

Predicted epitopes:

No.	Start Position	End Position	Peptide	Peptide Length
1	2	15	VQPADIDVPETPAR	14
2	45	45	P	1
3	48	48	A	1
4	55	55	A	1
5	66	78	GPASVYADGAPKL	13
6	110	117	TGTREYGR	8
7	128	134	HSDLPEV	7
8	139	139	M	1
9	143	164	DAVTAAPDGFDDVASSAGAPVA	22
10	181	189	EVMHTPHGQ	9
11	204	210	AQWTPAN	7
12	221	224	TQIG	4
13	234	239	GGVDSA	6
14	268	271	RAQV	4
15	298	307	GVSAPGKRR	10
16	325	327	LDG	3
17	340	355	PDVVESGGGSGTANIK	16
18	359	365	NVGGLPD	7
19	385	385	A	1
20	391	391	G	1
21	397	407	VARQPFPGPGL	11
22	436	437	AA	2
23	439	439	L	1
24	457	467	VGVQGDGRTYG	11
25	475	488	VSSDAMTADWTRV	14
26	505	505	A	1
27	515	524	TSKPPATIEW	10