Peptide	Unique	-10lgP ↓	Mass	Length	ppm	m/z	RT	Area Sample 1
A(+36.08)DQVSASLAK(+36.08)QGL	•	44.59	1358.8344	13	-0.5	680.4241	30.94	2.14E7
A(+32.06)DQVSASLAK(+32.06)QGL	•	34.53	1350.7958	13	-0.9	676.4045	30.98	9.5E6
A(+28.03)DQVSASLAK(+28.03)QGL	•	31.48	1342.7456	13	0.2	672.3802	31.40	1.05E7

>XP_021329127.1 microtubule-associated protein tau isoform X1

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481  PARKASVPSK AKAGAGATPE KKAGDAKTKT AGAKPQGVGA KIPAASRMEQ RRTGPGSIER ADSPKTPDRS GCSSPASRSS
561  TPGQQVKKVA VVRTPPKSPG SLRSRAQIAP VAPMPDLKNV KSKIGSTENL KHQPGGGKIQ IVHKKIDLSN VQSKCGSKAN
641  IHHKPGGGNV EIKSEKLDFK AQSKVGSLEN IGHVPGGGQR RIESHKLNFR DQAKARTDHG ADIVCKSPDI STDGSPRRLS
721  NVSSSGSLNM TDSPQLSTLA DQVSASLAKQ GL
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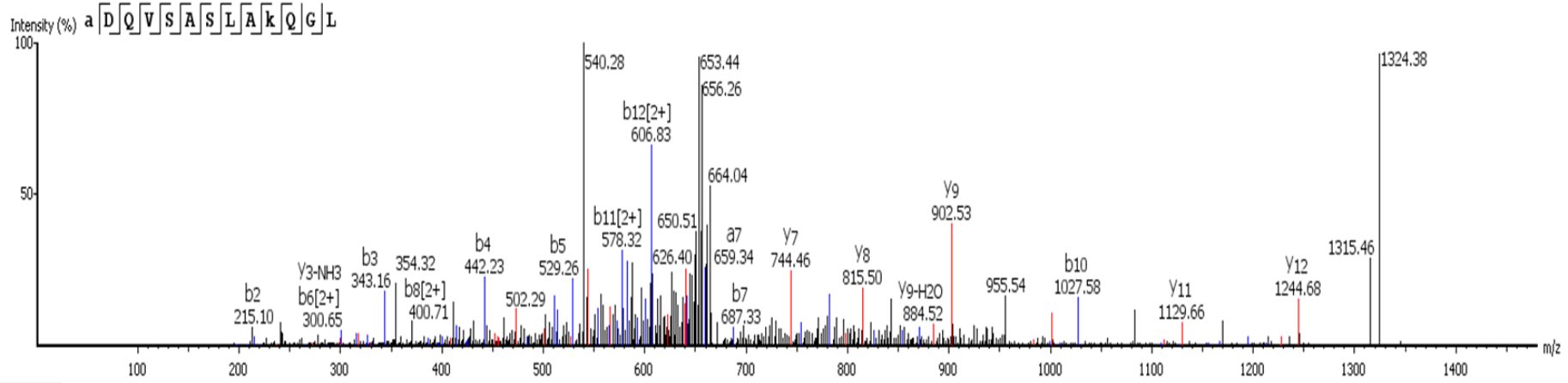

Ion Match

#	b	b-H2O	b-NH3	a	a-H2O	a-NH3	b (2+)	a (2+)	Seq	y	y-H2O	y-NH3	y (2+)	#
1	100.08	82.07	83.05	72.08	54.07	55.05	50.54	36.54	A(+28.03)					13
2	215.16	197.25	198.20	187.11	169.10	170.08	108.05	94.05	D	1244.68	1226.58	1227.79	622.69	12
3	343.27	325.39	326.34	315.26	297.16	298.14	172.08	158.08	Q	1129.61	1111.69	1112.51	565.44	11
4	442.29	424.28	425.36	414.32	396.32	397.36	221.19	208.12	V	1001.56	983.61	984.57	501.35	10
5	529.26	511.27	512.24	501.35	483.55	484.38	265.37	251.13	S	902.53	884.57	885.50	451.70	9
6	600.31	582.29	583.27	572.33	554.14	555.28	300.28	286.31	A	815.49	797.79	798.45	408.41	8
7	687.34	669.32	670.30	659.60	641.33	641.96	344.17	330.33	S	744.50	726.45	727.43	372.73	7
8	800.42	782.52	783.39	772.42	754.63	755.39	400.48	386.38	L	657.43	639.26	640.11	329.37	6
9	871.21	853.68	854.43	843.46	825.45	826.38	436.23	422.23	A	544.18	526.33	527.01	272.19	5
10	1027.51	1009.57	1010.55	999.69	981.57	982.56	514.37	500.29	K(+28.03)	473.32	455.04	456.17	237.24	4
11	1155.53	1137.63	1138.61	1127.81	1109.43	1110.35	578.20	564.34	Q	317.24	299.37	300.28	159.09	3
12	1212.68	1194.57	1195.63	1184.66	1166.57	1167.43	606.63	592.53	G	189.12	171.11	172.10	95.06	2
13									L	132.10	114.09	115.07	66.55	1