logo
NTDB id 1127514 FGL12 RS08915 WP 012130787.1 MENKKISLFTQMKPYIKGFQLPFVIAVTGAVISSIITVFGPYKLREITNLITDGLVGEMDLGAISNIGLFLAFLYIVGSL 80
NTDB id 370 SMU RS04260 WP 002263294.1 MENKKKSLLSQMAPYLKGYKALFGLAVIFTIVSSTITVIGPDRLKEMTDTMTKGLAGKIDLDKIGEIALTLALLYFAGAL 80
consensus
logo
NTDB id 1127514 FGL12 RS08915 WP 012130787.1 LNYAQGYAISDMIQHFSKRLRTAIAEKINRLPLSYFDSHSQGDTLSRVTNDVDTVGQSLTQSLGTLISSVLLLIAVLFMM 160
NTDB id 370 SMU RS04260 WP 002263294.1 VSYTASFIVSTLIQKFSQRLRNAIADKINKVPLKYFDSHSQGDTLSRVTNDVDLMTQSFNQSLVSMVAAIILLIGSIFMM 160
consensus
logo
NTDB id 1127514 FGL12 RS08915 WP 012130787.1 FYTNVVLSFVTIGSVLLGFVFSAVLMAKSQIYFQEQQSNLAAINGYVEEMYSGHNVITSYNGAEKTRLAFNALNQNLYTS 240
NTDB id 370 SMU RS04260 WP 002263294.1 IKTNGALAATAILSVFAGFVLSTVIMAKSQPLFKKQQANLADVSGYVEEVYSGHNVVSSYNAIQQSKKQFENLNDQLFAS 240
consensus
logo
NTDB id 1127514 FGL12 RS08915 WP 012130787.1 MWKSQFISGIMFPLMNFVGNFGYVMVVIVGATMAINGSISMGTIVAFMVYVRIFSQPLSQIAQGTTTLQSASAAMGRVFE 320
NTDB id 370 SMU RS04260 WP 002263294.1 MWKSQFFSGIMMPLMQFIGNFGYVMVCIVGATMAINGDITMGTIVAFMTYVRIFTQPIAQIAQGITQLQSANAAMGRVFE 320
consensus
logo
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NTDB id 370 SMU RS04260 WP 002263294.1 FLDEEEIEDENHKVKQLEKVEGNVNFDNVFFGYSPDKTIIHDFSAHAKAGQKIAIVGPTGAGKTTIVNLLMRFYEVDRGM 400
consensus
logo
NTDB id 1127514 FGL12 RS08915 WP 012130787.1 IMIDGVDIKDMQRSEVHDAFSMVLQDTWLFEGTIKENLIYNQKDITDQAVVEAAKAVGVHHFIMTLSDGYDTVLDDTVSL 480
NTDB id 370 SMU RS04260 WP 002263294.1 ISIDGVNIHDMTRKEVHDAFAMVLQDTWLFEGTVKENLIYNQKHITDEQVIAAAKAVGVHHFIKTLPKGYDTVLDDSVTL 480
consensus
logo
NTDB id 1127514 FGL12 RS08915 WP 012130787.1 SVGQKQLLTIARALLKDAPLLILDEATSSVDTRTEELIQKAMDKLMEGRTSFVIAHRLSTIRNADLILVMRDGNIIEQGS 560
NTDB id 370 SMU RS04260 WP 002263294.1 SVGQKQLLTIARALLKDAPLLILDEATSSVDTRTEELIQRAMDHLMEGRTSFVIAHRLSTIRNADLILVMRDGNIIEQGS 560
consensus

logo

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NTDB id 1127514 FGL12 RS08915 WP 012130787.1   HDQLMEQGGFYADLYNSQFVEEEQE   585
NTDB id 370 SMU RS04260 WP 002263294.1         HDQLMAENGFYADLYNSQFTEEVA.   584
consensus                                      !!!!!   !!!!!!!!!!!! !!  *
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X non conserved

X similar

X ≥ 50% conserved