

Predicted epitopes:

No.	Start Position	End Position	Peptide	Peptide Length
1	17	29	VADISPQDAAHRR	13
2	34	36	VEF	3
3	38	42	SRAIG	5
4	53	66	ARADGHDHAASAVA	14
5	78	79	VG	2
6	85	91	VPPVAAP	7
7	99	107	EHDNDGSGA	9
8	136	144	GSSGKTSTK	9
9	156	167	EVVAPPGSFNNE	12
10	169	169	G	1
11	202	206	IAPPS	5
12	227	227	I	1
13	231	241	KAELPQAVPHS	11
14	247	254	NADDPAVA	8
15	269	282	RDNTGDVWAGPVSL	14
16	297	298	AQ	2
17	309	310	HQ	2
18	327	327	S	1
19	336	343	TAAPPVSR	8
20	349	357	TTRGDGVTV	9
21	360	370	DAYNANPDSMR	11
22	382	388	QPEATRR	7
23	398	399	EL	2
24	401	406	EDAIAE	6
25	443	466	AWGSGEATADHGADRTAVNVADGD	24
26	477	477	R	1
27	489	492	AAGL	4