

Supplementary material for

An innovative germination-driven LAB fermentation strategy for allergenicity reduction in soymilk gels through spatiotemporal protease-mediated epitope disruption

Yaqiong Wang, Shijie Fang, Zhihao Fan, Yi Wang, Yue Wu, Xin Rui, Pei Wang,
Chong Xie, Runqiang Yang *

Sanya Institute of Nanjing Agricultural University, Whole Grain Food Engineering
Research Center, College of Food Science and Technology, Nanjing Agricultural
University, Nanjing, Jiangsu, 210095, China

* Corresponding authors contact information:

Runqiang Yang

College of Food Science and Technology, Nanjing Agricultural University

1 Weigang Road, Nanjing, Jiangsu, China

Tel: +86 13813927131; Fax: +86 25 84396293;

E-mail address: yangrq@njau.edu.cn

Supplementary methods

1. Antigenicity assay

100 μ L aliquots of each sample were dispensed into the wells of a microtiter plate and allowed to incubate at 25°C for 10 min. Subsequently, the wells were meticulously washed three times. Following this, 100 μ L of a horseradish peroxidase-conjugated secondary antibody was introduced into each well, and the plates were incubated again at the same temperature for an additional 10 min. After washing over three times, 100 μ L of substrate was added to each well and incubated for 10 min. The reaction was terminated by the addition of 100 μ L of stop solution, and the absorbance was then measured at a wavelength of 450 nm. The antigenicity was then calculated using a specific formula:

$$\text{Antigenicity (\%)} = \frac{\text{OD}_{450} [\text{S(S)M}]}{\text{OD}_{450} (\text{SM})} \times 100\%$$

2. IgE-binding capacity

Sera samples were obtained from individuals with varying severity of soybean allergies, supplied by Chongqing Wolcavi Biological Technology Co., Ltd., Chongqing, China. Prior consent was secured from all patients, and their sera were used in the study with appropriate acknowledgment. The patients included were aged 21, 51, 47, 31, and 28 years, with specific IgE levels of 67.5, 5.11, 6.07, 15.4, and 21.4 KU/L, respectively.

For the assay, 100 μ L of each serum sample was dispensed into wells of a 96-well ELISA plate and allowed to incubate overnight. The plates were then washed and treated with BSA-PBS to prevent nonspecific binding. Next, the plates were incubated

with diluted human serum and an anti-human IgE-peroxidase antibody. After further incubation and washing, 3,3',5,5'-Tetramethylbenzidine substrate was added, and the enzymatic reaction was halted with sulfuric acid. Finally, the absorbance was measured at a wavelength of 450 nm.

3. Peptidomic identification

Initially, peptides were extracted from F(S)SM samples through the addition of trichloroacetic acid, followed by centrifugation at $12,000 \times g$ for 15 min. The resulting supernatant, containing the peptide mixture, was then subjected to desalting by using a MonoSpin C₁₈ desalination column. Once desalted, the peptide solution was acidified with 0.1% formic acid (FA) and loaded onto a nano trap column. The elution process occurred at a flow rate of 4 $\mu\text{L}/\text{min}$, utilizing a loading buffer composed of 2% acetonitrile. Subsequently, the peptide solution was eluted onto an analytical column using a gradient elution ranging from 3% to 45% of buffer B (acetonitrile containing 0.1% FA) at a flow rate of 0.3 $\mu\text{L}/\text{min}$. The entire elution procedure lasted for 190 min. MS/MS acquisition in a data-dependent manner was performed utilizing an LTQ-Orbitrap XL mass spectrometer, with the spray voltage adjusted to 2.2 kV. The raw MS data files were subsequently processed using MaxQuant software (version 1.6.2.6), and these data were searched against the protein sequences available on the UniProt database. Carbamidomethylation of cysteine was set as a fixed modification, with protein N-terminal oxidation of methionine as a variable modification.

Supplementary Results

Table S1. The specific information of soy allergies.

Patients	S1	S2	S3	S4	S5
Classification			Allergic		
Skin prick Test type			Positive		
Methodology			ImmunoCAP		
			Fluorometric Enzyme Immunoassay Analysis (FEIA)		
Ethnicity	Asian	Caucasian	Other	Caucasian	Asian
Gender	Male	Male	Female	Female	Male
Blood Type	Not Known	Not Known	A+	A+	Not Known
PLI					
Permanent ID	33776-LL	33906-AH	34507-HA	34945-ND	28472-JT
Bleed Number	PL 26569	PL 27321	PL 31516	PL 29834	PL 28282
Birth Date	1973, 02, 01	1987, 09, 13	2003, 01, 16	1996, 10, 22	1993, 10, 14
Draw Date	2019, 12, 17	2021, 01, 08	2023, 10, 18	2022, 11, 09	2021, 11, 09
Diagnosis	Allergic rhinitis, Filariasis, No immunotherapy	Allergic rhinitis, Multi-allergic, No immunotherapy, Vaccine for SARS-COV-2	Allergic rhinitis, No immunotherapy	Allergic rhinitis, Allergy diagnosed by a physician, No immunotherapy, No prior history of cancer, Patient has clinically relevant allergic symptomatology	Allergic rhinitis, Allergy diagnosed by a physician, Atopic dermatitis, Eczema, Elevated IgE, Multi-allergic, No prior history of cancer, Patient has clinically relevant allergic symptomatology, Shingles History, Yes-immunotherapy

1 **Table S2. Particle size distribution of the fermented soymilk (FSM) and**
 2 **fermented sprouted seeds-derived soymilk (FSSM) collected at different *L.***
 3 ***plantarum* strains and fermented terminal pH values**

Fermentation conditions	Samples	D[4, 3]
<i>L. Plantarum</i> O3, pH 5.5	FSM	23.8 ± 0.4 μm ^d
	FSSM-1	21.3 ± 0.3 μm ^g
	FSSM-2	17.8 ± 0.4 μm ⁱ
	FSSM-3	12.5 ± 0.3 μm ⁿ
<i>L. Plantarum</i> P7, pH 5.5	FSM	22.4 ± 0.6 μm ^{ef}
	FSSM-1	15.9 ± 0.2 μm ^j
	FSSM-2	14.9 ± 0.2 μm ^l
	FSSM-3	11.3 ± 0.3 μm ^{pq}
<i>L. Plantarum</i> T7, pH 5.5	FSM	36.5 ± 1.2 μm ^a
	FSSM-1	32.8 ± 0.4 μm ^b
	FSSM-2	22.3 ± 0.2 μm ^f
	FSSM-3	12.2 ± 0.1 μm ^{no}
<i>L. Plantarum</i> O3, pH 4.5	FSM	23.1 ± 0.2 μm ^e
	FSSM-1	19.3 ± 0.4 μm ^h
	FSSM-2	11.7 ± 0.3 μm ^{op}
	FSSM-3	10.5 ± 0.3 μm ^{qr}
<i>L. Plantarum</i> P7, pH 4.5	FSM	20.6 ± 0.4 μm ^g
	FSSM-1	15.5 ± 0.1 μm ^k
	FSSM-2	11.4 ± 0.1 μm ^p
	FSSM-3	10.0 ± 0.2 μm ^r
<i>L. Plantarum</i> T7, pH 4.5	FSM	37.3 ± 0.5 μm ^a
	FSSM-1	26.9 ± 0.4 μm ^c
	FSSM-2	17.5 ± 0.2 μm ⁱ
	FSSM-3	12.0 ± 0.1 μm ^o

	FSM	20.6 ± 0.5 μm ^g
<i>L. Plantarum</i> O3, pH 4.0	FSSM-1	13.9 ± 0.5 μm ^m
	FSSM-2	10.9 ± 0.2 μm ^q
	FSSM-3	10.1 ± 0.1 μm ^r
	FSM	20.5 ± 0.5 μm ^g
<i>L. Plantarum</i> P7, pH 4.0	FSSM-1	13.8 ± 0.3 μm ^m
	FSSM-2	10.6 ± 0.4 μm ^{qr}
	FSSM-3	10.0 ± 0.3 μm ^r
	FSM	32.4 ± 0.3 μm ^b
<i>L. Plantarum</i> T7, pH 4.0	FSSM-1	22.8 ± 0.4 μm ^{ef}
	FSSM-2	15.1 ± 0.3 μm ^{kl}
	FSSM-3	11.6 ± 0.1 μm ^p

4 ¹ Values after ± represent the standard deviations.

5 ^{2 a-r} Values followed by different letters in the same column are significantly different ($p < 0.05$).

Table S3. The distrution of major allergenic epitope degradation of FS(S)M gel derived from soybeans at different germination times.

Soy allergen (Accession number)	Region	Allergenic epitope	Samples
Gly m 5.01 (P0DO16)	Region 1	E ₁₁₆ -E ₁₂₄	FSSM-1, FSSM-2, and FSSM-3
		H ₁₁₇ -E ₁₄₂	
	Region 2	S ₁₃₇ -E ₁₄₄	FSSM-3
		N ₂₀₁ -R ₂₁₅	
Gly m 5.02 (P11827)	Region 3	E ₃₆₂ -E ₃₇₈	FSSM-3
	Region 4	K ₃₈₁ -D ₄₀₃	FSSM-1, FSSM-2, and FSSM-3
		E ₃₈₂ -N ₄₀₇	
	Region 1	E ₁₁₈ -E ₁₂₄	FSSM-2 and FSSM-3
	Region 2	Q ₁₅₁ -E ₁₅₈	FSSM-1, FSSM-2, and FSSM-3
	Region 3	E ₁₇₃ -D ₁₈₈	FSSM-2 and FSSM-3
Region 4	H ₂₁₃ -R ₂₁₈	FSSM-3	
Gly m 5.03 (P25974)	Region 5	F ₅₇₅ -K ₅₈₀	FSSM-3
		L ₁₃₄ -A ₁₄₉	
	Region 1	D ₁₃₉ -K ₁₅₄	FSSM-1, FSSM-2, and FSSM-3
		K ₁₄₄ -D ₁₅₉	
		L ₁₄₈ -R ₁₅₇	
		A ₁₄₉ -S ₁₆₄	
	Region 2	Q ₂₀₃ -I ₂₁₀	FSSM-3
		R ₂₀₄ -R ₂₁₉	
V ₂₀₉ -R ₂₂₄			
V ₂₁₁ -I ₂₁₈			
	S ₂₁₄ -S ₂₂₉	FSSM-2 and FSSM-3	
	R ₂₁₉ -S ₂₃₄		

	Region 5	T ₂₉₂ -P ₂₉₇	
Gly m 8 (P19594)	Region 1	N ₃₉ -I ₅₀	FSSM-1, FSSM-2, and FSSM-3
	Region 2	Q ₉₅ -T ₉₉	
Gly m TI (P01070)	Region 1	R ₅₅ -G ₆₀	FSSM-1, FSSM-2, and FSSM-3
	Region 2	I ₈₉ -L ₁₀₀	FSSM-1, FSSM-2, and FSSM-3
		A ₉₃ -F ₁₀₂	
	Region 1	S ₁₁₅ -Q ₁₂₉	FSSM-3
Gly m Bd 30K (P22895)	Region 2	H ₂₀₂ -V ₂₀₉	FSSM-2 and FSSM-3
		C ₂₃₀ -I ₂₄₁	
	Region 3	N ₂₃₃ -Y ₂₄₄	FSSM-3
	Region 4	Q ₂₃₆ -L ₂₄₇	
		D ₂₅₁ -F ₂₆₂	FSSM-3
		T ₂₅₄ -A ₂₆₅	
Region 5	P ₂₉₆ -S ₃₁₀	FSSM-2 and FSSM-3	

Table S4. The number of epitopes and their distribution of major soybean allergen.

Soybean allergen	Number of epitopes	Amino acid sequence	Sencondary structure
Gly m 1	9	R ₅ -L ₁₁ , L ₁₈ -V ₂₅ , V ₂₅ -C ₂₉ , C ₂₈ -G ₃₃ , I ₃₈ -C ₄₅ , L ₄₄ -A ₅₀ , C ₆₇ -Y ₇₁ , R ₆₉ -A ₇₅ , N ₇₄ -R ₇₉	α-helix and random coil
Gly m 2	14	R ₄ -F ₉ , L ₁₅ -G ₂₈ , F ₁₈ -V ₂₃ , F ₁₈ -S ₃₃ , S ₂₀ -S ₃₅ , V ₂₄ -H ₃₆ , Q ₂₅ -H ₃₉ , S ₃₃ -H ₃₉ , S ₃₃ -A ₄₉ , H ₃₉ -H ₄₆ , R ₄₄ -C ₅₂ , L ₅₀ -E ₅₅ , E ₅₅ -G ₆₀ , S ₆₆ -F ₇₀	α-helix, β-sheet, and random coil
Gly m 3	7	A ₂₃ -D ₂₉ , D ₂₉ -S ₃₆ , W ₃₃ -G ₈₈ , T ₁₁₁ -N ₁₁₆ , V ₁₁₉ -L ₁₂₆ , G ₁₂₅ -D ₁₃₀ , I ₁₂₇ -Y ₁₃₁	α-helix, β-sheet, and random coil
Gly m 4	33	E ₉ -P ₁₃ , N ₁₁ -A ₁₅ , P ₁₃ -A ₁₇ , K ₃₉ -V ₄₄ , E ₄₂ -G ₄₆ , N ₄₃ -G ₄₆ , G ₄₉ -E ₆₃ , I ₅₃ -V ₆₇ , T ₅₇ -I ₇₁ , T ₅₇ -F ₆₆ , F ₅₈ -T ₆₄ , D ₆₁ -D ₇₅ , G ₆₂ -V ₆₇ , I ₇₄ -A ₇₇ , I ₇₄ -Y ₈₃ , A ₇₇ -L ₉₁ , L ₇₉ -Y ₈₃ , S ₈₂ -V ₈₆ , S ₈₄ -G ₈₇ , V ₈₅ -A ₉₀ , K ₉₇ -G ₁₁₁ , V ₁₀₅ -K ₁₁₉ , N ₁₀₉ -K ₁₁₅ , Y ₁₂₀ -G ₁₂₄ , E ₁₂₁ -D ₁₂₅ , G ₁₂₄ -P ₁₂₈ , A ₁₂₆ -E ₁₃₂ , Q ₁₃₀ -K ₁₃₄ , L ₁₃₃ -I ₁₄₇ , K ₁₃₇ -L ₁₅₁ , A ₁₄₂ -D ₁₅₆ , H ₁₅₄ -Y ₁₅₇	α-helix, β-sheet, and random coil
Gly m 7	9	K ₁₆ -R ₂₄ , K ₂₂ -K ₂₇ , H ₁₀₁ -R ₁₁₇ , K ₁₁₀ -A ₁₁₆ , G ₁₆₃ -K ₁₇₀ , E ₄₃₂ -L ₄₃₈ , Q ₄₄₅ -P ₄₅₃ , E ₄₅₀ -G ₄₅₈ , A ₆₃₄ -G ₆₃₉	α-helix and random coil
2S albumin (Gly m 8)	32	L ₁₂ -S ₂₀ , S ₂₀ -H ₂₆ , Q ₂₇ -R ₃₂ , S ₃₀ -L ₃₅ , C ₃₁ -P ₄₂ , K ₃₃ -N ₃₉ , N ₃₉ -I ₅₀ , N ₃₉ -K ₄₅ , I ₆₇ -I ₇₈ , T ₇₀ -K ₈₁ , E ₈₇ -C ₉₇ , E ₉₀ -L ₁₀₄ , H ₉₃ -P ₁₀₇ , Q ₉₅ -T ₉₉ , K ₉₆ -Q ₁₁₀ , T ₉₉ -A ₁₁₃ , T ₉₉ -L ₁₀₄ , S ₁₀₂ -A ₁₁₃ , S ₁₀₂ -K ₁₁₆ , K ₁₀₅ -D ₁₁₉ , I ₁₀₈ -S ₁₂₂ , K ₁₀₈ -I ₁₁₇ , C ₁₁₁ -L ₁₂₅ , C ₁₁₁ -K ₁₁₆ , N ₁₂₀ -K ₁₂₈ , E ₁₂₇ -I ₁₃₈ , K ₁₃₂ -F ₁₄₆ , K ₁₃₅ -M ₁₄₉ , I ₁₃₈ -C ₁₅₂ , A ₁₄₁ -S ₁₅₅ , M ₁₄₃ -P ₁₄₈ , I ₁₅₀ -S ₁₅₅	α-helix and random coil
Kunitz trypsin	6	R ₅₅ -G ₆₀ , I ₈₉ -L ₁₀₀ , A ₉₃ -F ₁₀₂ , C ₁₁₁ -T ₁₁₆ , T ₁₁₆ -V ₁₂₁ ,	β-sheet and random coil

inhibitor (Gly m TI) Lectin (Gly m Agglutini n)	4	S ₂₀₅ -R ₂₁₅ Q ₁₃₄ -Y ₁₃₉ , F ₁₆₀ -D ₁₆₅ , F ₂₆₀ -H ₂₆₆ , H ₂₆₆ -D ₂₇₂	β -sheet and random coil
7S α subunit (Gly m 5.01)	31	F ₁₅ -I ₂₅ , S ₄₈ -A ₅₆ , A ₅₆ -L ₆₁ , E ₆₄ -G ₇₁ , E ₆₇ - E ₉₅ , P ₇₆ -E ₈₆ , R ₁₀₂ -R ₁₂₇ , E ₁₁₆ -E ₁₂₄ , H ₁₁₇ -E ₁₄₂ , S ₁₃₇ -E ₁₄₄ , Q ₁₅₂ -R ₁₇₇ , D ₁₇₂ -L ₁₉₇ , E ₁₇₄ -E ₁₈₀ , S ₁₈₅ -G ₁₉₉ , N ₁₉₂ -F ₂₂₀ , K ₁₉₃ -F ₂₀₇ , N ₂₀₁ -R ₂₁₅ , N ₂₀₉ - R ₂₂₃ , L ₂₁₇ -R ₂₃₁ , Q ₂₂₆ -T ₂₄₄ , L ₂₂₇ -D ₂₅₂ , T ₂₆₃ -T ₂₉₁ , Y ₂₉₃ -S ₃₂₁ , F ₃₂₂ -E ₃₄₃ , E ₃₆₂ -E ₃₇₈ , K ₃₈₁ -D ₄₀₃ , E ₃₈₂ -N ₄₀₇ , S ₄₀₁ -N ₄₃₀ , D ₄₁₂ - D ₄₃₇ , N ₅₂₃ -Q ₅₄₃ , Q ₅₇₃ -R ₆₀₂ Y ₂₇ -Q ₃₁ , Q ₃₁ -K ₃₇ , R ₄₀ -E ₄₅ , K ₆₂ -Q ₇₃ , V ₆₃ - I ₇₄ , R ₇₆ -K ₉₂ , Q ₈₈ -P ₁₀₂ , E ₉₄ -R ₁₀₁ , D ₉₅ -R ₁₀₉ , P ₁₀₆ -Q ₁₁₀ , Q ₁₁₀ -E ₁₁₅ , E ₁₁₈ -E ₁₂₄ , K ₁₂₈ -G ₁₃₃ , S ₁₃₇ -E ₁₄₅ , E ₁₄₅ -H ₁₅₀ , Q ₁₅₁ -E ₁₅₈ , H ₁₆₀ -H ₁₆₉ , E ₁₇₃ -E ₁₇₇ , E ₁₇₇ - K ₁₈₈ , E ₁₉₀ -P ₂₀₃ , H ₂₁₃ -R ₂₁₈ , R ₂₁₈ -F ₂₂₃ , N ₂₂₅ -R ₂₃₁ , N ₂₃₇ -Q ₂₄₄ , S ₂₄₀ -R ₂₄₇ , L ₂₆₃ - D ₂₆₈ , S ₂₉₂ -Q ₂₉₆ , L ₃₀₁ -G ₃₀₆ , M ₃₂₂ -R ₃₃₄ , F ₃₃₉ -S ₃₅₄ , G ₃₇₅ -Q ₃₈₀ , E ₃₈₄ -E ₃₉₄ , K ₃₉₈ -L ₄₀₃ , T ₄₄₃ -D ₄₅₁ , D ₄₅₁ -K ₄₇₄ , V ₄₅₄ -D ₄₆₀ , E ₄₆₃ - F ₄₆₇ , I ₄₈₁ -N ₄₈₇ , I ₄₉₃ -Q ₄₉₈ , N ₅₃₃ -D ₅₃₇ , D ₅₈₁ - K ₅₈₇ , F ₅₇₅ -K ₅₈₀ , K ₅₈₇ -S ₅₉₂ , A ₅₉₇ -Q ₆₀₁ F ₁₆ -S ₂₃ , L ₂₄ -S ₃₉ , E ₂₈ -R ₃₇ , D ₂₉ -T ₄₄ , Y ₃₅ - N ₄₈ , S ₃₉ -R ₅₄ , T ₄₄ -F ₅₉ , Q ₄₉ -P ₆₄ , Q ₄₉ -Q ₅₇ , R ₅₄ -L ₆₉ , L ₅₅ - F ₅₉ , F ₅₉ -I ₇₄ , S ₆₃ -N ₆₈ , S ₆₃ -R ₇₀ , E ₆₇ -D ₇₁ , Y ₇₂ -K ₈₀ , I ₇₄ - H ₈₉ , I ₈₄ -L ₉₉ , H ₈₉ -I ₁₀₄ , F ₉₇ -L ₁₀₅ , L ₉₉ -D ₁₁₄ , A ₁₀₃ -R ₁₁₃ , I ₁₀₄ -H ₁₁₉ , S ₁₁₅ -P ₁₂₀ , H ₁₁₉ -L ₁₃₄ , D ₁₂₂ -P ₁₂₇ , I ₁₂₆ -K ₁₄₄ , G ₁₂₉ -K ₁₄₄ , Y ₁₃₃ -N ₁₄₂ , L ₁₃₄ -A ₁₄₉ , D ₁₃₉ -K ₁₅₄ , K ₁₄₄ - D ₁₅₉ , L ₁₄₈ -R ₁₅₇ , A ₁₄₉ -S ₁₆₄ , K ₁₅₄ -D ₁₅₉ , Y ₁₅₈ -R ₁₉₄ , Q ₁₇₄ - F ₁₈₉ , L ₁₈₁ -R ₁₉₄ , S ₁₈₄ -E ₁₉₉ , N ₁₉₃ -G ₁₉₈ , V ₁₉₅ -R ₂₀₄ , V ₁₉₅ -	α -helix, β -sheet, and random coil
7S α' subunit (Gly m 5.02)	44	E ₁₁₈ -E ₁₂₄ , K ₁₂₈ -G ₁₃₃ , S ₁₃₇ -E ₁₄₅ , E ₁₄₅ -H ₁₅₀ , Q ₁₅₁ -E ₁₅₈ , H ₁₆₀ -H ₁₆₉ , E ₁₇₃ -E ₁₇₇ , E ₁₇₇ - K ₁₈₈ , E ₁₉₀ -P ₂₀₃ , H ₂₁₃ -R ₂₁₈ , R ₂₁₈ -F ₂₂₃ , N ₂₂₅ -R ₂₃₁ , N ₂₃₇ -Q ₂₄₄ , S ₂₄₀ -R ₂₄₇ , L ₂₆₃ - D ₂₆₈ , S ₂₉₂ -Q ₂₉₆ , L ₃₀₁ -G ₃₀₆ , M ₃₂₂ -R ₃₃₄ , F ₃₃₉ -S ₃₅₄ , G ₃₇₅ -Q ₃₈₀ , E ₃₈₄ -E ₃₉₄ , K ₃₉₈ -L ₄₀₃ , T ₄₄₃ -D ₄₅₁ , D ₄₅₁ -K ₄₇₄ , V ₄₅₄ -D ₄₆₀ , E ₄₆₃ - F ₄₆₇ , I ₄₈₁ -N ₄₈₇ , I ₄₉₃ -Q ₄₉₈ , N ₅₃₃ -D ₅₃₇ , D ₅₈₁ - K ₅₈₇ , F ₅₇₅ -K ₅₈₀ , K ₅₈₇ -S ₅₉₂ , A ₅₉₇ -Q ₆₀₁ F ₁₆ -S ₂₃ , L ₂₄ -S ₃₉ , E ₂₈ -R ₃₇ , D ₂₉ -T ₄₄ , Y ₃₅ - N ₄₈ , S ₃₉ -R ₅₄ , T ₄₄ -F ₅₉ , Q ₄₉ -P ₆₄ , Q ₄₉ -Q ₅₇ , R ₅₄ -L ₆₉ , L ₅₅ - F ₅₉ , F ₅₉ -I ₇₄ , S ₆₃ -N ₆₈ , S ₆₃ -R ₇₀ , E ₆₇ -D ₇₁ , Y ₇₂ -K ₈₀ , I ₇₄ - H ₈₉ , I ₈₄ -L ₉₉ , H ₈₉ -I ₁₀₄ , F ₉₇ -L ₁₀₅ , L ₉₉ -D ₁₁₄ , A ₁₀₃ -R ₁₁₃ , I ₁₀₄ -H ₁₁₉ , S ₁₁₅ -P ₁₂₀ , H ₁₁₉ -L ₁₃₄ , D ₁₂₂ -P ₁₂₇ , I ₁₂₆ -K ₁₄₄ , G ₁₂₉ -K ₁₄₄ , Y ₁₃₃ -N ₁₄₂ , L ₁₃₄ -A ₁₄₉ , D ₁₃₉ -K ₁₅₄ , K ₁₄₄ - D ₁₅₉ , L ₁₄₈ -R ₁₅₇ , A ₁₄₉ -S ₁₆₄ , K ₁₅₄ -D ₁₅₉ , Y ₁₅₈ -R ₁₉₄ , Q ₁₇₄ - F ₁₈₉ , L ₁₈₁ -R ₁₉₄ , S ₁₈₄ -E ₁₉₉ , N ₁₉₃ -G ₁₉₈ , V ₁₉₅ -R ₂₀₄ , V ₁₉₅ -	α -helix, β -sheet, and random coil
7S β subunit (Gly m 5.03)	75	H ₈₉ -I ₁₀₄ , F ₉₇ -L ₁₀₅ , L ₉₉ -D ₁₁₄ , A ₁₀₃ -R ₁₁₃ , I ₁₀₄ -H ₁₁₉ , S ₁₁₅ -P ₁₂₀ , H ₁₁₉ -L ₁₃₄ , D ₁₂₂ -P ₁₂₇ , I ₁₂₆ -K ₁₄₄ , G ₁₂₉ -K ₁₄₄ , Y ₁₃₃ -N ₁₄₂ , L ₁₃₄ -A ₁₄₉ , D ₁₃₉ -K ₁₅₄ , K ₁₄₄ - D ₁₅₉ , L ₁₄₈ -R ₁₅₇ , A ₁₄₉ -S ₁₆₄ , K ₁₅₄ -D ₁₅₉ , Y ₁₅₈ -R ₁₉₄ , Q ₁₇₄ - F ₁₈₉ , L ₁₈₁ -R ₁₉₄ , S ₁₈₄ -E ₁₉₉ , N ₁₉₃ -G ₁₉₈ , V ₁₉₅ -R ₂₀₄ , V ₁₉₅ -	α -helix, β -sheet, and random coil

			<p>K₂₁₅, L₁₉₆-E₂₀₂, Q₂₀₃-I₂₁₀, R₂₀₄-R₂₁₉, V₂₀₉-R₂₂₄, V₂₁₁-I₂₁₈, S₂₁₄-S₂₂₉, R₂₁₉-S₂₃₄, R₂₂₄-P₂₃₉, S₂₃₄-Y₂₄₉, P₂₃₉-G₂₅₄, S₂₄₄-I₂₅₉, Y₂₄₉-N₂₆₄, G₂₅₄-D₂₆₉, I₂₅₉-L₂₇₄, N₂₆₄-I₂₇₉, D₂₆₉-L₂₈₄, D₂₆₉-K₂₉₂, I₂₇₂-N₂₈₀, L₂₇₄-F₂₈₉, I₂₇₉-I₂₉₄, F₂₈₉-A₃₀₄, V₂₉₅-E₃₀₁, A₃₄₄-L₃₅₉, A₃₄₅-V₃₅₀, S₃₅₄- Q₃₆₉, A₃₇₄-Q₃₈₉, N₃₇₉-P₃₉₄, Q₃₈₉-L₄₀₄, P₃₉₄-E₄₀₉, D₃₉₉-D₄₁₄, L₄₀₄-Q₄₁₉, Q₄₁₉-I₄₃₄, S₄₂₄-Y₄₃₉ T₃₅-S₄₉, L₆₅-N₇₉, I₇₅-R₈₉, P₈₅-T₉₉, K₁₀₅- K₁₁₉, S₁₁₅-Q₁₂₉, P₁₃₅-Q₁₄₉, E₁₉₂-G₂₀₀, E₁₉₃-E₂₀₇, N₁₉₉- W₂₀₈, H₂₀₂-V₂₀₉, K₂₂₆-T₂₄₀, C₂₃₀-I₂₄₁, N₂₃₃-Y₂₄₄, Q₂₃₆-L₂₄₇, D₂₄₂-S₂₅₃, D₂₅₁-F₂₆₂, T₂₅₄-A₂₆₅, P₂₉₆-S₃₁₀, E₃₂₅-T₃₃₉, A₃₆₆-L₃₇₉</p>	
P34 (Gly m Bd 21 34K)				<p>α-helix, β-sheet, and random coil</p>
P28 (Gly m Bd 3 28K)			<p>G₃₁-S₄₅, L₅₉-I₇₃, E₁₅₇-H₁₇₁</p>	<p>α-helix, β-sheet, and random coil</p>
11S glycinin G1 (Gly m 6.01)	34		<p>S₁₄-F₂₀, Q₂₅-Q₃₄, P₄₁-E₄₆, R₄₄-G₄₉, N₈₆- E₉₀, I₁₀₃-E₁₁₃, P₁₀₅-S₁₂₃, E₁₁₃-G₁₂₁, G₁₂₁-D₁₂₈, P₁₂₆-Q₁₃₁, I₁₄₂-G₁₄₇, E₁₉₁-G₂₀₀, F₁₉₂-E₁₉₈, Q₁₉₆-Q₂₀₃, H₂₀₂-G₂₀₇, K₂₀₈-E₂₁₃, G₂₁₇-L₂₂₉, S₂₂₂- D₂₃₆, G₂₂₃-F₂₂₈, G₂₂₃-V₂₃₅, F₂₂₈-A₂₃₂, E₂₃₀-D₂₃₆, N₂₄₂-E₂₄₆, Q₂₄₄-E₂₄₈, G₂₅₃-I₂₆₅, L₂₆₂-P₂₆₇, P₂₆₈-Q₂₇₄, Q₂₇₄-E₂₇₉, E₂₈₁-K₂₈₇, E₂₈₆-C₂₉₀, K₂₉₃-C₂₉₇, E₃₁₄-T₃₁₈, F₃₇₀-L₃₇₆, F₄₈₂-K₄₉₁ M₁-G₁₅, K₃-F₁₇, C₉-A₂₃, L₁₂-A₁₈, L₁₉- Q₃₁, N₄₀-G₄₆, S₄₄-I₄₉, K₅₇-A₆₅, K₅₇-L₇₁, A₆₅-P₇₉, R₇₃- E₈₇, Y₈₁-G₉₅, N₈₃-F₉₇, Y₈₉-G₁₀₃, Q₉₁-P₁₀₅, F₉₇-P₁₁₁, I₁₀₀-Y₁₀₈, E₁₁₀-Q₁₁₆, Q₁₁₂-G₁₁₈, G₁₁₈-Q₁₂₈, Q₁₂₁- K₁₂₉, Q₁₂₈-F₁₃₃, V₁₃₀-V₁₄₁, G₁₃₆-E₁₅₃, P₁₄₂-W₁₄₇, E₁₆₉-Q₁₇₄, N₁₇₀-N₁₈₄, R₁₇₈-Y₁₉₂, E₁₈₆-L₁₉₀, E₁₈₆-S₂₀₀, E₁₈₈-Q₁₉₇, F₁₈₉-Q₁₉₅, L₁₉₀-K₂₀₄, Q₁₉₄-Q₂₀₈, Q₁₉₅- Q₂₀₁, G₁₉₉-G₂₀₅, G₂₁₄-F₂₂₅, S₂₁₉-N₂₃₃, S₂₄₉-A₂₆₃, K₂₅₆-V₂₆₁, G₂₅₇-Q₂₇₁, Q₂₈₃-</p>	<p>α-helix, β-sheet, and random coil</p>
11S glycinin G2 (Gly m 6.02)	53			<p>α-helix, β-sheet, and random coil</p>

C₂₉₁, G₃₂₉-W₃₄₃, L₃₃₇-Y₃₅₁, L₃₄₅-M₃₅₉,
 S₃₅₃-N₃₆₇, M₃₅₉-Y₃₇₃, V₃₆₁-L₃₇₅, N₃₆₇-
 V₃₈₁, N₃₆₉-V₃₈₃, V₄₄₉-Q₄₆₃, Q₄₅₁-K₄₆₅,
 K₄₅₇-S₄₇₁

11S		C ₁₇ -F ₂₂ , F ₂₀ -Q ₂₇ , R ₈₁ -T ₈₅ , E ₉₀ -G ₉₆ , Y ₉₂ - I ₉₉ , E ₁₁₃ -G ₁₁₈ ,	
glycinin	22	E ₁₁₃ -H ₁₂₇ , Q ₁₁₅ -R ₁₂₂ , D ₁₂₅ -F ₁₃₃ , I ₁₃₀ -E ₁₃₅ , I ₁₃₉ -F ₁₄₅ ,	β -sheet and random coil
G3 (Gly m 6.03)		F ₁₆₈ -Q ₁₇₄ , P ₁₇₆ -Y ₁₈₀ , R ₁₇₇ -L ₁₉₀ , G ₁₈₃ - Q ₁₈₇ , F ₁₈₉ -Q ₁₉₅ , G ₂₅₉ -P ₂₆₆ , P ₂₆₆ -Q ₂₇₂ , R ₂₇₄ -P ₂₈₁ , E ₂₇₈ -E ₂₈₅ , Q ₂₉₁ -N ₂₉₆ , T ₃₁₄ -F ₃₂₀ E ₃₁ -N ₃₆ , L ₄₀ -H ₄₄ , V ₄₆ -G ₅₁ , N ₅₇ -K ₇₂ , S ₅₈ - L ₆₃ , G ₆₇ -K ₇₂ ,	
11S		N ₇₈ -P ₈₃ , K ₁₂₈ -D ₁₃₃ , N ₁₄₂ -V ₁₄₈ , G ₁₅₂ -A ₁₆₇ , Q ₂₁₅ -Q ₂₂₀ ,	α -helix, β -sheet, and random coil
glycinin	23	I ₂₆₂ -Q ₂₇₇ , G ₂₆₈ -P ₂₇₄ , Q ₂₇₉ -E ₂₈₄ , D ₂₉₂ -K ₃₀₇ , G ₃₃₆ -E ₃₄₁ , P ₃₆₄ -R ₃₇₀ , A ₃₉₄ -A ₃₉₉ , A ₃₉₄ -I ₄₀₉ , A ₃₉₉ -S ₄₁₄ , S ₄₁₄ -Y ₄₂₉ , Q ₄₂₁ -Y ₄₂₈ , A ₅₀₈ -Y ₅₁₂ N ₇₇ -H ₈₂ , N ₇₉ -S ₈₅ , L ₈₃ -Y ₈₉ , Y ₈₆ -Q ₉₁ , A ₁₀₅ - P ₁₁₀ ,	
11S		G ₁₀₈ -E ₁₁₄ , H ₁₄₀ -G ₁₄₄ , L ₁₄₇ -G ₁₅₂ , E ₁₆₃ -I ₁₆₈ , D ₁₉₄ -T ₂₀₀ , K ₂₁₄ -E ₂₂₂ , G ₂₂₆ -L ₂₃₇ , K ₂₅₀ - D ₂₅₆ , S ₂₇₂ -Q ₂₇₆ , W ₂₇₅ -D ₂₈₂ , D ₂₈₄ -G ₂₉₀ ,	α -helix, β -sheet, and random coil
glycinin	33	T ₂₉₂ -P ₂₉₇ , R ₂₉₉ -H ₃₀₅ , E ₃₁₁ -P ₃₂₀ , E ₃₁₃ -P ₃₂₄ , Q ₃₁₇ -H ₃₂₂ , P ₃₂₀ -Q ₃₂₅ , Q ₃₂₅ -E ₃₃₄ , Q ₃₃₂ - G ₃₃₇ , R ₃₃₆ -Q ₃₄₁ , G ₃₃₉ -N ₃₄₄ , A ₄₃₃ -E ₄₃₈ , T ₄₆₉ -A ₄₇₃ , V ₄₇₄ -K ₄₇₉ , L ₄₉₀ -N ₄₉₅ , N ₄₉₂ - G ₄₉₇ , N ₅₀₉ -N ₅₁₅ , G ₅₁₀ -N ₅₁₄	
