Table S6. Multiple Reaction Monitoring transitions for amino acid analysis.

Amino acid	Transition (*m*/*z*)	RT (min)	Fragmentor	CE (V)
Trp	205.1 -> 146	3.71	25	15
Arg	175.1 -> 70.1	8.27	75	28
Phe	166.1 -> 120.1	3.51	25	5
His	156.1 -> 110	7.44	25	12
Met	150.1 -> 56.1	4.38	22	9
Glu	148.1 -> 84.1	7.04	75	16
Lys	147.1 -> 84	8.71	50	16
Ile	132.1 -> 86.1	4.10	25	8
Leu	132.1 -> 86.1	3.81	25	8
Hyp	132.1 -> 86	5.65	79	27
Cr	132.1 -> 44	5.56	28	10
Thr	120.1 -> 56.2	5.71	25	10
Val	118.1 -> 72.1	4.97	25	8
Pro	116.1 -> 70.1	5.03	50	16
Ser	106 -> 60.1	6.21	25	8
Ala	90.1 -> 44.1	5.63	25	8
Sar	90.1 -> 44	5.48	44	19
Gly	76 -> 30.1	6.03	25	5