

Predicted epitopes:

No	Start Position	End Position	Peptide	
1	1	70	MSRAAPRRASQSQSTRPARGLRPPGAQEVGQRKRPGKTQKARQAQEATKSRPATRSDVAPAGRST RARR	70
2	77	80	VGTR	4
3	119	120	AQ	2
4	122	122	A	1
5	125	125	L	1
6	127	141	VTDVQPAARGSIVDR	15
7	167	183	EEARKKTSAAPDPQQRL	17
8	189	189	E	1
9	191	201	AGKLNNKPDAA	11
10	221	223	DPA	3
11	233	251	PEVGAERQDLRQYPGGSLA	19
12	257	261	GIDWD	5
13	277	294	AGTDGSVTYDRGSDGVVI	18
14	297	301	SYRNR	5
15	304	305	AV	2
16	326	335	VQQAKNLGA	10
17	356	387	DNTFDPSQDIGRQGDQLGNPAVSSPFEPGSV	32
18	401	409	LSSPDEVLQ	9
19	413	416	SIQM	4
20	430	433	MPYT	4
21	435	443	TGVFGKSSN	9
22	453	457	VGPER	5
23	470	492	RTGVGLPGESAGLVPPIDQWSGS	23
24	494	494	F	1
25	519	524	NDGVRV	6
26	533	549	VAPDGSRTEEPRPDDIR	17
27	554	557	QTAQ	4
28	569	602	RDPMGYQQGTGPTAGVPGYQMAGKTGTAQQINPG	34
29	619	625	ATADNPR	7
30	635	647	PARNSDGAPGHS	13
31	663	675	NVPLSPDPGPPLV	13