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NTDB id 1347 HI 0435 AAC22094.1		0
NTDB id 1373 NTHI RS02645 WP 011272046.1		0
NTDB id 38678 A464 RS17195 WP 042917093.1		0
NTDB id 1433 BW25113 RS17570 WP 000815987.1		0
NTDB id 1193 PA5040 NP 253727.1	MNSGLSRLGIALLAAMFAPALLAADLEKLDVAALPGDRVELKLQFDEPVAAPRGYTIEQPARIALDLPGVQNKLGTKNRE	80
NTDB id 1186 GTF74 RS01320 WP 000788426.1		0
NTDB id 1175 A1552VC RS12105 WP 000788426.1		0
NTDB id 1408 DSB67 RS13980 WP 010643500.1		0
consensus		0

Accession	Protein	Length	Score	Identity	Positives	Negatives	Gap	Consensus
NTDB id 1465	ASU2 RS06170 WP 014991934.1	160	100	100	160	0	0	0
NTDB id 1347	HI 0435 AAC22094.1	160	100	100	160	0	0	0
NTDB id 1373	NTHI RS02645 WP 011272046.1	160	100	100	160	0	0	0
NTDB id 38678	A464 RS17195 WP 042917093.1	160	100	100	160	0	0	0
NTDB id 1433	BW25113 RS17570 WP 000815987.1	160	100	100	160	0	0	0
NTDB id 1193	PA5040 NP 253727.1	160	100	100	160	0	0	0
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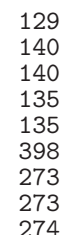
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NTDB id 1175 A1552VC RS12105 WP 000788426.1	NQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVV	116
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consensus		

NTDB id 1465	ASU2 RS06170 WP 014991934.1	MKRM.....ILLIF.F.....ILPPVIAHAFSISLKNAPTAKILSYLAEEHGKNIIVLSDNITETNT	55
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consensus		* * * * *	

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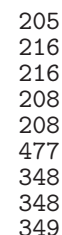
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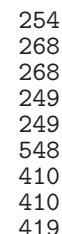
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consensus

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