

**Predicted epitopes:**

<b>No.</b>	<b>Start Position</b>	<b>End Position</b>	<b>Peptide</b>	<b>Peptide Length</b>
1	2	9	LTGNKPAV	8
2	27	68	SSNPLANFAPGYPTIEPAQPAVSPPTSQDPAGAVRPLSGHP	42
3	85	92	PGADSAAP	8
4	114	123	AAALTSDDHG	10
5	139	144	SSGHTA	6
6	150	154	DAAHT	5
7	161	162	RR	2
8	173	174	DG	2
9	181	195	KNPAVDPASGAATVA	15
10	209	210	TQ	2
11	212	212	N	1
12	220	249	GQTSVTTIGADGHAQQALRAGQGATTMAAD	30
13	279	288	PVRQAPYGLA	10
14	298	302	QTASN	5
15	312	316	GIPVE	5
16	318	336	VRYPTVQQPNSLAFDETS	19
17	343	348	GSGAGV	6