

**Predicted epitopes:**

No.	Start Position	End Position	Peptide	Peptide Length
1	1	8	MTTRSASP	8
2	25	26	AN	2
3	28	28	V	1
4	44	48	RAAQA	5
5	59	67	EASAGNAAA	9
6	79	88	CLTTSGPGFL	10
7	96	97	NA	2
8	108	115	SGSSSRPM	8
9	120	126	RGDYQDL	7
10	128	129	QL	2
11	147	150	DIGR	4
12	158	170	TATSGRPPGGVYLD	13
13	174	177	DVLG	4
14	180	207	VEASAASGAIWRPVDPAPRLLPAPEAID	28
15	228	231	AYAQ	4
16	257	267	LPDSHPQSAAA	11
17	292	300	NGESPQWSA	9
18	311	321	ASEFDSNRPIV	11
19	323	323	P	1
20	325	327	TGD	3
21	329	329	G	1
22	340	362	ADRSSVASAAWTGELADRKARNS	23
23	371	371	D	1
24	373	373	H	1
25	390	390	Q	1
26	395	398	VYVV	4
27	402	403	AN	2
28	421	424	LDSG	4
29	426	426	W	1
30	441	451	VETGRPVAIE	11
31	453	453	D	1
32	455	456	AF	2
33	482	487	GVYRGD	6
34	497	507	PVWRHDPAPTV	11
35	521	537	FGGKGYHVSTPTELESA	17
36	542	549	LASNGPSL	8
37	554	562	LDPADGVES	9