

NTDB id 1384 A4U84 RS08525 WP 010787006.1	MS.....QYTIIC.....EQQTERIFDIISPSRW	22
NTDB id 993534 AAIB13 RS06535 WP 014668226.1	MFHENPLCTV.....SMTGTEVFEIQQALW	25
NTDB id 1366 NTHI RS01950 WP 011271972.1	MTSYALLHTQRV.....TAQNGEIFTISPDLW	27
NTDB id 1355 HI 0298 AAC21962.1	MTSYALLHTQRV.....TAKNGEVFTISPDLW	27
NTDB id 1168 A1552VC RS11070 WP 000957200.1	MLTINLVAITLRQAELTSATQEQAVVTQVSASGTSVPEALLLELSIFHAQELTEQLSHIFGLPETDLSRYDYANL	72
NTDB id 1293 VP RS12240 WP 005479695.1	MHSNLSTILRQKGLITFSQEESLIEQVKASGISMPEALLSSGGFTTSSELAEHLSSIFGLSQPELSQYEEYASL	72
NTDB id 1401 DSB67 RS12665 WP 010643259.1	MLTINLSTVLRQAGILITFSQEESLIEQVKASGISMPEALLSSGGFFQSHELTEHLSSVFGLSCTSLNQYEEYASL	72
NTDB id 1251 GC085 RS07725 WP 011946523.1	MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVKNKILSAEQIALTAQAQNFVGPMLDINCIDVGTI	80
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MSALHTSPKFTGFFRRLVEEKHVSAAATMQTALDAAKRAKQDITVAYLIEEVHLSPLSLAETISAEFAEYPFDLDVYDTSQI	80
NTDB id 1018 ACIAD RS01685 WP 004920473.1	MSAFTTPPKFSGFIRRLVEEGYVNAQNMQALLEKAKKFKQDITVPYLLIDNFSISPLTIAEIIISLEFGEPLLDLVGFDPALF	80
consensus	*.....***.....*	

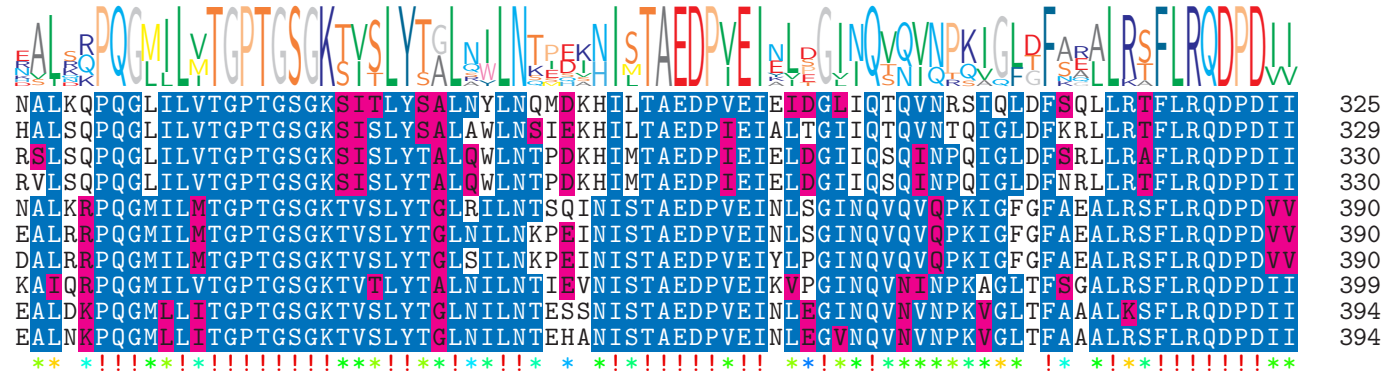
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NTDB id	1355 HI 0298 AAC21962.1	ERNQQQQSLLLRYFALPLKEENNRLWLGVDLSNLSACETIAFITGKPVEPILLESSQLKELLQLTPNQMQVEEQVK..	105
NTDB id	1168 AI552VC RS11070 WP 000957200.1	CQQLGLRELITRYDALPTAKQGNNLLLVASDPTLLQAEEEFRAFATGLQVELALADHRALQAATIRRLYGRSIQGAANQGG.	151
NTDB id	1293 VP RS12240 WP 005479695.1	CQQLGLRELITRHNALPLHRTPTSTLLLAVADPTNQQAEDDFRAFATGLQVELVLADFRELSTAIRRLYGRSLSHKESGLK.	151
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consensus		***** *** ** * ! *	! ***** * * * ! ***** ! *** * * *

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NTDB id	1366 NTHI RS01950 WP 011271972.1	FYCHQEETHFEQEDDEPVIIRLLNQIFESALQKNASDIHLETADQFQVRFRIDGVLQPPLISKIF	170
NTDB id	1355 HI 0298 AAC21962.1	FYQHQEETHFEQEDDEPVIIRLLNQIFESALQKNASDIHLETADQFQVRFRIDGVLQPPLISKIF	170
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NTDB id	1018 ACIAD RS01685 WP 004920473.1	.ELD...LDVEVDPHTTDDDDEDDKLKDPAIVKYINKLLIDAIRMSASDLHFEPYEKSYYRVRYVDGVLRLIATPPLQL	234
consensus		* * ** * * !*****!!*!*!*** ****!	

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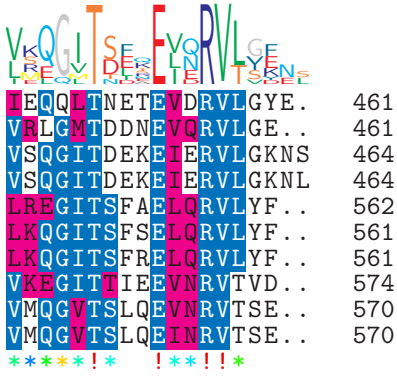
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consensus



☐ non conserved
☒ similar
☒ >= 50% conserved