

peptide	species_1
IVIPVQILVHNESIDQLGYR	BRAHW
IVIPVQILVHNESIDQLGYR	BRAHW
LDNTYGNMIGISLDPQIR	BRAHW
NYTQEVLR	BRAHW
MYGDQDDWIDFLTDGNQFR	BRAHW
SYDEEFFAR	BRAHW
VNHYSWCAYAELYITPVK	BRAHW
VNHYSWCAYAELYITPVK	BRAHW
ALDSMFEPIEGYSSDRPVSSQANINAK	BRAHW
MDQGFVVLGNSTIK	BRAHW
NYEEFFSR	BRAHW
FYFLNTTVGNVTINPFIK	BRAHW
FYFLNTTVGNVTINPFIK	BRAHW
NYEEELFSR	BRAHW
VAYNTALHGVGTMIR	BRAHW
ANTLINGSILNTGNK	BRAHW
MNQLGFTLGNGTIK	BRAHW
YYTGIDAFNYIR	BRAHW, BRAPL
AGEAVQNTVSGYHPSNPNYK	BRAHW
TADGANEAFQSLGFEAR	BRAHW
VAYNTALQGYNR	BRAHW
DYLGISTDIQIR	BRAHW
IAIPIQIAASK	BRAHW
TIGNVNINPFIK	BRAHW
FMVGVSGETAGVLLDNLDR	BRAHW
FMVGVSGETAGVLLDNLDR	BRAHW
QGGLAFNGSTGITWK	BRAHW
QGGLAFNGSTGITWK	BRAHW
TANGASEYFAQSLGFEAR	BRAHW
GGINTFRPAAIAGFGYK	BRAHW
GGINTFRPAAIAGFGYK	BRAHW
GYFIAATDPVLVEPQIR	BRAHW
GYFIAATDPVLVEPQIR	BRAHW
LDNTYDNYMIGISLDPQIR	BRAHW
LDNTYDNYMIGISLDPQIR	BRAHW
YYTGIDLLNQVR	BRAHW
LDVHTPVLAVALANNLR	BRAHW
FMVGVLGETAGVLLDNLDR	BRAHW
FMVGVLGETAGVLLDNLDR	BRAHW

accession	rank	name	recursor charge
COR1E3, F1CJQ1, O30571	2	vspA	nd
COR1E3, F1CJQ1, O30571	2	vspA	nd
C0QZU2	6	vspA	3
COR1E3	4	vspA	2
COR1E1	12	vspB	2
COR1E1, O30572	5	vspB	nd
O30572	2	vspB	nd
O30572	2	vspB	nd
COR1E0, O68156	7	vspC	3
COR1E0, O68156	11	vspC	2
COR1E0, F1CJQ2, O68156	8	vspC	2
COR1D9, F1CJQ3, O68157	1	vspD	nd
COR1D9, F1CJQ3, O68157	1	vspD	nd
COR1D9, F1CJQ3, O68157	6	vspD	2
COR1D9, F1CJQ3, O68157	4	vspD	2,3
COR1T2, F1CJQ4, Q9AEX8	3	vspE	2
COR1T2, F1CJQ4	4	vspE	2
COR1T2, D8IFY2, F1CJQ4, J9UHW9, KOJK05, Q9AEX8	4	vspE	2
Q9AEX7	6	vspF	2,3
COR1T3, F1CJQ5	6	vspF	2,3
COR1T3, F1CJQ5	3	vspF	2
Q9AEX6	3	vspG	nd
Q9AEX6	2	vspG	nd
Q9AEX6	1	vspG	nd
C0QX33	6	vspH	nd
C0QX33	6	vspH	nd
C0QX33	2	vspH	nd
C0QX33	2	vspH	nd
Q9AEX5	11	vspH	2,3
C0QX33, F1CJQ7	1	vspH, vspJ	nd
C0QX33, F1CJQ7	1	vspH, vspJ	nd
C0QX33, F1CJQ7	2	vspH, vspJ	nd
C0QX33, F1CJQ7	2	vspH, vspJ	nd
F1CJQ6	4	vspI	nd
F1CJQ6	4	vspI	nd
C0QZU2, F1CJQ6	4	vspA, vspI	2
C0QZU2, F1CJQ6	1	vspA, vspI	3
F1CJQ7	10	vspJ	nd
F1CJQ7	10	vspJ	nd

elected charg	mass	m/z mono	final mass	WA1		WA1 sept
				psms	av psms	psms
2	2468,3572	1235,1786	2470,3572			
3	2468,3646	823,7882	2471,3646			
3	2069,0106	690,6702	2072,0106	ND*		
2	1092,571	547,2855	1094,571	ND*		
2	2334,9998	1168,4999	2336,9998	32	11	13
2	1162,5076	582,2538	1164,5076		0	
2	2326,1602	1164,0801	2328,1602		0	
3	2326,1676	776,3892	2329,1676		0	
3	2925,4032	976,1344	2928,4032	1	0	8
2	1555,785	778,8925	1557,785	61*	20	18*
2	1219,5292	610,7646	1221,5292	3	1	
2	2087,1238	1044,5619	2089,1238	4	1	7
3	2087,1309	696,7103	2090,1309		0	
2	1185,5448	593,7724	1187,5448	ND	nd	
2	1601,8494	801,9247	1603,8494	18	6	6
2	1528,8354	765,4177	1530,8354	9	3	3
2	1492,7854	747,3927	1494,7854	50	17	8
2	1494,7288	748,3644	1496,7288	ND	nd	
2	2132,0068	1067,0034	2134,0068	ND*	nd	
2	1853,869	927,9345	1855,869	2550*	850	764*
2	1368,6932	685,3466	1370,6932	62	21	15
2	1392,7394	697,3697	1394,7394		0	
2	1123,711	562,8555	1125,711		0	
2	1328,7598	665,3799	1330,7598		0	
2	1992,0132	997,0066	1994,0132		0	
3	1992,0204	665,0068	1995,0204		0	
2	1535,7878	768,8939	1537,7878		0	
3	1535,7951	512,9317	1538,7951		0	
2	1917,9002	959,9501	1919,9002	ND	nd	
2	1738,93	870,465	1740,93		0	
3	1738,9374	580,6458	1741,9374		0	
2	1888,024	945,012	1890,024		0	
3	1888,0314	630,3438	1891,0314		0	
2	2127,0088	1064,5044	2129,0088		0	
3	2127,0162	710,0054	2130,0162		0	
2	1453,771	727,8855	1455,771	13	4	2
3	1929,1014	644,0338	1932,1014	19*	6	5*
2	2018,0652	1010,0326	2020,0652		0	
3	2018,0724	673,6908	2021,0724		0	

B78 psms	INF1 psms	WA1 detected	B78 detected	INF1 detected
25	7	nd	nd	nd
14		nd	nd	nd
59*		y	y	nd
1		y	nd	nd
14	6			
44	20	y	y	y
		y	y	nd
1		nd	nd	nd
285		nd	nd	nd
		nd	nd	nd
8	16*			
3	13	no conclusive	no conclusive	y
		nd	nd	nd