1 Title

- 2 Separation and Purification of antioxidant peptide from fermented whey protein by
- 3 Lactobacillus rhamnosus B2-1

4 **Running title**

5 Antioxidant peptides from fermented whey protein

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19 **ARTICLE**

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Separation and Purification of antioxidant peptide from fermented whey protein by *Lactobacillus rhamnosus* B2-1

24 Abstract

In this study, a new antioxidant activity peptide was separated and purified 25 from whey protein fermented by Lactobacillus rhamnosus. The fermentation 26 sample was separated by microporous resin D101 and Sephadex G-15. The 27 collected fractions were tested for antioxidant and antitumor activities. 28 In order to test the antioxidant activity of fractions, Hydroxyl (·OH), 2,2'-azino-29 bis (3-ethylbenzothiazoline-6-sulfonic acid (ABTS), and Oxygen Radical 30 Absorbance Capacity (ORAC) were used. The final purified peptide B11 showed 31 highest ABTS and \cdot OH radical scavenging rate by 84.36 ± 1.89% and 62.43 ± 32 2.64%, respectively, and had an ORAC activity of $1726.44 \pm 2.76 \mu M$ Trolox 33 equivalent/g at 1.0 mg/mL. Further, the effects of B11 on LoVo human colon 34 cancer cells, KB and Cal-27 human oral cancer cells were enhanced with 35 increasing concentrations of B11. B11 contains 51.421% amino acids, with Glu 36 and Asp being the major constituents. This study obtained a new bifunctional 37 peptide B11, which is promising for development. 38

39 Keywords: peptide, separation, purification, antioxidant, antitumor

40 Introduction

The presence of free radicals is known to contribute to various chronic and 41 degenerative diseases, such as diabetes, coronary heart disease, inflammation, 42 stroke, and cancer (Cheng et al., 2003). Under normal physiological conditions, 43 the body constantly generates free radicals, but also strictly regulated by the 44 antioxidant defense system, so as to maintain a healthy balance of free radicals and 45 the normal functioning of multiple body functions (Laguerre et al., 2007), 46 however, excessive accumulation of free radicals can occur when the internal 47 defense mechanism is out of balance or external harmful stimuli cause the 48 imbalance of free radicals to occur, then cause a series of diseases such as cancer 49 (Leanderson et al., 1997), cardiovascular disease, and inflammatory disease (Wang 50 et al., 2016). 51

Antioxidants are defined as any substance that prevent the adverse effects of 52 oxygen. It is a class of substances that can help capture and neutralize free radicals 53 to remove their damage to the human body (Theansungnoen et al., 2014). 54 Meanwhile, antioxidants are increasingly used in the field of dietary supplements 55 and should be widely used for disease prevention purposes. However, due to the 56 disadvantages of synthetic antioxidants such as harm to health, high cost, and 57 single mechanism of action (Shi et al., 1996), the development of green, pollution-58 free and high-safety antioxidants that can replace artificial synthesis has 59 development and application prospects. 60

Bioactive peptides (BAPs) refer to peptide compounds which are beneficial 61 to biological activities or have physiological effects. As a basic nutrient, active 62 peptides have high bioavailability and are more easily absorbed than amino acids. 63 Meanwhile, they have the function of regulating human physiological function, 64 which is not possessed by original proteins and amino acids composed of them 65 (Hartmann and Meisel, 2007). They have a variety of biological functions, such as 66 hormone effect, immune regulation, anti-hypertension, antitumor effect, 67 antioxidant effect, cholesterol lowering, inhibition of bacteria and viruses, etc. 68 (Samaei et al., 2021; Sarabandi and Jafari, 2020). Among them, antioxidant 69 peptides act as antioxidants by scavenging free radicals and relieving oxidative 70 stress (Giordano et al., 2017; Pandya et al., 2019; Yi et al., 2022). It is found that 71 certain bioactive peptides do indeed have antioxidative properties and can be used 72 as a natural substitute to synthetic antioxidants to improve health (Nadalian et al., 73 2019). Due to their low toxicity and high efficiency, antioxidant peptides are 74 widely used by the product processing industry and in health care products 75 (Tadesse and Emire, 2020; Wen et al., 2020). 76

Food protein from dairy products by fermentation method is a source of antioxidant peptides (Anukam and Reid, 2009). The fermentation method uses dairy products as raw materials, inserts protease-producing microorganisms, and utilizes protease produced in the fermentation process of microorganisms to hydrolyze milk protease into peptides with different amino acid sequences and molecular weights under aerobic or anaerobic conditions (Agbor et al., 2011;

83	Chandrasekara and Shahidi, 2011), which are microbial cells themselves, or direct
84	metabolites or secondary metabolites (Ali et al., 2019; Li et al., 2012). Lactic acid
85	bacteria (LAB) are the dominant group of protease-producing microorganisms,
86	among which Lactobacillus spirochetes, Lactobacillus bulgaricus and
87	Lactobacillus paracasei have been reported (Sun et al., 2010). The abundance and
88	characteristics of antioxidant peptides released from milk and dairy products are
89	strain-dependent (Ayyash et al., 2020; Rubak et al., 2020).
90	Due to LAB's ability to release bioactive peptides in fermented dairy
91	products, and the fact that it is "generally recognized as safe" (GRAS) for use in
92	food, LAB has become more common for certain strains to produce fermented
93	dairy products with certain functional properties (Kim et al., 2017). Therefore, the
94	objective of this study was to obtain antioxidant peptides using Lactobacillus
95	rhamnosus B2-1 from fermented whey protein. It has important practical
96	significance for developing new whey protein products and improving the
97	comprehensive utilization rate of whey protein.
98	

99 Materials and Methods

100	Materials

Dimethyl Sulfoxide (DMSO) was purchased from Sangon Biotech (Shanghai) Co., Ltd. 2,2-amino-di (2-ethyl-benzothiazoline sulphonic acid-6) ammonium salt (ABTS), 3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) and Trolox were purchased from Sigma-Aldrich Co. Fetal bovine serum, penicillin streptomycin, DMEM, and dimethyl sulfoxide were purchased from Thermo
Scientific. Macroporous resin D101 was purchased from Amicogen Biopharm Co.,
Ltd. Sephadex G-15 gel was purchased from GE healthcare Bio-Sciences AB Co.
Ltd. All other chemicals and reagents used were of analytical grade.

109

Culture medium

Whey protein-MRS medium: According to the modified MRS (DeMan Rogosa-Sharpe) medium, the specific components are as follows: desalinated whey powder 10 g, beef extract 10 g, yeast extract 5 g, glucose 20 g, Twain's 80 1 mL, dipotassium hydrogen phosphate 2 g, sodium acetate 5 g, sodium citrate 2 g, magnesium sulfate 0.2 g, manganese sulfate 0.05 g, add water to 1 L, 121°C, sterilization for 15 min.

116

Lactic acid bacteria strain

Lactobacilli was obtained from the soaking liquid of corn. The soaking 117 solution was inoculated into 5% desalted whey powder medium and cultured for 118 48 h, centrifuged at 3500 r/min for 10 min, and the supernatant was taken for 119 preliminary screening. After initial screening, the bacteria fermentation broth was 120 expanded and enriched at 37°C for 48 h. Single colonies were isolated and 121 selected after spreading the serially diluted fermentation broth on MRS agar plate 122 and incubating them at 37°C for 48 h. The selected strains were streaked on MRS 123 agar plate and observed by Meilan staining. 124

By extracting the genome of the strain with properties, and identifying the 16S rRNA by Bioengineering Biotechnology (Shanghai) Co. Ltd., the data was

127 submitted to National Biotechnology Information Center by basic local queue128 search methods for homology comparison.

129

Fermentation and sample preparation

130 Lactobacillus rhamnosus was cultured in 1 mL of sterilized whey protein-

131 MRS medium and incubated under anaerobic conditions at 37°C for 48 h. After

132 continuous expansion for 2 times, the culture was inoculated into 5% whey

protein-MRS medium for 48 h, centrifuged at 6000 r/min for 10 min, and the

134 supernatant was freeze-dried.

135

Peptide isolation and purification

Peptides for whey protein fermentation (PWPF) were preliminary isolated by 136 macroporous resin D101. PWPF (20 mL, 150 mg/mL) were loaded on 137 macroporous resin D101 column (25 mm \times 460 mm), and stepwise eluted with a 138 series of gradient ethanol solutions at a flow rate of 2.5 mL/min. Automatic 139 collectors were used to collect the eluate (2 min/tube). In order to obtain the 140 elution curve of the sample solution, the absorbance of the fractions was measured 141 at 214 nm, 254 nm and 280 nm. Besides, the peptide content and ABTS radical 142 clearance rate were measured in every three tubes. The absorption peaks were 143 collected, pooled, rotary evaporation and lyophilized. The purified fractions, 144 labeled F1, F2, F3, and F4, were stored at -20°C until needed. 145 A Sephadex G-15 chromatography was used to purify the fraction with strong 146

radical scavenging activity. The column was pre-equilibrated with ultrapure water.
 The purified fraction was formulated into a 0.7 mg/mL solution with deionized

water, and then the peptide solution (4 mL) was loaded onto the column (16 mm ×
100 cm). Deionized water was used to elute the column at a volume of 4 mL/tube,
with absorbance monitoring at 214 nm, 254 nm and 280 nm. The fractions were
lyophilized for further use.

153

Tricine SDS-PAGE

The concentrations of acrylamide in the decomposing and stacking gels were 154 16 and 4%, respectively. The sample was mixed with one volume of $1 \times$ sample 155 buffer [12% SDS (w/v), 6% mercaptoethanol (v/v), 30% glycerol (w/v), 0.05% G-156 250 (w/v) and 150 mM Tris/HCl (pH 7.0)]. After boiling for 3 min, the samples 157 were cooled and loaded onto acrylamide gels, which were pre-run at 60 V for 20 158 min before the sample was applied. The electrophoresis was performed by 159 stacking and breaking down the gel at constant voltages of 60 V and 100 V, 160 respectively. After electrophoresis, the gel was removed, the fixative was fixed 161 for 30 min, stained for 1 h, and it was decolorated. 162

163

Amino acid composition

The sample was mixed with 10mL of 6 mol/L HCl (containing 0.1% phenol), and nitrogen was filled into the prepared hydrolysis tube to protect the samples from oxidation during the hydrolysis process. The hydrolysis tube was hydrolyzed at 110°C under vacuum for 24 h. After drying, sodium citrate buffer solution (1 mL, pH 2.2) was added into the test tube and dissolved. The absorbed solution was transferred to a sample bottle for detection by the amino acid analyzer after passing through a 0.22 μ m filter membrane.

ABTS radical scavenging activity

Antioxidant activity was measured by measuring the radical-scavenging ability of ABTS. The sample (40 μ L, 1 mg/mL) was mixed with the ABTS reagent (160 μ L) and reacted at room temperature for 6 min in the dark. The absorbance was measured with a microplate reader at 734 nm. As a blank control, 50 mmol/L of Tris-HCl buffer was used, and Vc as a positive control. The calculation formula is:

ABTS radical scavenging ability
$$/\% = \left(1 - \frac{A_{\text{sample}}}{A_{\text{blank}}}\right) \times 100\%$$

179

Oxygen radical absorbance capacity

To determine the oxygen radical scavenging ability, ORAC method was used. The sample (20 μ L) was added to FL working solution (80 μ L), followed by 200 mmol/L AAPH added quickly. The fluorescence intensity of each well was measured every 6 min at 37°C at excitation wavelength of 485 nm and emission wavelength of 538 nm using a microplate analyzer. The measurement time is usually set after the fluorescence attenuation has reached the baseline. With Trolox as positive control, the calculation formula is as follows:

Relative ORAC value =
$$\left[\frac{(AUC_{sample} - AUC_{blank})}{AUC_{Trolox} - AUC_{blank}}\right] \times (c_{Trolox}/\omega_{sample})$$

188

AUC_{sample} and AUC_{blank} in the formula represent sample solution and blank
 solution respectively.

191 Hydroxyl radical scavenging activity

The hydroxyl (\cdot OH) radical scavenging activity was determined using the salicylic acid method. The sample (2 mL) was incubated with ferrous sulfate solution (0.5 mL, 9 mmol/L), salicylic acid solution (2 mL, 9 mmol/L in absolute ethanol) and H₂O₂ solution (8.8 mmol/L) at 37°C for 30 min, and the absorbance was measured at 510 nm. In the control, deionized water was used as a substitute for H₂O₂, while deionized water was also used for the blank, and Vc was used as a positive control. The calculation formula is:

¹⁹⁹ ·OH radical scavenging activity
$$/\% = \left(1 - \frac{A_{\text{sample}} - A_{\text{blank}}}{A_{\text{control}}}\right) \times 100\%$$

200

201 Cell lines and culture conditions

LoVo human colon cancer cells, KB human oral cancer cells and Cal-27 human oral cancer cells were purchased from the Type Culture Collection of the Chinese Academy of Sciences, Shanghai, China. Three types of cells were cultivated in DMEM containing 10% FBS, ampicillin sodium, and streptomycin sulfate. All cells were cultivated at 37°C in a humidified atmosphere containing 5% CO₂.

208

Cell proliferation assay

MTT assay was used to evaluate the effects of samples on cell proliferation. Cells were harvested during logarithmic growth phase and seeded at a density of 2 $\times 10^4$ cells per well in 96-well plates. During overnight growth, the culture medium was replaced by various concentrations of the sample for 48 h. MTT was added to each well for 4 h. The supernatant was removed and dissolved in 150 µL of DMSO. ELISA plate readers were then used to measure absorption at 570 nm.

²¹⁵ Cell inhibition rate (percentage) was calculated using the following formula:

216

217

Inhibition rate/ %= $\left(1 - \frac{A_{\text{sample}}}{A_{\text{blank}}}\right) \times 100\%$

218 Statistical analysis

Statistical analyses were performed using the variance test analysis (SPSS Statistics 19 software). Significant differences between the groups with different treatments were examined using T-test. P < 0.05 was set as statistically significant difference.

223

224 **Results**

Identification of the fermented bacterial strain

In the experiment, the fermented bacterial strain was isolated from the soaking liquid of corn. The strain isolate was genotyped by using 16S rRNA gene sequences. The alignment of 16S rRNA gene sequences showed 99.15% identity with *Lacticaseibacillus rhamnosus* strain NBRC 3425. *Lactobacillus rhamnosus* B2-1 was identified based on these results. All the raw reads have been submitted to the NCBI SRA under the accession number SRR19749980.

232

Macroporous resin technology

The lyophilized powder from the fermentation broth of whey protein

metabolites of *Lactobacillus rhamnosus* B2-1 was separated by macroporous resin

235 D101 column, and its elution was detected by ultraviolet detector. As can be seen

from the Fig. 1A, four peaks were successively separated, namely F1, F2, F3 and
F4. The yield of each component was 6.03%, 21.50%, 16.66% and 9.69%,
respectively.

The results of Tricine-SDS-PAGE electrophoresis were shown in Fig. 1B. Controlling with α -lactoglobulin and β -chylous protein, the results showed that the molecular weight of F1 mainly concentrated around 35 kD, while the molecular weight of F3 and F4 mainly concentrated around 14 kD, 20 kD and 35 kD, showing obvious bands in these three molecular weights.

244

Amino acid composition of fractions

The amino acid compositions of F1, F2, F3 and F4 are represented in Table 1. 245 Of the four fractions, the highest amino acid content was F4, followed by F3, F2 246 and F1. The results of amino acid compositions are in good agreement with the 247 results of Tricine-SDS-PAGE of the above fractions. F4 was found to contain both 248 essential and non-essential amino acids, with the main amino acids being Glu, Pro, 249 Thr, Asp and Ile. Hydrophobic amino acids (Ala, Val, Met, Ile, Leu, Phe, Pro, and 250 Tyr) were 8.738% of F4 (Qian et al., 2008). Thus, F4 contained the highest 251 proportion of hydrophobic amino acids. 252

253

Antioxidant activities of fractions

Antioxidant activity of protein fractions was determined by ABTS radical scavenging activity, oxygen radical absorbance capacity and •OH radical scavenging activity.

Fig. 2A shows that ABTS radical scavenging activity of protein fractions. As shown in Fig. 2A, the ABTS radical scavenging activity of F4 was $34.46 \pm 1.71\%$, which was significantly higher than other fractions at the same concentration. The scavenging rate of ABTS radical was F4 > F1 > F3 > F2. These results indicated that the F4 and F1 has potential antioxidant abilities.

Fig. 2B shows that the results of the fluorescence natural decay curves of 262 protein fractions. Based on Fig. 2B, the antioxidant capacity of ORAC was 263 measured for different fractions of the same concentration and compared with the 264 positive control Trolox that was used to calculate ORAC value. The relative 265 ORAC values of F1, F2, F3 and F4 were $1060.36 \pm 1.52 \mu$ M Trolox equivalent/g, 266 $704.82 \pm 1.55 \ \mu\text{M}$ Trolox equivalent/g, $631.00 \pm 0.96 \ \mu\text{M}$ Trolox equivalent/g and 267 $1388.80 \pm 2.34 \,\mu\text{M}$ Trolox equivalent/g, respectively. The results showed that F4 268 and F1 have good oxygen radical absorbance capacity. 269

Fig. 2C shows that \cdot OH radical scavenging activity of protein fractions. As shown in Fig. 2C, \cdot OH radical scavenging rates of F1, F2, F3 and F4 were 22.27 ± 1.89%, 16.38 ± 0.86%, 26.82 ± 1.45% and 35.63 ± 1.11%, respectively. Overall, \cdot OH radical scavenging rates of each fraction were F4 > F3 > F1 > F2 in descending order. These data suggested that the fraction F4 protected cells effectively against damage and oxidative lipid peroxidation.

276

Antitumor activity of fractions

On Fig. 3A–C, the effect of various concentrations on the inhibition of LoVo cells (A), KB cells (B), and Cal-27 cells (C) can be seen. With the increase of

279	concentration, the inhibitory effect of F1 and F4 on tumor cells was significantly
280	enhanced, while F2 and F3 were less effective than F1 and F4. At 8 mg/mL, F1
281	inhibited the growth of LoVo cells, KB cells, and Cal-27 cells by $58.28 \pm 0.89\%$,
282	21.71 \pm 0.53%, and 53.86 \pm 2.43%, respectively. At the same concentration, the
283	inhibition rate of F4 was 66.23 \pm 0.61%, 57.14 \pm 1.67%, and 47.91 \pm 1.05%,
284	respectively. These results confirmed that F1 and F4 inhibited cancer cell growth
285	at low concentrations.

Gel filtration chromatography

The fraction of F4 from macroporous resin D101 column with the highest antitumor and antioxidation effects was separated by gel filtration chromatography (Sephadex G-15). As shown in Fig. 4, two fractions (B11 and B2) were separated from F4. Due to the excessive conductivity of fraction B2, which contained more salt and less polypeptides, B11 was only used as the study material.

292

Amino acid composition of antioxidant peptide

The results of the amino acids composition are presented in Table 1. The 293 amino acid content of B11 is 51.421%, nearly three times that of F4 (18.949%). 294 Acidic amino acids Glu and Asp, which are capable of scavenging radicals via 295 proton transfer, were the primary amino acids contained in the peptide, accounting 296 for 11.180% and 6.013%, respectively. There were seven kinds of essential amino 297 acids (Thr, Val, Met, Ile, Leu, Phe, and Lys) in B11 accounting for 21.871% of 298 total amino acids. Therefore, B11 not only had a high antioxidant activity, but also 299 had higher nutritional value. 300

Antioxidant activities of antioxidant peptide

Fig. 5A shows that ABTS radical scavenging activity of B11. The ABTS radical clearance rates of B11 increased with the increase of sample concentration after purification, and when the concentration was 1.0 mg/mL, the ABTS radical scavenging rate was the highest, and the antioxidant activity was the strongest, reached $70.36 \pm 1.89\%$. In the control group, the ABTS radical scavenging rate gradually increased with increasing concentration, reaching 98% at the maximum concentration.

Fig. 5B shows that B11 has strong antioxidant capacity in a dose-dependent manner. The B11 (1.0 mg/mL) showed a high oxygen radical absorbance activity of 1726.44 \pm 2.76 μ M Trolox equivalent/g. The ORAC values for the remaining concentrations (0.4, 0.6 and 0.8 mg/mL) of B11 were 465.66 \pm 3.05 μ M Trolox equivalent/g, 982.84 \pm 1.65 μ M Trolox equivalent/g and 1614.07 \pm 2.41 μ M Trolox equivalent/g, respectively.

Fig. 5C shows that \cdot OH radical scavenging activity of B11. As shown in Fig. 5C, B11 demonstrated significant \cdot OH radical scavenging activity, in which the activity increased significantly with increasing concentration, but all concentrations were significantly weaker than Vc. At 1.0 mg/mL, B11 demonstrated scavenging activity of $62.43 \pm 2.64\%$ against \cdot OH radicals. At other concentrations, B11 displayed scavenging activity of $51.04 \pm 1.95\%$, $46.33 \pm 2.07\%$, $28.63 \pm 2.33\%$, respectively.

322 Antitumor activity of antioxidant peptide

323	As shown in Fig. 5D, the inhibition rate of cell proliferation gradually
324	increased with increasing concentrations of B11 from 0.4 to 1.0 mg/mL, which
325	demonstrates a dose-dependent relationship. For LoVo cells, the inhibition rate of
326	B11 from 0.4 to 1.0 mg/mL was 20.78 \pm 1.65%, 30.47 \pm 1.63%, 44.41 \pm 1.56%,
327	$52.36 \pm 2.36\%$, respectively. At this concentration, the cell activities of KB cells
328	were $13.67 \pm 1.78\%$, $18.14 \pm 3.87\%$, $30.21 \pm 2.34\%$, and $48.42 \pm 3.42\%$,
329	respectively, while that of Cal-27 cells were $18.97 \pm 2.87\%$, $24.10 \pm 2.96\%$, 35.74
330	\pm 2.78%, and 54.57 \pm 2.14%, respectively. Based on the above data, it can be can
331	conclude that compared with the other tumor cells, B11 has a significant
332	inhibitory effect on the proliferation of LoVo cells.

334 **Discussion**

Antioxidant peptides can be prepared through fermentation using microbial 335 enzyme systems, to date, most studies have primarily concentrated on obtaining 336 antioxidant peptides by fermenting whey protein with potential probiotics. Cui et 337 al. (Cui et al., 2022) found that the polypeptides of whey protein generated by 338 Lactobacillus reuteri WQ-Y1 had the potential antioxidant activity. Virtanen et al. 339 (Virtanen et al., 2007) investigated the production of antioxidant activity during 340 fermentation with 25 lactic acid bacterial strains, and all strains showed the 341 antioxidant activity. In the present study, Lactobacillus rhamnosus B2-1 isolated 342 from the soaking liquid of corn was selected as the fermentation strain, and 343 peptides with antioxidant activity were also obtained. It might mean that the 344

antioxidant activity of peptides depends on the fermenting strains and the
development of antioxidant activity is strain-specific.

The polypeptides produced by fermentation are generally a mixture, so it is 347 particularly important to separate the purified parts. Macroporous resin adsorption 348 technology and gel filtration chromatography have been widely used for natural 349 products purification (Puchalska et al., 2015; Yan et al., 2015). Macroporous resin 350 has the characteristics of distributed apertures, easy desorption regeneration 351 ability, selectivity on the adsorbate and other features (Xu et al., 2010). As 352 different volume fractions of ethanol have different polarity and affinity for 353 antioxidant peptides with different hydrophobicity, gradient elution may produce 354 antioxidant peptides with different hydrophobicity, and the physiological activity 355 of these antioxidant peptides is closely related to their hydrophobicity. Therefore, 356 in this study, the four fractions obtained by gradient elution with different volume 357 fractions of ethanol had different antioxidant activities. The results showed that 358 the most hydrophobic elution fraction F4 was indeed the best antioxidant activity 359 among the four fractions, so it was completely feasible to use macroporous resin 360 for preliminary separation and purification. Some researchers have reported that 361 macroporous resin D101 can be used to separate and obtain hydrophobic active 362 ingredients (Gao et al., 2018; Gu et al., 2019). 363

With the rise and in-depth study of antioxidant peptides, a variety of methods have been established to evaluate the activity of antioxidant peptides *in vitro*. The antioxidant activity of peptides may be classified into two groups, depending on

the chemical reaction involved: 1) those peptides are based on the hydrogen atom 367 transfer reaction, which are mainly analyzed by kinetic curves, and 2) those 368 peptides are based on electron transfer reactions, which are mainly reflected by 369 color changes (Moayedi et al., 2017; Ohata et al., 2016; Yang et al., 2017b). In the 370 first case, it consists of ORAC assay and ABTS radical scavenging assay, while in 371 the second case it consists of ·OH and DPPH radical scavenging assays (Shen et 372 al., 2010; Tang et al., 2010). In this study, three in vitro assays were used to 373 determine the antioxidant activity of protein fractions. The results showed that the 374 antioxidant activity of the target peptide B11 gradually increased as the fermented 375 solution was separated by different separation methods. 376

Researchers reported that most antioxidant peptides contained one or more 377 hydrophobic amino acids since hydrophobic amino acid residues can act as 378 hydrogen donors to break off free radical peroxide chain reactions with their 379 aromatic residues (Xing et al., 2016). In the fraction B11, 23.349% of the amino 380 acids were hydrophobic, which would lead to better oxidation inhibition than in 381 F4. In this study, the content of Met and Tyr in B11 was higher than that of other 382 fractions (F1, F2, F3 and F4). Polypeptides with Met and Tyr could effectively 383 scavenge free radicals (Yang et al., 2017a), which was also one of the reasons for 384 385 the improved ABTS free radical scavenging ability of B11. Wattanasiritham et al. (Wattanasiritham et al., 2016) reported that peptides containing Pro, Gly, Ala, Val, 386 and Leu had intrinsic antioxidant activity. Amino acids such as His, Leu, Gly and 387 Pro were suggested to play an important role in radical caging (Scholljegerdes et 388

al., 2005). Moreover, peptides containing Tyr, Pro, Phe, and His, displayed strong 389 antioxidant properties (Ren et al., 2008; Shen et al., 2010). The results of the 390 present study supported the above statement by the strong antioxidant properties 391 of B11 with the presence of intrinsic antioxidant amino acids. As a whole, the 392 strong antioxidant properties of B11 may be attributed to their constituent amino 393 acids, which can provide protons or act as electron donors to the free radicals and 394 react with them to convert them to more stable products and terminate radical 395 chain reaction (Zielinski et al., 2009). All of these reasons may explain the strong 396 antioxidant activity of B11. Results from this study indicated that it was feasible to 397 produce natural antioxidants from the whey protein fermented by Lactobacillus 398 rhamnosus. 399

Cells have an innate ability to modulate apoptosis using various redox 400 systems, in which cancer cells are more dependent on antioxidant systems than 401 other non-transformed cells, so they are especially vulnerable to increased 402 oxidative stress as this redox balance is frequently deregulated in cancer cells 403 (Eliassen et al., 2002; Trachootham et al., 2009). It was identified that antioxidants 404 had been proposed as potential candidates in the prevention and treatment of 405 diseases associated with active oxygen species, particularly cancer diseases (Leng 406 407 et al., 2005). In this study, the results showed that B11 could significantly inhibit the proliferation of cancer cells, indicating that that the antioxidant and 408 antiproliferative activities of the peptide were interrelated. Therefore, it might be 409 proposed that higher antioxidant activity leads to better antitumor activity, and the 410

possible reason was the initial antitumor effect by regulating the level of oxidative
stress in cancer cells (Kannan et al., 2008). Similar results were observed
previously, Sun et al. (Sun et al., 2017) demonstrated that the polypeptide from *P*. *eryngii* mycelium exerted significant antioxidant effects and could inhibit the
growth of cervical, breast, and gastric cancer cells, which was also a bifunctional
bioactive peptide.

417

418 Conclusion

In summary, our study reports the antioxidant activity of peptides derived 419 from the whey protein fermented by Lactobacillus rhamnosus B2-1 by 420 macroporous resin and gel chromatography purification for the first time. After 421 whey protein fermented from Lactobacillus rhamnosus, the obtained fermented 422 solution was separated by macroporous resin D101 to four fractions (F1, F2, F3 423 and F4). Fraction F4 showed significantly greater antioxidant activity than the 424 other three fractions. F4 was re-separated using the Sephadex G-15 425 chromatography, and the final antioxidant fraction, B11, was obtained. Fraction 426 B11 exhibited potent in vitro antioxidant activity in ABTS, •OH radical 427 scavenging activity and ORAC assay, and antitumor activity on LoVo, KB and 428 Cal-27 cancer cells proliferation. The high antioxidant activity of B11 can be 429 attributed to the presence of intrinsic antioxidant amino acids. Thus, B11 might be 430 a good source of antioxidant peptide that may act as a food additive and bioactive 431 material, and open up new possibilities for the utilization of whey by-products. 432

433	
434	Conflicts of interest
435	There are no conflicts to declare.
436	
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442	Author contributions
443	Conceputualization: Hong Guan. Data curation: Huo Hao. Formal analysis:
444	Lei Fan. Methodology: Lin Ding. Validation: Wenqin Yang. Investigation:
445	Chuangang Zang. Writing-Original draft: Hao Guo.
446	
447	Ethics Approval
448	This article does not require IRB/IACUC approval because there are no
449	human and animal participants.
450	
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564 Figure Legends

- 565
- 566 FIGURE 1. Macroporous resin D101 column chromatography (25 mm × 460 mm)
- ⁵⁶⁷ and Tricine-SDS-PAGE results of each fraction.
- (A) Macroporous resin D101 column chromatography. Eluent: A solution (water),
- ⁵⁶⁹ B solution (ethanol). Flow rate: 2.5 mL/min, Fraction: 2 mL/tube. Fractionation:
- ⁵⁷⁰ F1 (tube on, 10-16), F2 (tube on, 19-34), F3 (tube on, 67-79), F4 (tube on, 80-88).
- (B) M: Marker, 1: F1, 2: F2, 3: F3, 4: F4, 5: Sterilized whey protein, 6: PWPF, 7:
- 572 Unsterilized whey protein, α : α -lactoglobulin protein, β : β -chylous protein.

573

- 574 FIGURE 2. Antioxidant activity of each fraction at a concentration 1 mg/mL.
- 575 (A) ABTS radical clearance rate.
- 576 (B) Fluorescence natural decay curves.
- 577 (C) Hydroxyl radical clearance rate.
- 578

579 FIGURE 3. Effect of various concentrations of each fraction on cancer cells

- 580 inhibition rate.
- 581 (A) LoVo cells.
- 582 (B) KB cells.
- 583 (C) Cal-27 cells.

584

585 FIGURE 4. Chromatogram profiles of separation of F4 by Sephadex G-15 (16 mm

- 586 × 100 cm). Eluent: Water. Flow rate: 0.5 mL/min, Fraction: 4 mL/tube,
- ⁵⁸⁷ Fractionation: B11 (tube on, 14-18), B2 (tube on, 22-25).

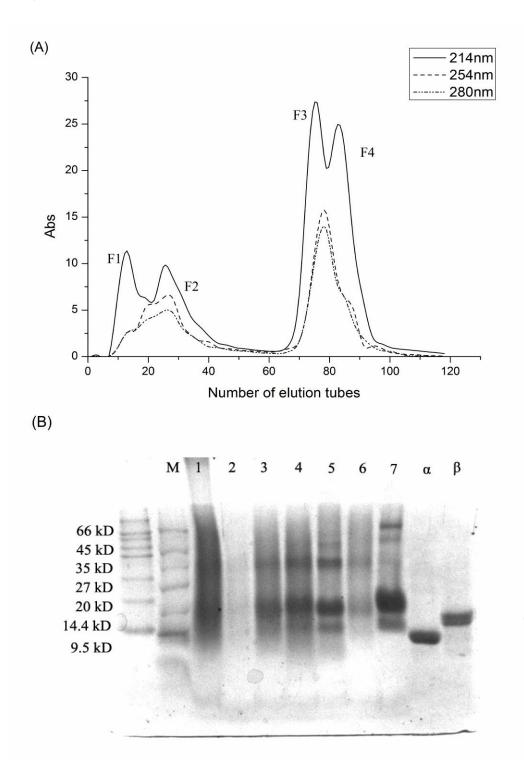
- 589 FIGURE 5. Antioxidant and antitumor activity of the fraction B11.
- 590 (A) ABTS radical clearance rate.
- 591 (B) Fluorescence natural decay curves.
- 592 (C) Hydroxyl radical clearance rate.
- 593 (D) Effect of various concentrations on cancer cells inhibition rate.

594 Table

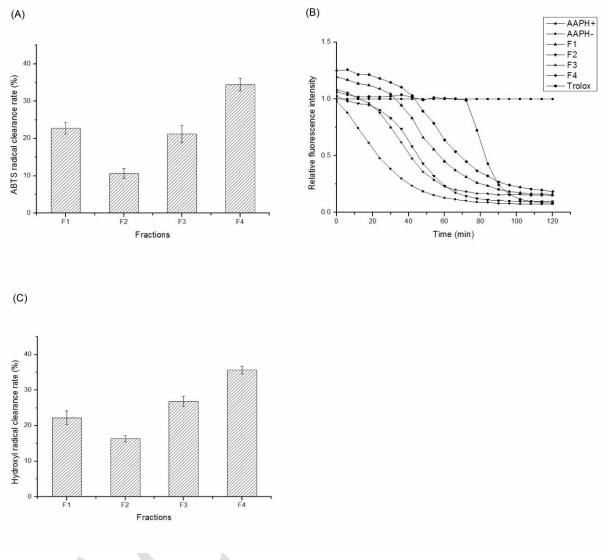
595	Table 1. Amino aci	d compositior	n of each fra	ction.	
	Amino acids		Content	ts of amino ad	cids (%)
	Ammo actus	E 1	БJ	E2	E4

Amino soids	Contents of amino acids (%)				
Amino acids -	F1	F2	F3	F4	B11
Asp	0.311	0.031	0.808	2.145	6.013
Thr	0.183	0.017	0.553	2.346	4.281
Ser	0.157	0.017	0.373	1.320	2.950
Glu	0.555	0.058	1.434	4.551	11.180
Gly	0.070	0.016	0.129	0.437	1.011
Ala	0.152	0.019	0.344	0.961	2.865
Cys	0.023	0.004	0.088	0.084	0.575
Val	0.156	0.016	0.392	1.125	3.208
Met	0.052	0.005	0.095	0.410	1.229
Ile	0.149	0.014	0.463	2.021	3.608
Leu	0.308	0.025	0.693	1.078	5.629
Tyr	0.037	0.003	0.185	0.213	1.533
Phe	0.102	0.009	0.211	0.484	1.641
His	0.185	0.075	0.210	0.925	2.181
Lys	0.073	0.009	0.197	0.679	2.275
Arg	0.036	0.004	0.058	0.169	1.243
Pro	0.165	0.165	0.468	2.446	3.636
Total amino acid	2.550	0.322	6.234	18.949	51.421













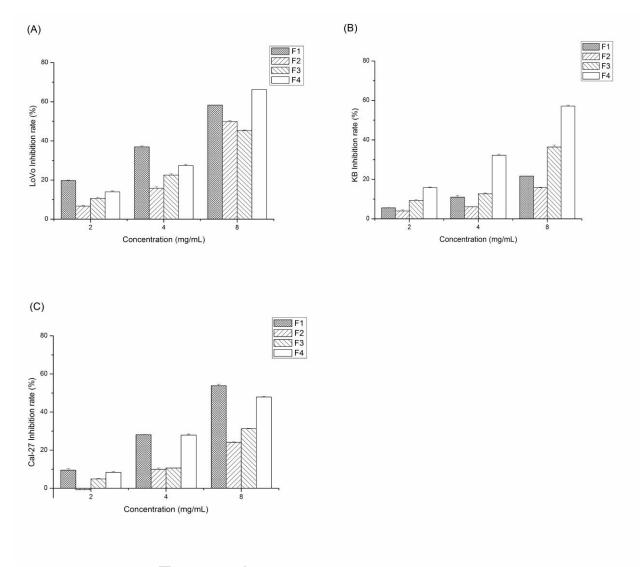




Figure 4

