

[illegible]

logo

NTDB id 1168 A1552VC RS11070 WP 000957200.1 CQQGLGRELITRYDALPIAKQGNLLLAVSDPTLLQAEEEFRFATGLQVELALADHRA LQAAIRRLYGRSIQGAANQGKE 152

NTDB id 1251 GC085 RS07725 WP 011946523.1 PVNLVNEKLIKRRHAMVPLFSRGNTNLYLATDDPSKQASLKEIQFHTGLNTHAIVETDKLSALIDNLLTAKESQGLSEYVE 160

NTDB id 1039666 ACCQ18 RS05745 WP 005915819.1 0

NTDB id 1018 ACIAD RS01685 WP 004920473.1 LKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNIEAMD A IRFNSK LKVEPTIVEHDKLERLISEHFVEETHFNFDTE.. 158

NTDB id 1058 ABD1 RS01615 WP 001274986.1 PKDLVDQKLILKHRILPLIQRGQILYVATSNPSNIEAIDAIRFNSKLLVEPVIVEHHKLEKVLGQHFAESSFDFNDE.. 158

consensus

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NTDB id 1251 GC085 RS07725 WP 011946523.1
NTDB id 1039666 ACCQ18 RS05745 WP 005915819.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1

consensus

ISQD...ELANLVKVSDDDELQSIEDLSQDDSPVSRFINQVLLDAVRKGASDIHFEPYENQYRIRLRCDGILVETQQPASHL 230
DSGDLEGLEISAD.DEDQDIDTATSVTDDAPIVKFVNKILLDAIRQGASDIHFEPYERFYRIRYRQDGI LHEVATPPASL 239
.....
.ELD...LDVEVDPHTTDDDEDKCLKDEAPIVKYINKLLIDAIRMSASDLHFEPYEKSYRVRYRV DGVLRLLIATPPLQL 234
.EFD...LDVNL DGP T AQ EEE EAPQGDEAPIVKYINKLLIDAIRMGASDLHFEPYEKSYRVRYRV DGVLRQIANPPLQL 234

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NTDB id 1168 A1552VC RS11070 WP 000957200.1	MVGEIRDLETAETIAKAAQTGHLVSTLHTNSAAETLRLRMGVPSFNIAISVNLIAQRLARRLCSQCKIPYDYPNQ	467
NTDB id 1251 GC085 RS07725 WP 011946523.1	MVGEIRDLETAETIAKAAQTGHLVSTLHTNSAAETLRLRMGVPSFNIAISVNLIAQRLARRLCKHCKIAVRP...S	479
NTDB id 1039666 ACCQ18 RS05745 WP 005915819.1	MVGEIRDLETAETIAKAAQTGHLVSTLHTNSAAETLRLRMGVPSFNIAISVNLIAQRLARRLCKHCKIAVRP...S	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	MVGEIRDLETAETIAKAAQTGHLVSTLHTNSAAETLRLRMGVPSFNIAISVNLIAQRLARRLCSQCKRPI.QVPER	473
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MVGEIRDLETAETIAKAAQTGHLVSTLHTNSAAETLRLRMGVPSFNIAISVNLIAQRLARRLCSQCKIPA.DIPKQ	473
consensus	*****	

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NTDB id 1168 A1552VC RS11070 WP 000957200.1	SLLEMGFTPSDLAPNLYEFPVGCNCTEGYKGRVGIYEVMTPEISKIMEDGNALQAAALQALGNTLRRSGL	543
NTDB id 1251 GC085 RS07725 WP 011946523.1	ALLQSQFAFQPNEL...LYEANAAAGNCTGTGYSGRVGIYEVMAFNTELAEAIMQRASIHQIERLAKANGMQLQESGL	555
NTDB id 1039666 ACCQ18 RS05745 WP 005915819.1	GLIE..LGFKEADLV..NLKLYKAVGCEQCTSGYRGRVGLFEVLPMTKELGQLIMSGGNSLDILKLAQSEGMLTIFQSGI	20
NTDB id 1018 ACIAD RS01685 WP 004920473.1	SLLE..MGFTPEDLAQPEFQIFEPVGCHDCREGYKGRVGIYEVMTITPEISKIIMEDGNALEIAATAETLGFNNLRRSGL	551
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SLLE..MGFTEQDLAHPDFRVEFQPVGCLCREGYKGRVGIYEVMTITPEISKIIMEDGNALEIAAASEKLGFNLRRLSGL	551
consensus	*** ****	

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NTDB id 1168 A1552VC RS11070 WP 000957200.1	EKIREGITSFAELQRVLYF	562
NTDB id 1251 GC085 RS07725 WP 011946523.1	EKVKEGITTIEEVNRVTVD	574
NTDB id 1039666 ACCQ18 RS05745 WP 005915819.1	MKAAHGVTSLAEINRVTKD	39
NTDB id 1018 ACIAD RS01685 WP 004920473.1	KKVMQGVTSLQEINRVTS	570
NTDB id 1058 ABD1 RS01615 WP 001274986.1	KKVMQGVTSLQEVNRVTSE	570
consensus	!* !*!*** !*!!* *	

- X non conserved
- X similar
- X ≥ 50% conserved