logo
NTDB id 196588 BHS01 RS09505 WP 109835333.1 MSTETQTSRRPAGRGFGPGRGLGRGGEKPKNIKKALSDLLKYVKPWYGGIFVAAIAACGGAYLTIIGPDRLSEITNKIIS 80
NTDB id 370 SMU RS04260 WP 002263294.1 .....MENKKKSLLSQMAPYLKGYKALFGLAVIFTIVSSTITVIGPDRLKEMTDTMTK 53
consensus

logo
NTDB id 196588 BHS01 RS09505 WP 109835333.1 GLQASFIPKGYPGHGDIDLDKIMSIGMLLIAFYAGSAILSYSQGFIMATVTQRASKKLRQDISTKINKLPLRYFDGHPYG 160
NTDB id 370 SMU RS04260 WP 002263294.1 GLA..........GKIDLDKIGEIALTLALLYFAGALVSYTASFIVSTLIQKFSQRLRNAIADKINKVPLKYFDSHSQG 122
consensus

logo
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NTDB id 370 SMU RS04260 WP 002263294.1 DTLSRVTNDVDLMTQSFNQSLVSMVAAIILLIGSIFMMIKTNGALAATAILSVFAGFVLSTVIMAKSQPLFKKQQANLAD 202
consensus

logo
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NTDB id 370 SMU RS04260 WP 002263294.1 VSGYVEEVYSGHNVVSSYNAIQQSKKQFENLNDQLFASMWKSQFFSGIMMPLMQFIGNFGYVMVCIVGATMAINGDITMG 282
consensus

logo
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NTDB id 370 SMU RS04260 WP 002263294.1 TIVAFMTYVRIFTQPITAQIAQGITQLQSANAAMGRVFEFLDEEEIEDENHKVKQLEKVEGNVNFDNVFFGYSPDKTIIHD 362
consensus

logo
NTDB id 196588 BHS01 RS09505 WP 109835333.1 FSAKAEPGQKIAIVGPTGAGKSTMVNLLMRFYDTTSGQIIIDGTPIKEITREAVHDQFSMVLQDTWLFEGTVKENLIYNQ 480
NTDB id 370 SMU RS04260 WP 002263294.1 FSAHAKAGQKIAIVGPTGAGKTTIVNLLMRFYEVDRGMISIDGVNIHDMTRKEVHDAFAMVLQDTWLFEGTVKENLIYNQ 442
consensus

logo
NTDB id 196588 BHS01 RS09505 WP 109835333.1 ANISDQQVVDAAKAVGIDHFINTLPSGYDSMLKDADELSVGERQLLTIARALLKDAPMLILDEATSSVDTRTEELIQKAM 560
NTDB id 370 SMU RS04260 WP 002263294.1 KHITDEQVIAAAKAVGVHHFIKTLPKGYDTVLDDSVTLSVGQKQLLTIARALLKDAPLLILDEATSSVDTRTEELIQRAM 522
consensus

logo

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NTDB id 196588 BHS01 RS09505 WP 109835333.1  DKLMEGRTSFVIAHRLSTIKNADLILVMKDGDIIESGDHKALLAQDGFYADLYNSQFERVA.  621
NTDB id 370 SMU RS04260 WP 002263294.1       DHLMEGRTSFVIAHRLSTIRNADLILVMRDGNIIEQGSHDQLMAENGFYADLYNSQFTEEVA  584
consensus                                    !*!!!!!!!!!!!!!!!!!*!!!!!!!!!*!! !!! ! !  !*!  !!!!!!!!!!!     *
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X non conserved
X similar
X ≥ 50% conserved