

Predicted epitopes:

No.	Start Position	End Position	Peptide	Peptide Length
1	1	14	MTTSESPDAYTESF	14
2	19	44	IVKPAGPPRVGQPSWNPQRASSMPVN	26
3	46	53	YRPF AEEV	8
4	61	68	RTWPDRVI	8
5	80	82	RDG	3
6	84	94	QALIDPMSPAR	11
7	112	121	VGFP SASQTD	10
8	132	137	GAIPDD	6
9	158	161	CSGA	4
10	190	199	AIATDGARKC	10
11	201	226	EQA AKYPGTQWRFEYSPESYTGTELE	26
12	238	246	VIAP TPERP	9
13	255	264	VEM TTPNVYA	10
14	275	275	A	1
15	287	294	HND RGTAV	8
16	305	307	ADR	3
17	316	319	GERT	4
18	336	341	VDPQID	6
19	361	368	HERHPYGG	8
20	376	384	SGSHQDAIN	9
21	394	400	ADA ADCD	7
22	411	421	LPIDPRDVGRT	11
23	430	436	SQSGKGG	7
24	465	480	AEGTAGEGGEVSPKEM	16
25	490	490	A	1
26	502	515	VDAADDDGGTTSIT	14
27	524	534	ETEISGSGNGP	11
28	559	568	MSAGDDAQAA	10
29	570	570	Y	1
30	576	616	TIASPAQPGEAGRHASDPVTIASPAQPGEAGRHASDPVTSK	41
31	618	618	V	1
32	624	628	APSIT	5