

Predicted epitopes:

No.	Start Position	End Position	Peptide	Peptide Length
1	9	10	PA	2
2	18	22	QEHVT	5
3	43	50	PRGCGKTS	8
4	61	71	CAQGPTANPCG	11
5	79	87	LAPNAPGSI	9
6	94	106	AASHGGVDDTREL	13
7	143	146	EEPP	4
8	156	162	TEPEKVL	7
9	195	196	VV	2
10	208	216	AGGGSPRDT	9
11	229	232	DTHV	4
12	272	281	IDGGHDPRRF	10
13	299	316	PDAASRGVVDAPEDALDR	18
14	319	319	E	1
15	322	323	AR	2
16	341	347	GEMRGAT	7
17	365	369	ASDAE	5
18	386	427	SIPAPQAVPRPSAAAAEPKHQPAREPRPVLAPTPASSEPTVA	42
19	435	435	T	1
20	471	472	AP	2
21	478	483	SEQRNA	6
22	503	542	ETGEPAAAASPVGGGANVATAKAVNPAPTANSTQRDEEEH	40
23	547	560	AGRGDPSPRRDPEE	14
24	575	575	I	1