

Predicted epitopes:

No.	Start Position	End Position	Peptide
1	9	44	GTSDTDGSQTRGAEPVEGQRTGPEEASNPGSARPRT
2	81	89	VRSYDDDGS
3	142	149	IGKEANGS
4	160	161	MQ
5	163	163	S
6	210	215	PDLGQT
7	255	259	GYRSD
8	265	279	LNPENDPQDSGYQAR
9	290	298	IFGDGLGQG
10	300	300	A
11	307	307	N
12	345	348	SRSA
13	390	396	AGGTSTA
14	409	416	ARHEPEAV
15	421	426	AGRDDK
16	436	442	PEPYLPP
17	449	512	DRKRANPQPAQTQPARKTPRTAPGQPARQMGLPPRPGSPRTADP PVRRSVHHGAGQRYAGQRRT