logo
NTDB id 1111 NGFG RS09220 WP 003689814.1 MSDLS...VLSPFA.VPLAAVLGLLVGSFLNVVIYRVPVMMERGWTVFAKEHLNLPLTDDESRTFNLMKPDSCCPKCRVP 76
NTDB id 1061 ABD1 RS18470 WP 001152280.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPKMMEQDWQQECQMLLNPEQPIIDHERLTLNKPASSCPACQQP 80
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 MQDIIAYFIQNLTALYIAVALVSLCIGSFLNVVIYRTPRMMEQDWQQECQMLLNPEQPIIDHEKLTLSKPASSCPACHQP 80
NTDB id 83174 L6418 RS12090 WP 237247175.1 MTELFQLFKSEPLVFAIACGLLGLVVGSFLNVVIHRLPRMMERDWHAQCAELNGQS..PEPAPRYNLLVPRSACPHCGHA 78
NTDB id 1403 DSB67 RS12675 WP 010643256.1 ....MEVFQYYPWLFVVFATIFGLIVGSFLNVVIYRLPKIMELEWRRECAESFPEYKIEPPKETLTLSVPRSSCQQCGTQ 76
NTDB id 1170 A1552VC RS11080 WP 000418747.1 ....MELFYFYPWLFPVLATLFGLIVGSFLNVVIYRLPKIMEREWRAECAASFPEYGITPPEGKLTLSLPRSTCPHCQTP 76
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NTDB id 1111 NGFG RS09220 WP 003689814.1 IRAWQNIPIVSYLLLRGKCASCQTKISIRYPLIELLTGVLFGLVAWQYGWSWITLGGLILTAFLISLTFIDADTQYLPDS 156
NTDB id 1061 ABD1 RS18470 WP 001152280.1 IRWYQNIPVISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPDR 160
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 IRWYQNIPLISWLVLRGKCGHCQHPISIRYPAIELLTMLCSLVVVMVFGPTIQMLFGLVLTWVLIALTFIDFDTQLLPDR 160
NTDB id 83174 L6418 RS12090 WP 237247175.1 ISAAENIPVLSYLLLRGKCKGCGAAISPRYPTVEAVSGILSAYAAWHFGFGIAAVAALVFIWALLALTFIDFDTQLLPDD 158
NTDB id 1403 DSB67 RS12675 WP 010643256.1 IRIRDNIPVISWLLLRGKCHNCQAPINARYPLIELLTAFCSGFIAFHFGFSYFTVALVFFTFVLIAATFIDLDTMLLPDQ 156
NTDB id 1170 A1552VC RS11080 WP 000418747.1 IRVIDNIPLLSWLALRGQCSHCKAPISARYPLIELLTALMSLVIATHFPFGVFAVALLFFSYVLIAATFIDFDTLLLPDQ 156
consensus
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NTDB id 1111 NGFG RS09220 WP 003689814.1 MTLPLIWLGLIFNLDGGF.VPLQSAVLGAVAGYSSLWLLCAVYKLLTGKTGMGNGDFKLIAALGAWLGISALPVLIFVSS 235
NTDB id 1061 ABD1 RS18470 WP 001152280.1 FTLPLAALGLGINTFNIY.TSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLSS 239
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 FTLPLAALGLGINTFNIY.TSPNSAIWGYLIGFLCLWIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLIVLLSS 239
NTDB id 83174 L6418 RS12090 WP 237247175.1 ITLPLLWLGLLLNLNGVY.TSLPNAVIGAAAGYLCLWSVYWAFKLTTGKEGMGYGDFKLLAVIGAWLGWQLLPLVILLSS 237
NTDB id 1403 DSB67 RS12675 WP 010643256.1 LTLPLMWTGIALALAGISPVSLQDSIIGAMAGYLCLWSVYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQSLPMIILLSS 236
NTDB id 1170 A1552VC RS11080 WP 000418747.1 LTLPLLWGGIALALLGFSPVSLSDAVIGAMAGYLSLWSIYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQQLPVIVLLSS 236
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NTDB id 1111 NGFG RS09220 WP 003689814.1 LIGLVAAIVM....RVAKGRHFAFGPALTVSGWIIFTANDSVWRAVNWWLTHPVR... 286
NTDB id 1061 ABD1 RS18470 WP 001152280.1 LLGAIIGIILLKLRN..DNQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 1080 FDQ49 RS18305 WP 022575844.1 LLGAIIGIILLKLRN..DNQPFAFGPYIAIAGWVAFLWGDQI...MKIYLGG..... 286
NTDB id 83174 L6418 RS12090 WP 237247175.1 LVGAVVGIALILFAKHGRNIPIPFGPYLAGGGLIALFWGLAL...TQSYLQLLAL... 289
NTDB id 1403 DSB67 RS12675 WP 010643256.1 VVGVIFGLIQLRLQKQGIEKAFPFGPYLAIAGWVSLIWGHQI...LNWYFTSILGV.. 289
NTDB id 1170 A1552VC RS11080 WP 000418747.1 VVGVIFGLIQLRQQKKGIDMAFPFGPYLAIAGWFALLWGDKV...IDWYFTTWVGQPL 291
consensus
X non conserved
X similar
X ≥50% conserved