA (nmol mg ⁻¹)	1.2 \pm 0.31	0.63 \pm 0.19**	0.91 \pm 0.31	1.12 \pm 0.47##	1.07 \pm 0.37##
IL-6 (pg mg ⁻¹)	11.05 \pm 11.23	23.57 \pm 15	34.26 \pm 11.03**	29.08 \pm 8.26**	11.04 \pm 4.44&&@
IL-8 (pg mg ⁻¹)	6.1 \pm 1.24	11.44 \pm 3.72**	12.59 \pm 3.5**	7.24 \pm 1.43##&	7.18 \pm 2.09##&
Caspase-3 (ng mg ⁻¹)	0.09 \pm 0.05	0.16 \pm 0.06	0.19 \pm 0.07*	0.08 \pm 0.03##&	0.05 \pm 0.03##&
sIgA (μ g mg ⁻¹)	0.23 \pm 0.04	0.3 \pm 0.06**	0.25 \pm 0.07	0.22 \pm 0.04##	0.25 \pm 0.05

^a Data are shown as mean \pm SD. * $p < 0.05$, ** $p < 0.01$ vs. C; # $p < 0.05$, ## $p < 0.01$ vs. M; & $p < 0.05$, && $p < 0.01$ vs. E; @ $p < 0.05$, @@ $p < 0.01$ vs. Z. C, no exercise control group; M, no exercise with low dose group; E, exercise control group; Z, exercise with low dose group; D, exercise with high dose group.



Table 6 Effects of wheat peptide on serum biochemical indexes in rats^a

	C	M	E	Z	D
CK (U L ⁻¹)	794.75 ± 273.71	644.44 ± 136.95	598.38 ± 263.13	718.88 ± 259.59	741.79 ± 175.28
TTE (ng mL ⁻¹)	0.51 ± 0.11	0.67 ± 0.29	0.55 ± 0.05	0.55 ± 0.03	0.6 ± 0.13
COR (ng mL ⁻¹)	296.65 ± 25.37	290.11 ± 24.34	299.37 ± 8.29	298.84 ± 17.16	285.06 ± 17.47
ET (EU mL ⁻¹)	0.71 ± 0.17	0.84 ± 0.06	0.78 ± 0.23	0.72 ± 0.14	0.64 ± 0.15 [#]
IL-6 (pg mL ⁻¹)	192.19 ± 29.15	198.23 ± 22.31	209.23 ± 17.23	197.53 ± 18.11	191.84 ± 21.65 [#]
IL-8 (pg mL ⁻¹)	44.59 ± 6.79	46.15 ± 7.74	50.82 ± 5.63	43.6 ± 10.89	42.08 ± 5.83 [#]

^a Data are shown as mean ± SD. [#]*p* < 0.05 vs. M; [&]*p* < 0.05 vs. E. C, no exercise control group; M, no exercise with low dose group; E, exercise control group; Z, exercise with low dose group; D, exercise with high dose group.

Table 7 Effects of wheat peptide on skeletal muscle tissue related indexes in rats^a

	C	M	E	Z	D
SOD (U mg ⁻¹)	3.52 ± 0.99	3.49 ± 0.66	3.32 ± 0.84	4.2 ± 0.62 ^{&}	4.62 ± 0.83 ^{**##&}
MDA (nmol mg ⁻¹)	0.82 ± 0.09	0.7 ± 0.12 [*]	0.89 ± 0.21 ^{##}	0.71 ± 0.11 ^{&}	0.78 ± 0.18
GSH-Px (U mg ⁻¹)	106.56 ± 27.07	78.28 ± 7.17	70.94 ± 13.34 [*]	93.4 ± 25.76	102.6 ± 18.69 ^{##&}
MG (mg g ⁻¹)	3.1 ± 0.88	2.49 ± 0.47	3.32 ± 0.67 [#]	3.2 ± 0.92 [#]	3.38 ± 0.77 ^{##}
IL-6 (pg mg ⁻¹)	6.61 ± 2.45	8.37 ± 0.85	7.93 ± 2.67	8.11 ± 2.76	9.37 ± 4.38

^a Data are shown as mean ± SD. ^{*}*p* < 0.05, ^{**}*p* < 0.01 vs. C; [#]*p* < 0.05, ^{##}*p* < 0.01 vs. M; [&]*p* < 0.05, ^{&&}*p* < 0.01 vs. E. C, no exercise control group; M, no exercise with low dose group; E, exercise control group; Z, exercise with low dose group; D, exercise with high dose group.

GSH-Px activity in group E was significantly lower than that in group C (*p* < 0.05), and group D was significantly higher than that in groups M (*p* < 0.05) and E (*p* < 0.01). The activities of SOD and GSH-Px in skeletal muscle of exercise group rats fed with wheat peptide were significantly higher than the exercise control groups. MDA content was significantly lower than that of exercise control group, which had an important effect on scavenging free radicals in skeletal muscle tissue. Rats not fed with wheat peptide demonstrated significantly lower activity of GSH-Px in the exercise group and the content of MDA was significantly higher than those in the quiet group. This indicated that the antioxidant enzyme system of skeletal muscle tissue was affected by exercise stress, while its function was significantly improved with wheat peptide supplementation. These results suggest that long-term exercise training would increase the level of oxidative stress in skeletal muscle and increase oxygen free radicals, and was an important factor in the occurrence of exercise-induced fatigue. The supplement of wheat peptide significantly improved the function of antioxidant enzyme system of the body, and thus the oxygen free radicals were eliminated in time.

Exhaustive experiments are the most direct way to test the exercise capacity of animals. In this study, exhaustive experiment was conducted in E, Z, and D groups, with 12 rats in each group. Results showed that the exhaustive time of group D with high dose wheat peptide supplementation was longer than group E (*p* = 0.028) and group Z (*p* = 0.055), but no significant differences between groups Z and E were observed (Fig. 4). Although there were large individual differences, the effects of high-dose wheat peptide supplementation on prolonging the exhaustive time and improving the exercise ability were obvious. The positive effects may be due to the improvement of

the situation of oxidative stress in the skeletal muscle system. Yu *et al.*¹⁴ found that soybean peptide had a strong free radical scavenging ability. Rats fed with 400 mg kg⁻¹ body weight of soybean peptide daily for 20 days demonstrated an increase of 70% in the swimming exhaustive time of rats than the control group. In combination with the results of exhaustive experiments in this study, it can be concluded that the enhancement of exercise capacity in wheat peptide nutrition group was related to the enhancement of antioxidant capacity of skeletal muscle, suggesting that anti-fatigue may be related to anti-oxidation.

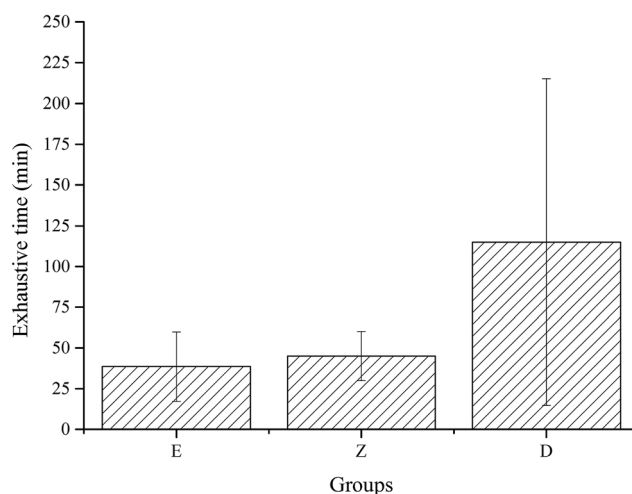


Fig. 4 Effect of wheat peptide on exhaustive time in exercise rats. The data are shown as mean ± SD. C, no exercise control group; M, no exercise with low dose group; E, exercise control group; Z, exercise with low dose group; D, exercise with high dose group.



Table 8 Effects of wheat peptide on brain tissue related indexes in rats^a

	C	M	E	Z	D
AchE ($\mu\text{g mg}^{-1}$)	23.49 \pm 5.74	25.76 \pm 6.61	21.2 \pm 6.58	29.25 \pm 3.4 ^{*&&}	27.74 \pm 5.93 [*]
5-HT (pg mg^{-1})	13.18 \pm 3.65	18.82 \pm 5.12 [*]	14.12 \pm 5.19 [#]	10.3 \pm 2.3 ^{##}	19.14 \pm 5.74 ^{*&&@}

^a Data are shown as mean \pm SD. * p < 0.05, ** p < 0.01 vs. C; # p < 0.05, ## p < 0.01 vs. M; & p < 0.05, && p < 0.01 vs. E; @ p < 0.05, @& p < 0.01 vs. Z. C, no exercise control group; M, no exercise with low dose group; E, exercise control group; Z, exercise with low dose group; D, exercise with high dose group.

Long-term exercise stress can lead to the dysfunction of central nervous system. 5-HT as a neurotransmitter is mainly involved in the neurological regulation of mood, energy, appetite, sleep, body temperature and so on. Studies found that lower levels of 5-HT were more prone to anxiety, depression, suicide, impulse, alcoholism, assault and violence.^{45,46} Acetylcholine is a common neurotransmitter in the central nervous system and the peripheral nervous system. It participates in nerve conduction in the autonomic nervous system and the kinetic nervous system. Acetylcholine is released when the nerve endings are excited by stimulation. AchE is a key enzyme in the conduction of biological nerve impulse, and mainly exists in the synaptic space of cholinergic nerve endings. AchE mainly hydrolyzes acetylcholine, terminate excitatory function of neurotransmitters on the synaptic membrane, and ensure the normal transmission of nerve signals in organisms.⁴⁷ In Table 8, the AchE of group E was significantly lower than that of groups Z (p < 0.01) and D (p < 0.05), and group C was significantly lower than that in group Z (p < 0.05). 5-HT of group M was significantly higher than that of groups C, E (p < 0.05) and Z (p < 0.01), while group D was significantly higher than that of groups C, E (p < 0.05) and Z (p < 0.01). In this study, there were no significant changes in AchE and 5-HT in the exercise control group compared with the quiet control group. But the levels of the two substances in the exercise group with wheat peptide supplementation were significantly higher than the non-supplemented group. It is possibly due to the long-term intake of glutamine-rich wheat peptide, which may improve the serotonergic system and cholinergic system of central nervous system, and also make emotions stable, ease the movement stress, and ensure the normal functioning of the central nervous system. This study demonstrated that supplementation of wheat peptide may modulate neurotransmitters in the central nervous system. According to the previous reports,¹⁶ soybean peptides regulated neurotransmitters, and also enhancing the brain function.

3.3 Effect of wheat peptides on intestinal immune function

Long-term exercise stress can cause declination in body function, affecting the immune function, and is considered as an important law of motor immunology. Especially, intestinal immune function has an important impact on the body's immune system. Intestine is the largest organ of bacteria and endotoxin repository. Under normal circumstances, the bacteria and endotoxins were harmless to the body, which is entirely dependent on the body's intestinal mucosal barrier

function. However, under stress conditions, the intestinal immune system is affected, which decrease the epithelial cell integrity, and increase the intestinal permeability. This in turn lead to intestinal bacteria and endotoxin breakthrough of the intestinal mucosal barrier into the mesenteric lymph node or portal vein system, and get into the organs away from the intestine, resulting in the translocation of intestinal bacteria, and increase the blood endotoxins, and further arising the blood immune response.^{48,49} As shown in Table 6, the endotoxin levels of exercise control group showed no significant difference compared with the quiet control group from the serum test, indicating no obvious occurrence of intestinal bacterial translocation during exercise. There was no significant difference between the nutrition group and control group of endotoxin level, which indicated that edible wheat peptide had no significant effect on the inhibition of endotoxin level and intestinal bacterial translocation. Endotoxin is a component of the Gram-negative bacterial cell wall, and its main chemical ingredient is lipopolysaccharide, which is toxic to the host. Studies have shown that antibacterial agents do not necessarily be effective against endotoxins.⁵⁰ As a result, wheat peptide might not necessarily have significant anti-endotoxin effects in the blood.

Caspase-3 is the most important terminal cleavage enzyme in the process of apoptosis and plays an irreplaceable role in apoptosis.⁵¹ In Table 5, the activity of caspase-3 was significantly higher in group E than that in group C (p < 0.05), which indicated that exercise stress could accelerate the process of intestinal epithelial cell apoptosis, and was consistent with the results of Hoffman-Goetz *et al.* study.⁵² The activity of caspase-3 in the exercise group with supplementation of wheat peptide was significantly lower than that in the control group, which indicated that wheat peptide supplementation had an important effect on the apoptosis of intestinal epithelial cells. The mechanism of wheat peptide inhibition on caspase-3 activity may be that the wheat peptide is a pseudo-substrate of caspase-3, and after cleavage by caspase-3, wheat peptide still binds to active protease and hinders the active site of caspase-3. Intestinal epithelial cells acts as an important barrier in the intestine. This study showed that exercise may have adverse effects on the intestinal barrier, while wheat peptide can protect the intestinal barrier during exercise. Ostaszewska *et al.*⁵³ showed that compared with the wheat protein supplementation alone, the mixture of wheat protein and lysine-glycine (Lys-Gly) dipeptide could play a significant inhibitory effects on the apoptosis of intestinal caspase-3 positive cells, suggesting that supplementation of small peptides may be beneficial for slowing the apoptosis of intestinal epithelial cells.



IgA is classified as serotype IgA and secretory IgA (sIgA). Serotype IgA does not show significant immune function in the serum, while sIgA is the main antibody that is produced against the local mucosal anti-infection immunity.⁵⁴ High intensity exercise showed adverse effects on the intestinal immune system, leading to changes in the ultrastructure of intestinal mucosa, decreased IgA levels, decreased intestinal mucosal immunity and digestive system function, and diminished the capacity of resistance to pathogen.⁵⁵ In addition, after high-intensity exercise, skeletal muscle protein catabolism is increased, and several amino acids were released, where glutamine accounted for the largest proportion. Under stress conditions, the demand for glutamine increases in the intestinal tissue, resulting in a sharp declination in the blood glutamine concentration. Glutamine deficiency also resulted in a significant decrease in the sIgA content in the intestine, which was associated with a decrease in the number of plasma cells secreting sIgA in the intestinal mucosa. Therefore, glutamine supplementation can significantly improve the body's immune function after high-intensity exercise.²¹ Glutamine can improve the body's immune function and protect the intestinal mucosal barrier. The most abundant amino acids in wheat peptide are glutamine and glutamic acid, which constitutes to 37.7 g/100 g in this study. Therefore, supplementation of wheat peptide may play an important role in improving the immune function of the body. Although sIgA in this study was not significantly associated with exercise, wheat peptide can significantly enhance the level of intestinal sIgA during the absence of exercise (Table 5). Zhou *et al.*⁵⁶ showed that alanyl-glutamine dipeptide significantly increased sIgA activity in calf intestine and reduced the incidence of diarrhea. Fan *et al.*⁵⁷ also found that supplementation of glutamine increased the number of intestinal lamina IgA plasma cells, and improved the immune function of the intestinal mucosa.

Interleukin is produced by activated monocytes – macrophages and lymphocytes, which is a cytokine that regulates immunity and eliminates inflammation. The main functions of interleukin-6 (IL-6) are: mediating T and B cell activation, proliferation and differentiation, induction of acute phase response during protein synthesis, promoting bone marrow hematopoietic function. Interleukin-8 (IL-8) is a strong chemotactic factor of specific and non-specific immune cells, as well as induces chemotaxis and activation of neutrophils.⁵⁸ Table 5 showed that IL-6 in group E and group Z was significantly higher than that in groups C and D ($p < 0.01$). The levels of IL-8 in groups M and E were significantly higher than those in group C ($p < 0.01$), and groups Z and D were significantly lower than those in groups M and E ($p < 0.05$ or 0.01). These results indicated that the inflammatory response of intestinal tract was more obvious after the exercise intervention, and the inflammatory reaction was obviously eliminated after supplementing wheat peptide. In addition, IL-6 and IL-8 levels of group D were significantly lower than those in group E ($P < 0.05$, Table 6). Serum test results showed that exercise did not cause significant increase in the inflammatory response in blood, but the addition of wheat peptide significantly reduced the levels of blood inflammatory factors. Teixeira-Lemos *et al.*⁵⁹ reported

that long-term high-intensity exercise may lead to systemic inflammatory response, and glutamine supplementation can reduce the inflammatory response, which is consistent with the results of our study. In addition, the activity of SOD showed no significant differences in each group. The content of MDA in group M was significantly lower than that in groups C, Z and D ($p < 0.01$, Table 5). The present study showed that the levels of SOD and MDA in the intestinal tissue without wheat peptide supplementation were not significantly changed after exercise, which means that the level of oxidative stress on the intestinal tissue was not affected by training. After wheat peptide supplementation, the level of MDA was reduced in the absence of exercise intervention, but the antioxidant effect was lost in the training groups. The above results indicated that wheat peptide could play a significant role in slowing the inflammatory response caused by long-term exercise, which may have no association with oxidative stress in the intestine.

4. Conclusion

In summary, from the results of exercise capacity and anti-fatigue ability analysis showed that long-term supplementation of wheat peptide could effectively improve the exercise capacity of one-time exhaustive exercise, remove free radicals in skeletal muscle in time, maintain long-term emotional stability, but could not improve the ability of glycogen storage in skeletal muscle or prevent skeletal muscle injury. The intestinal immune function analysis showed that long-term supplementation of wheat peptide inhibited apoptosis of intestinal epithelial cells, improved the intestinal sIgA level at rest, and alleviated the inflammatory response in the intestine and in the blood effectively. However, wheat peptide supplementation could not reflect the ability to improve the antioxidant capacity of the intestinal tract at the state of motion or prevent the displacement of bacteria.

Conflicts of interest

All the authors declare that they have no conflict of interest.

Abbreviations

SOD	Superoxide dismutase
MDA	Malondialdehyde
GSH-Px	Glutathione peroxidase
TTE	Testosterone
COR	Cortisol
CK	Creatine kinase
ET	Endotoxin
MG	Muscle glycogen
IL-6	Interleukin-6
IL-8	Interleukin-8
Caspase-3	Cysteinyl aspartate specific proteinase-3
sIgA	Secretory immunoglobulin A
5-HT	5-Hydroxytryptamine
AchE	Acetylcholinesterase



Acknowledgements

This work was supported by a key project of the Chinese Army (No. AX110C002). We would like to thank participants of this study without whom this research wouldn't have been accomplished.

References

- 1 H. Wieser, *Food Microbiol.*, 2007, **24**, 115–119.
- 2 M. C. Gianibelli, O. R. Larroque, F. MacRitchie and C. W. Wrigley, *Cereal Chem.*, 2001, **78**, 635–646.
- 3 S. Žilić, G. Akilhoğlu, A. Serpen, M. Barać and V. Gökmen, *Food Res. Int.*, 2012, **49**, 1–6.
- 4 K. X. Zhu, H. M. Zhou and H. F. Qian, *Process Biochem.*, 2006, **41**, 1296–1302.
- 5 J.-s. Wang, M.-m. Zhao, Q.-z. Zhao and Y.-m. Jiang, *Food Chem.*, 2007, **101**, 1658–1663.
- 6 J. Jia, H. Ma, W. Zhao, Z. Wang, W. Tian, L. Luo and R. He, *Food Chem.*, 2010, **119**, 336–342.
- 7 B. G. Thewissen, A. Pauly, I. Celus, K. Brijs and J. A. Delcort, *Food Chem.*, 2011, **127**, 1653–1658.
- 8 A. Pihlanto-Leppä, P. Koskinen, K. Piilola, T. Tupasela and H. Korhonen, *J. Dairy Res.*, 2000, **67**, 53–64.
- 9 H. Motoi and T. Kodama, *Nahrung*, 2003, **47**, 354–358.
- 10 C. Ziadrou, R. A. Streaty and W. Klee, *J. Biol. Chem.*, 1979, **254**, 2446–2449.
- 11 S. Fukudome, Y. Jinsmaa, T. Matsukawa, R. Sasaki and M. Yoshikawa, *FEBS Lett.*, 1997, **412**, 475–479.
- 12 I. Calzuola, F. Giavarini, P. Sassi, L. De Angelis, G. L. Gianfranceschi and V. Marsili, *Peptides*, 2005, **26**, 2074–2085.
- 13 H. J. Jeong, J. B. Jeong, D. S. Kim, J. H. Park, J. B. Lee, D.-H. Kwon, G. Y. Chung, E. W. Seo and B. O. de Lumen, *Cancer Lett.*, 2007, **255**, 42–48.
- 14 B. Yu, Z.-X. Lu, X.-M. Bie, F.-X. Lu and X.-Q. Huang, *Eur. Food Res. Technol.*, 2008, **226**, 415–421.
- 15 W. Yuanxiu, S. Xiaoyan, Z. Guixiang, S. Naxin and Z. Mingyang, *International Conference on Biomedical Engineering and Biotechnology*, 2012.
- 16 D. Yimit, P. Hoxur, N. Amat, K. Uchikawa and N. Yamaguchi, *Nutrition*, 2012, **28**, 154–159.
- 17 N. Mach and D. Fuster-Botella, *J. Sport Health Sci.*, 2017, **6**, 179–197.
- 18 R. Z. Gu, W. Y. Liu, F. Lin, Z. T. Jin, L. Chen, W. X. Yi, J. Lu and M. Y. Cai, *Food Res. Int.*, 2012, **49**, 326–333.
- 19 X. Li, J. Lin, Y. Gao, W. Han and D. Chen, *Chem. Cent. J.*, 2012, **6**, 140.
- 20 R.-Z. Gu, C.-Y. Li, W.-Y. Liu, W.-X. Yi and M.-Y. Cai, *Food Res. Int.*, 2011, **44**, 1536–1540.
- 21 L. M. Castell, *Nutrition*, 2002, **18**, 371–375.
- 22 K. Saito, D. Jin, T. Ogawa, K. Muramoto, E. Hatakeyama, A. Tadashi Yasuhara and K. Nokihara, *J. Agric. Food Chem.*, 2003, **51**, 3668.
- 23 M. B. Reid, *Free Radicals Biol. Med.*, 2008, **44**, 169–179.
- 24 K. Suetsuna and J. R. Chen, *Food Sci. Technol. Res.*, 2002, **8**, 227–230.
- 25 G. Hang, Y. Kouzuma and M. Yonekura, *Food Chem.*, 2009, **113**, 238–245.
- 26 E. Varlet-Marie, F. Maso, G. Lac and J. F. Brun, *Clin. Hemorheol. Microcirc.*, 2004, **30**, 211–218.
- 27 A. L. T. Uusitalo, M. Valkonen-Korhonen, P. Helenius, E. Vanninen, K. A. Bergstrom and J. T. Kuikka, *Int. J. Sports Med.*, 2004, **25**, 150–153.
- 28 E. P. Murolo, R. C. Derk and Y. Akgul, *Reprod. Toxicol.*, 2006, **21**, 148–153.
- 29 A. Urhausen, T. Kullmer and W. Kindermann, *Eur. J. Appl. Physiol. Occup. Physiol.*, 1987, **56**, 528–533.
- 30 A. Urhausen, H. H. Gabriel and W. Kindermann, *Med. Sci. Sports Exercise*, 1998, **30**, 407–414.
- 31 R. Rahimi, H. Rohani and M. Ebrahimi, *Apunts Medicina De Lesport*, 2011, **46**, 145–149.
- 32 L.-L. Huang, C.-T. Huang, M.-L. Chen and I. F. Mao, *Chin. J. Physiol.*, 2010, **53**, 254–261.
- 33 P. Brancaccio, N. Maffulli and F. M. Limongelli, *Br. Med. Bull.*, 2007, **81–82**, 209–230.
- 34 J. Lee, 김주영, 이철현 and 박광훈, *Korean Journal of Sport Science*, 2010, **21**, 1298–1314.
- 35 B. Knechtel, P. Knechtel, C. Mrazek, O. Senn, T. Rosemann, R. Imoberdorf and P. Ballmer, *J. Int. Soc. Sports Nutr.*, 2011, **8**, 6.
- 36 K. Sakamoto, D. E. W. Arnolds, I. Ekberg, A. Thorell and L. J. Goodyear, *Biochem. Biophys. Res. Commun.*, 2004, **319**, 419–425.
- 37 M. Morifuji, A. Kanda, J. Koga, K. Kawanaka and M. Higuchi, *Nutrition*, 2011, **27**, 833–837.
- 38 J. L. Ivy, P. T. Res, R. C. Sprague and M. O. Widzer, *Int. J. Sport Nutr. Exercise Metab.*, 2003, **13**, 382–395.
- 39 G. McConell, R. J. Snow, J. Proietto and M. Hargreaves, *J. Appl. Physiol.*, 1999, **87**, 1083–1086.
- 40 Y. Lecarpentier, *J. Appl. Physiol.*, 2007, **103**, 1917–1918.
- 41 S. K. Powers and M. J. Jackson, *Physiol. Rev.*, 2008, **88**, 1243–1276.
- 42 R. A. Nazeer, N. S. S. Kumar and R. J. Ganesh, *Peptides*, 2012, **35**, 261–268.
- 43 J.-F. Ding, Y.-Y. Li, J.-J. Xu, X.-R. Su, X. Gao and F.-P. Yue, *Food Hydrocolloids*, 2011, **25**, 1350–1353.
- 44 J. J. Wang, M. J. Shieh, S. L. Kuo, C. L. Lee and T. M. Pan, *Appl. Microbiol. Biotechnol.*, 2006, **70**, 247–253.
- 45 L. A. W. Jans, W. J. Riedel, C. R. Markus and A. Blokland, *Mol. Psychiatry*, 2007, **12**, 522–543.
- 46 G. P. A. Placidi, M. A. Oquendo, K. M. Malone, Y. Y. Huang, S. P. Ellis and J. J. Mann, *Biol. Psychiatry*, 2001, **50**, 783–791.
- 47 K. A. Modesto and C. B. R. Martinez, *Chemosphere*, 2010, **78**, 294–299.
- 48 G. A. P. Nieuwenhuijzen, E. A. Deitch and R. J. A. Goris, *Eur. J. Surg.*, 1996, **162**, 259–273.
- 49 B. J. Rowlands, C. V. Soong and K. R. Gardiner, *Br. Med. Bull.*, 1999, **55**, 196–211.
- 50 K. Brandenburg, L. Heinbockel, W. Correa and K. Lohner, *Biochim. Biophys. Acta, Biomembr.*, 2016, **1858**, 971–979.
- 51 G. Nunez, M. A. Benedict, Y. M. Hu and N. Inohara, *Oncogene*, 1998, **17**, 3237–3245.



- 52 L. Hoffman-Goetz and P. A. Spagnuolo, *J. Neuroimmunol.*, 2007, **187**, 94–101.
- 53 T. Ostaszewska, K. Dabrowski, M. Kamaszewski, P. Grochowski, T. Verri, M. Rzepkowska and J. Wolnicki, *Comp. Biochem. Physiol., Part A: Mol. Integr. Physiol.*, 2010, **157**, 158–169.
- 54 B. Moldt, K. Saye-Francisco, N. Schultz, D. R. Burton and A. J. Hessen, *Methods*, 2014, **65**, 127–132.
- 55 M. Godinez-Victoria, M. Elisa Drago-Serrano, H. Reyna-Garfias, M. Vilorio, E. Lara-Padilla, A. A. Resendiz-Albor, L. E. Sanchez-Torres, T. R. Cruz-Hernandez and R. Campos-Rodriguez, *Brain, Behav., Immun.*, 2012, **26**, 1300–1309.
- 56 Y. Zhou, P. Zhang, G. Deng, X. Liu and D. Lu, *Vet. Immunol. Immunopathol.*, 2012, **145**, 134–142.
- 57 J. Fan, G. Li, L. Wu, S. Tao, W. Wang, Z. Sheng and Q. Meng, *Nutrition*, 2015, **31**, 766–774.
- 58 S.-M. Chen, C.-P. Lin, J.-D. Tsai, Y.-H. Chao and J.-N. Sheu, *Pediatrics & Neonatology*, 2014, **55**, 120–126.
- 59 E. Teixeira-Lemos, J. Oliveira, L. P. Teixeira-Lemos, M. J. Reis-Lima and J. P. Pinheiro, in *Nutraceuticals*, Academic Press, 2016, pp. 669–714, DOI: 10.1016/B978-0-12-804305-9.00017-8.

