

**Predicted epitopes:**

<b>No.</b>	<b>Start Position</b>	<b>End Position</b>	<b>Peptide</b>	<b>Peptide Length</b>
1	12	21	LYGPTTGDR I	10
2	34	49	EDRCGGPGLAGDEAVF	16
3	54	54	V	1
4	59	75	MGQGRASRADGAPDTV I	17
5	102	110	GKAGNPDI M	9
6	119	120	VG	2
7	123	126	TEII	4
8	156	156	A	1
9	160	181	TIIGGGTGPAEGTKATTVTPGE	22
10	193	193	G	1
11	204	211	GNTVNPDA	8
12	217	221	RGGAS	5
13	228	235	DWGSTPAA	8
14	256	259	TLNE	4
15	261	263	GFV	3
16	279	290	HTEGAGGGHAPD	12
17	297	315	QPNVLPSSSTNPTRPHTVNT	19
18	332	334	RIP	3
19	344	350	IRPSTIA	7
20	366	372	SDSQAMG	7
21	393	409	GALEGDPSGSQAADNNR	17
22	468	483	MGDANASIP TPQPVLP	16
23	487	489	FGA	3
24	491	496	AATAAA	6
25	506	508	SID	3
26	522	522	A	1
27	528	532	RAVGK	5
28	534	538	DLPLN	5
29	541	551	LPSIEVDPDTF	11
30	558	569	QVWQPQPA AELP	12