

Supplementary Material

Table S1. Amino acids monoisotopic masses. MRMs. retention times (t_R) and conditions in the mass spectrometer along with their respective molecular formulas.

Analyte	Molecular Formula	Monoisotopic mass	Precursor Ion	Product Ion	Collision Energy (V)	Tube Lens	t_R standard solutions (min) (Mean \pm Sd*)	t_R hydrolysed sample (min) (Mean \pm Sd*)
Phe	C ₉ H ₁₁ NO ₂	165.19	166.1	120.2	14	47	2.50 \pm 0.06	3.12 \pm 0.13
Trp	C ₁₁ H ₁₂ N ₂ O ₂	204.23	205.1	188.1	12	52	2.58 \pm 0.06	-
Ile	C ₆ H ₁₃ NO ₂	131.17	132.1	86.3	6	40	2.67 \pm 0.06	3.18 \pm 0.15
Leu	C ₆ H ₁₃ NO ₂	131.17	132.1	86.3	11	49	3.19 \pm 0.07	3.38 \pm 0.17
Asn	C ₄ H ₈ N ₂ O ₃	132.12	133.1	87.2	8	49	2.74 \pm 0.05	-
Met	C ₅ H ₁₁ NO ₂ S	149.21	150.1	56.4	17	46	3.15 \pm 0.07	3.70 \pm 0.12
Val	C ₅ H ₁₁ NO ₂	117.15	118.1	72.3	11	46	3.22 \pm 0.03	3.99 \pm 0.14
Pro	C ₅ H ₉ NO ₂	115.13	116.1	70.3	17	51	3.56 \pm 0.07	4.02 \pm 0.16
Tyr	C ₉ H ₁₁ NO ₃	181.19	182.1	136.2	14	46	3.53 \pm 0.05	4.24 \pm 0.12
Ala	C ₃ H ₇ NO ₂	89.094	90.1	44.5	10	40	4.55 \pm 0.10	5.26 \pm 0.11
Thr	C ₄ H ₉ NO ₃	119.12	120.1	74.3	11	46	4.91 \pm 0.03	5.73 \pm 0.11
Gly	C ₂ H ₅ NO ₂	75.07	76.1	30.6	12	40	5.22 \pm 0.04	5.85 \pm 0.15
Ser	C ₃ H ₇ NO ₃	105.09	106.1	60.4	12	60	5.74 \pm 0.07	6.47 \pm 0.13
Glu	C ₅ H ₉ NO ₄	147.13	148.0	84.0	15	70	5.94 \pm 0.09	6.85 \pm 0.11
Asp	C ₄ H ₇ NO ₄	133.11	134.1	88.3	11	70	7.86 \pm 0.12	8.81 \pm 0.16
Arg	C ₆ H ₁₄ N ₄ O ₂	174.20	175.1	70.3	23	54	8.35 \pm 0.11	9.18 \pm 0.11
Gln	C ₅ H ₁₀ N ₂ O ₃	146.15	147.1	130.2	9	45	8.78 \pm 0.12	9.54 \pm 0.12
Lys	C ₆ H ₁₄ N ₂ O ₂	146.11	147.1	84.3	17	48	8.79 \pm 0.11	9.54 \pm 0.14
His	C ₆ H ₉ N ₃ O ₂	155.16	156.1	110.2	15	49	9.20 \pm 0.10	9.92 \pm 0.17
Cys ₂	C ₆ H ₁₂ N ₂ O ₄ S ₂	240.29	241.0	152.0	13	76	10.24 \pm 0.08	-
Cys	C ₃ H ₇ NO ₂ S	121.16	122.1	59.3	20	44	1.84 \pm 0.06	-

* For S_d calculation n=6.

Table S2. Precision and trueness data for the analysis of AA in spiked B-90 bread wheat sample.

Target Amino acid	Added (mg/100 g)	Repeatability (n=3)			Intermediate precision (n=3)		
		Found (mg/100 g)	Recovery (%)	RSD (%)	Found (mg/100 g)	Recovery (%)	RSD (%)
Gly	20.0	19.0	94.8	8.7	19.2	96.0	7.9
	40.0	43.9	109.8	2.9	44.5	111.2	6.6
Ala	20.0	19.9	99.6	6.2	20.2	100.8	4.8
	40.0	46.5	116.2	4.4	47.0	117.6	3.4
Ser	20.0	21.3	106.3	2.3	21.52	107.6	1.4
	40.0	40.5	101.3	6.5	41.04	102.6	3.5
Pro	20.0	20.1	100.7	7.2	20.4	101.9	3.2
	40.0	39.2	98.1	13.9	39.7	99.3	9.8
Val	20.0	20.1	100.3	2.9	20.3	101.5	3.6
	40.0	39.7	99.3	6.8	40.2	100.5	8.7
Thr	20.0	19.3	96.5	4.6	19.5	97.7	2.8
	40.0	41.9	104.7	6.6	42.4	106.0	6.7
Asp	20.0	20.6	103.2	0.9	20.9	104.5	2.7
	40.0	46.2	115.4	1.7	46.7	116.8	4.3
Glu	20.0	18.8	93.8	0.6	19.0	95.0	1.8

	40.0	35.6	88.9	4.8	36.0	90.0	7.2
Ile	20.0	19.4	97.0	2.3	19.6	98.2	5.2
	40.0	39.3	98.2	5.3	39.8	99.4	8.9
Leu	20.0	19.4	97.0	2.3	19.6	98.2	6.1
	40.0	48.7	121.8	2.8	49.3	123.3	5.4
Gln	20.0	18.9	94.4	6.5	19.1	95.6	5.6
	40.0	38.9	97.3	9.2	39.4	98.5	8.7
Lys	20.0	20.8	104.2	3.7	21.1	105.5	3.1
	40.0	34.3	85.7	5.6	34.7	86.8	13.7
Met	20.0	20.6	103.0	9.8	20.9	104.3	5.5
	40.0	48.1	120.2	5.7	48.7	121.7	7.6
His	20.0	18.8	93.9	1.8	19.0	95.1	2.3
	40.0	44.5	111.3	5.5	45.1	112.7	2.7
Phe	20.0	20.6	102.9	2.3	20.8	104.2	8.4
	40.0	40.1	100.2	11.1	40.6	101.4	11.3
Arg	20.0	19.1	95.3	7.3	19.3	96.5	5.6
	40.0	39.0	97.6	10.2	39.5	98.8	4.3
Tyr	20.0	20.4	102.0	4.0	20.7	103.3	3.8
	40.0	39.9	99.9	8.3	40.4	101.1	9.7