

Predicted epitopes:

No.	Start Position	End Position	Peptide	Peptide Length
1	1	3	MTA	3
2	12	21	CPSPTGTPHV	10
3	36	37	HT	2
4	47	56	DTDAQRDSEE	10
5	72	90	WDEGPEVGGPYGPYRQSQR	19
6	108	119	YHAFSTPEEVEA	12
7	122	133	VAAGRNPCLGYD	12
8	140	140	T	1
9	149	153	AEGRQ	5
10	160	160	M	1
11	162	166	DDDLA	5
12	172	172	R	1
13	176	182	TFAAGSV	7
14	184	184	D	1
15	188	194	TRASGDP	7
16	218	225	LLPSTPRQ	8
17	251	266	LGEGTKKLSKRDPQSN	16
18	274	275	GF	2
19	292	292	A	1
20	311	327	DVNSSPARFDQKKADAL	17
21	391	409	YVIDPKAAAKELGPDGAAV	19
22	421	423	TDW	3
23	446	448	KAF	3
24	453	463	VAATGTTVSPP	11