

logo

NTDB id 1161 ABDM36 RS11380 WP 000340098.1

NTDB id 1147 A1552VC RS01790 WP 000340098.1

NTDB id 35644 VEA RS05055 WP 031780576.1

NTDB id 1291 VP RS12200 WP 005479697.1

consensus

MDSIEKRPRTRLSPQKRKQLMEIALEVFAKRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDDVLNFVVRQYSNFL

MDASIEKRPRTRLSPQKRKQLMEIALEVFAKRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDDVLNFVVRQYSNFL

.MDSIAKRPRTRLSPQKRKQLMEIALEVFAKRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDDVLNFVVRQYSNFL

.MDSIAKRPRTRLSPLKRKQLMEIALEVFAKRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDDVLNFHVVRQFSNFL

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80807979

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NTDB id 1147 A1552VC RS01790 WP 000340098.1

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consensus

TDHIDLDLHVRIQLTITNEVVKLVSTDCHWLKVFWEFSASTRDEVWPLFVSTNRTNQLLYQNMFKAMERGEYCEQHDV

SDHIDLDLDAKEITANTCKAMTELAHQDCHWLKVFWEFSASTRDEVWPLFVSTNRTNQLLYQNMFKAMERGEYCEQHDV

TDHIDLDLDVKTNIQTLCKEMVKLAMTDCHWLKVFWEFSASTRDEVWPLFVSTNRTNQLLYQNMFKAMERGEYCEQHDV

SDNIDLDLHARENIANITNAMIELVVSQDCHWLKVFWEFSASTRDEVWPLFVSTNRTNQLLYQNMFKAMERGEYCEQHDV

SDNIDLDLHARENIANITNAMIELVVSQDCHWLKVFWEFSASTRDEVWPLFVSTNRTNQLLYQNMFKAMERGEYCEQHDV

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consensus

ENMASLFHGI FYSIFLQVNRLEKEQEEVYKLVSSYLDMLCIYNREH

DNMASLFHGI FYSIFLQVNRLEKEQEEVYKLVSSYLDMLCIYNREH

DNMASLFHGI FYSIFLQVNRLEKEQEEVYKLVSSYLDMLCIYNREH

EHLANLFHGICYSLFVQANRFKGEAELKELVSAAYLDMLCIYNREH

EHLANLFHGICYSLFVQANRFKGEAELKELVSAAYLDMLCIYNREH

***! !!!!!*!*!*! !* * * *! *!! !!!!!!! **

203203204204

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved