

Predicted epitopes:

No.	Start Position	End Position	Peptide	Peptide Length
1	1	17	MADPKTKGRGSGGNGSG	17
2	26	30	PTKAR	5
3	41	42	IV	2
4	47	47	G	1
5	53	68	PRAASDVPAKYKSQPW	16
6	76	78	DAD	3
7	87	93	PEKRSTV	7
8	110	117	TDGDREGE	8
9	143	155	EPAIRAAAEHPRD	13
10	163	163	A	1
11	188	193	PKLSAG	6
12	195	195	V	1
13	227	240	ASVSDPDAAPPTFS	14
14	250	258	RVATGRDFD	9
15	263	265	LRK	3
16	275	279	GSATA	5
17	284	286	LDG	3
18	289	289	L	1
19	291	308	VASAEKPYARRPYPPFM	18
20	312	315	LQQE	4
21	317	317	S	1
22	345	345	T	1
23	347	352	STTLSE	6
24	355	360	INAART	6
25	364	366	QLY	3
26	369	414	EYVAPAPRQYTRKVKNAQEAHEAIRPAGETFATPDAVRRELDGPNI	46
27	430	435	SQMADA	6
28	448	449	SG	2
29	477	489	ELVGGEADDAERR	13
30	491	496	PHLTPG	6
31	504	518	LTPDGHATNPPARYT	15
32	531	537	IGRPSTY	7
33	549	552	YVHK	4
34	556	556	A	1
35	585	585	A	1
36	587	601	EDELDEIAAGNERRT	15
37	612	622	DHGVPDSVARS	11

38	636	636	I	1
39	638	639	AR	2
40	649	653	DTHGR	5
41	660	665	GKNGPY	6
42	671	692	AGDTGEPTPQRANLSDSITPDE	22
43	702	721	FATPQQGRTLGLDPETGHEI	20
44	726	730	GRFGP	5
45	735	764	ILPEPAADAAAAAQGVKKRQKAAGPKPRTG	30
46	791	821	GVDPASGEEITAQNGRYGPYLKRGNDSRSLV	31
47	840	868	EPKRRGRQSASAPPLRELGTDPASGKPMV	29
48	873	900	RFGPYVTDGETNASLRKGDDVASITDER	28