

logo

NTDB id 1160447 ACI6PA RS09510 WP 057867685.1
NTDB id 370 SMU RS04260 WP 002263294.1
consensus

MENKKKSLLSQMAPYLKGYKALFGLAVIFTIVSSTITVIGPDRLKEMTDTMTKGLAGKIDLDKIGEIALTLALLYFAGAL
.....
MENKKKSLLSQMAPYLKGYKALFGLAVIFTIVSSTITVIGPDRLKEMTDTMTKGLAGKIDLDKIGEIALTLALLYFAGAL

0
80

logo

NTDB id 1160447 ACI6PA RS09510 WP 057867685.1
NTDB id 370 SMU RS04260 WP 002263294.1
consensus

VSYTASFIVSTLIQKFSQRLRNAIADKINKVPLKYFDSHSQGDLSRVTNDVDLMTQSFNQSLVSMVAAILLLIGSIFMM
.....
VSYTASFIVSTLIQKFSQRLRNAIADKINKVPLKYFDSHSQGDLSRVTNDVDLMTQSFNQSLVSMVAAILLLIGSIFMM

0
160

logo

NTDB id 1160447 ACI6PA RS09510 WP 057867685.1
NTDB id 370 SMU RS04260 WP 002263294.1
consensus

IKTNGALAATAILSVFAGFVLSTVIMAKSQPLFKKQQANLADVSGYVEEVYSGHNWVSSYNAIQSKKQFENLNDQLFAS
.....
IKTNGALAATAILSVFAGFVLSTVIMAKSQPLFKKQQANLADVSGYVEEVYSGHNWVSSYNAIQSKKQFENLNDQLFAS

0
240

logo

NTDB id 1160447 ACI6PA RS09510 WP 057867685.1
NTDB id 370 SMU RS04260 WP 002263294.1
consensus

MWKSQFFSGIMMPLMQFIGNFGYVMCIVGATMAINGDITMGTIVAFMTYVRIFTQPIAQIAQGITQLQSANAAMGRVFE
.....
MWKSQFFSGIMMPLMQFIGNFGYVMCIVGATMAINGDITMGTIVAFMTYVRIFTQPIAQIAQGITQLQSANAAMGRVFE

0
320

logo

NTDB id 1160447 ACI6PA RS09510 WP 057867685.1
NTDB id 370 SMU RS04260 WP 002263294.1
consensus

FLDEEEIEDENHKVKQLEKVEGNVFNDFVFFGYSPDKTIIHDFSAAHAKAGQKIAIVGPTGAGKTTIVNLLMRFYEVDRGM
.....
FLDEEEIEDENHKVKQLEKVEGNVFNDFVFFGYSPDKTIIHDFSAAHAKAGQKIAIVGPTGAGKTTIVNLLMRFYEVDRGM

0
400

logo

NTDB id 1160447 ACI6PA RS09510 WP 057867685.1
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consensus

ISIDGVNIHDMTRKEVHDAFAMVLQDTWLFEGTVKENLIYNQKHITDEQVIAAAKAVGVVHFIKTLPKGYDTVLDDSVTL
.....
ISIDGVNIHDMTRKEVHDAFAMVLQDTWLFEGTVKENLIYNQKHITDEQVIAAAKAVGVVHFIKTLPKGYDTVLDDSVTL

0
480

logo

NTDB id 1160447 ACI6PA RS09510 WP 057867685.1
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consensus

SVGQKQLLTARALLKDAPLLILDEATSSVDTRTEELIQRAMDHLMGRTSFVIAHCFLLIKDADLLVWRDGNIEERGT
.....
SVGQKQLLTARALLKDAPLLILDEATSSVDTRTEELIQRAMDHLMGRTSFVIAHCFLLIKDADLLVWRDGNIEERGT

27
560

logo

NTDB id 1160447 ACI6PA RS09510 WP 057867685.1
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consensus



non conserved
 similar
 ≥ 50% conserved