

Predicted epitopes:

No.	Start Position	End Position	Peptide	Peptide Length
1	1	3	MSD	3
2	5	7	STF	3
3	48	67	TDTGAAPGLDSLPPAASEAE	20
4	90	96	YYDTHTP	7
5	108	123	AWYTAYTPYQPEISQG	16
6	148	156	MLDEGTAAA	9
7	167	169	RGP	3
8	180	180	F	1
9	205	210	AGLPDG	6
10	220	227	PGASGRIT	8
11	237	238	HD	2
12	257	264	PPGEIGAD	8
13	266	268	AFG	3
14	272	272	R	1
15	274	274	G	1
16	278	284	GFGGPHA	7
17	295	296	RQ	2
18	306	313	VDSDGTPA	8
19	324	332	HIRRDKATS	9
20	352	355	HGAG	4
21	393	398	PGRADE	6
22	423	428	DEATTD	6
23	442	452	AAAPAHTDIAT	11
24	463	468	FTQYRT	6
25	506	506	M	1
26	509	509	I	1
27	511	528	WPEFGRQHPFAPASDTAG	18
28	548	561	AVSLQPNAGSQGEY	14
29	572	578	ASRGEPH	7
30	587	595	SAHGTNAAS	9
31	609	616	HDNGDVDL	8
32	621	621	A	1
33	624	626	GEH	3
34	639	644	STHGVY	6
35	656	664	HDAGGQVYV	9
36	678	683	PGKFGG	6
37	697	708	PHGGGGPGVGPV	12
38	719	741	PGHPFAPELPGYYPVSSAPYGSA	23

39	759	759	L	1
40	786	786	T	1
41	789	791	NGM	3
42	823	825	FHA	3
43	828	831	MSFP	4
44	838	847	VEPTESESLA	10
45	864	883	DKVGAGEWPVDDNPLRGAPH	20
46	893	905	WDHPYTREQAAYP	13
47	907	907	G	1
48	909	911	AFR	3
49	915	915	W	1
50	922	926	DGAYG	5
51	937	939	VEA	3