

**Predicted epitopes:**

<b>No.</b>	<b>Start Position</b>	<b>End Position</b>	<b>Peptide</b>	<b>Peptide Length</b>
1	1	35	MKDTVSQPAGGRGATAPRPADAASPCGSSPSADS	35
2	43	51	GGTAGHVEP	9
3	53	53	M	1
4	82	82	V	1
5	96	107	PMPRKPGGDLAR	12
6	121	121	D	1
7	123	125	LDD	3
8	149	156	LPLPPRRR	8
9	168	171	ARAG	4
10	177	180	GAHT	4
11	186	194	SAVPDSGLR	9
12	224	227	GFPD	4
13	239	242	QGAV	4
14	249	254	SGAAAD	6
15	267	269	GPQ	3
16	277	285	RAQGDPPYV	9
17	328	335	PIGNGEQR	8
18	343	344	NA	2
19	348	348	M	1
20	354	355	AL	2
21	384	393	VGHRDAAGQV	10