



**Fig. SI-1** PAGE analysis of the purified Selp preparation

# FRAGMENTATION of *m/z* 3945 peptide

(a) mono charged fragments

#1	Seq.		#2		
1	E		30		
2	S	1267.16906	29	<b>b<sup>+</sup></b>	<b>y<sup>+</sup></b>
3	C		28	<b>b<sup>2+</sup></b>	<b>y<sup>2+</sup></b>
4	Q		27	<b>b<sup>3+</sup></b>	<b>y<sup>3+</sup></b>
5	C		26	<b>b<sup>4+</sup></b>	<b>y<sup>4+</sup></b>
6	R		25	<b>b<sup>5+</sup></b>	<b>y<sup>5+</sup></b>
7	L		24		
8	P	777.80907	23	<b>b-H<sub>2</sub>O<sup>+</sup></b>	<b>y-H<sub>2</sub>O<sup>+</sup></b>
9	P		22	<b>b-H<sub>2</sub>O<sup>2+</sup></b>	<b>y-H<sub>2</sub>O<sup>2+</sup></b>
10	A	729.52869	21	<b>b-H<sub>2</sub>O<sup>3+</sup></b>	<b>y-H<sub>2</sub>O<sup>3+</sup></b>
11	A		20	<b>b-H<sub>2</sub>O<sup>4+</sup></b>	<b>y-H<sub>2</sub>O<sup>4+</sup></b>
12	C- KB_S_sugar		19	<b>b-H<sub>2</sub>O<sup>5+</sup></b>	<b>y-H<sub>2</sub>O<sup>5+</sup></b>
13	Q	769.98529	18	<b>b-NH<sub>3</sub><sup>+</sup></b>	<b>y-NH<sub>3</sub><sup>+</sup></b>
14	I	732.97461	17	<b>b-NH<sub>3</sub><sup>2+</sup></b>	<b>y-NH<sub>3</sub><sup>2+</sup></b>
15	S	689.60441	16	<b>b-NH<sub>3</sub><sup>3+</sup></b>	<b>y-NH<sub>3</sub><sup>3+</sup></b>
16	Q	990.38696	15	<b>b-NH<sub>3</sub><sup>4+</sup></b>	<b>y-NH<sub>3</sub><sup>4+</sup></b>
17	Q	934.87094	14	<b>b-NH<sub>3</sub><sup>5+</sup></b>	<b>y-NH<sub>3</sub><sup>5+</sup></b>
18	L	431.66783	13		
19	I	814.29962	12		
20	P	757.75759	11		
21	T		10		
22	E		9		
23	A		8		
24	S		7		
25	A		6		
26	S		5		
27	C-KBSecAu		4		
28	R		3		
29	C		2		
30	K		1		

(b) doubly charged fragments

#1		Seq.		#2
1		E		30
2		S	760.70434	29
3		C		28
4		Q		27
5		C		26
6	689.24939	R		25
7	820.34402	L		24
8		P	777.80907	23
9	996.43898	P		22
10		A	729.52869	21
11		A		20
12		C- KB_S_sugar	694.01014	19
13		Q	769.98529	18
14		I	732.97461	17
15	966.38628	S	689.60441	16
16	1030.90758	Q	990.38696	15
17	547.97207	Q	934.87094	14
18	773.66387	L	580.89686	13
19		I	814.29962	12
20	628.77729	P	757.75759	11
21		T		10
22	686.29986	E		9
23	704.05914	A		8
24	580.85517	S		7
25	990.77147	A		6
26	765.33443	S	471.11967	5
27	881.33930	C-KBSecAu		4
28	920.61058	R		3
29	757.29176	C		2
30		K		1

(c) triply charged fragments

#1		Seq.		#2
1		E		30
2		S	1267.16906	29
3		C		28
4	431.12311	Q	1209.49750	27
5		C		26
6	689.24939	R		25
7		L		24
8		P	777.80907	23
9	996.43898	P	758.04852	22
10		A	729.52869	21
11		A		20
12		C- KB_S_sugar	698.26677	19
13		Q	769.98529	18
14		I	732.97461	17
15	966.87829	S	689.60441	16
16	412.96740	Q	990.38696	15
17	547.97207	Q	934.87094	14
18	767.98836	L	431.66783	13
19		I		12
20	628.77729	P	757.75759	11
21		T		10
22		E	658.70737	9
23	704.05914	A		8
24	580.85517	S		7
25	991.09948	A		6
26	765.33443	S		5
27		C		4
28	740.09523	R		3
29	757.29176	C-KBSecAu		2
30		K		1

(d) +4 charged fragments

#1		Seq.		#2
1		E		30
2		S	760.70434	29
3		C		28
4		Q	1209.49750	27
5		C		26
6	689.24939	R		25
7		L		24
8		P	777.80907	23
9	996.43898	P	758.04852	22
10		A	729.52869	21
11		A		20
12		C- KB_S_sugar	694.01014	19
13		Q	769.98529	18
14		I	732.97461	17
15	966.38628	S	689.60441	16
16	1030.90758	Q	990.38696	15
17	547.97207	Q	934.87094	14
18	773.66387	L	580.89686	13
19		I		12
20	628.77729	P	757.75759	11
21		T		10
22	686.29986	E	658.70737	9
23	704.05914	A		8
24	580.85517	S		7
25	990.77147	A		6
26	765.33443	S	471.11967	5
27		C		4
28		R		3
29	757.29176	C-KBSecAu		2
30		K		1

**FRAGMENTATION of *m/z* 3403 peptide**

#1		Seq.		#2		
1		G-Acetyl		29		
2		S	1096.52536	28		<b>b<sup>+</sup></b>
3		V		27		<b>b<sup>2+</sup></b>
4	369.17686	T	1034.49188	26		<b>b<sup>3+</sup></b>
5		V	1000.80932	25		<b>b<sup>4+</sup></b>
6		V		24		<b>b<sup>5+</sup></b>
7		A		23		
8		L		22		<b>b-H<sub>2</sub>O<sup>+</sup></b>
9		L	659.55094	21		<b>y-H<sub>2</sub>O<sup>+</sup></b>
10		Q	835.36729	20		<b>b-H<sub>2</sub>O<sup>2+</sup></b>
11	1064.59864	A		19		<b>y-H<sub>2</sub>O<sup>2+</sup></b>
12		S		18		<b>b-H<sub>2</sub>O<sup>3+</sup></b>
13	651.79581	C- S_Se_subst	745.99491	17		<b>y-H<sub>2</sub>O<sup>3+</sup></b>
14		Y	1034.49860	16		<b>b-H<sub>2</sub>O<sup>4+</sup></b>
15	789.86951	L	952.96694	15		<b>y-H<sub>2</sub>O<sup>4+</sup></b>
16	937.35664	C-KB_Pt		14		<b>b-H<sub>2</sub>O<sup>5+</sup></b>
17		I	747.95376	13		<b>y-H<sub>2</sub>O<sup>5+</sup></b>
18	1059.44599	L	700.41701	12		<b>b-NH<sub>3</sub><sup>+</sup></b>
19	743.64376	Q	429.58574	11		<b>y-NH<sub>3</sub><sup>+</sup></b>
20		A		10		<b>b-NH<sub>3</sub><sup>2+</sup></b>
21	796.33348	S		9		<b>y-NH<sub>3</sub><sup>2+</sup></b>
22	839.03180	K	500.81112	8		<b>b-NH<sub>3</sub><sup>3+</sup></b>
23	657.79668	L		7		<b>y-NH<sub>3</sub><sup>3+</sup></b>
24		E	759.43593	6		<b>b-NH<sub>3</sub><sup>4+</sup></b>
25		D		5		<b>y-NH<sub>3</sub><sup>4+</sup></b>
26	747.08508	L		4		<b>b-NH<sub>3</sub><sup>5+</sup></b>
27	786.11036	R		3		<b>y-NH<sub>3</sub><sup>5+</sup></b>
28		V		2		
29		K		1		