

Predicted epitopes:

No.	Start Position	End Position	Peptide	Peptide Length
1	10	10	G	1
2	12	12	T	1
3	20	37	EGGDPVVVANSEGSRTTP	18
4	49	61	LVGQPAKNQAVTN	13
5	64	64	R	1
6	72	72	H	1
7	74	74	G	1
8	78	80	SIE	3
9	82	90	DGKKYTAPE	9
10	104	110	EAYLGED	7
11	114	114	A	1
12	116	116	I	1
13	120	136	AYFNDAQRQATKDAGQI	17
14	147	151	EPTAA	5
15	158	162	DKGEK	5
16	174	174	G	1
17	191	206	RATSGDNHLGGDDWDQ	16
18	217	219	KGT	3
19	221	224	GIDL	4
20	236	240	EAAEK	5
21	246	252	SSSQSTS	7
22	260	264	VDADK	5
23	290	292	RKP	3
24	302	303	IS	2
25	317	320	TRMP	4
26	330	345	TGGKEPNKGVNPDEVV	16
27	376	376	E	1
28	378	378	K	1
29	380	380	G	1
30	390	410	TTIPTKRSETFTTADDNQPSV	21
31	417	417	G	1
32	419	422	REIA	4
33	434	445	TGIPPAPRGIPQ	12
34	462	476	AKDKGTGKENTIRIQ	15
35	478	484	GSGLSKE	7
36	494	512	EAHAEEDRKRREEADVRNQ	19
37	523	543	FVKEQREAEGGSKVPEDTLNK	21
38	546	550	AAVAE	5

39	553	558	AALGGS	6
40	560	561	IS	2
41	569	577	KLGQESQAL	9