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logo
NTDB id 1017 ACIAD RS02540 WP 011182074.1   MSKWTINKTLGCAAIVGAFILSGCQTVPT.TKDPEKGVKVRTQLAAEYIKSGDLDSAKRALDQALKVNPKDSSANMMMGVLLQQE   84
NTDB id 127671 IX88 RS04740 WP 187406181.1  .MSTPKLKTMLCMGVAVALLASGCQTSQTVKKDPEKAVKVRTQLAAEYIRSGDLDSAKRSLDQALSVDSRDATANMMMGILLQQE   84
NTDB id 1051 ABD1 RS02465 WP 004738086.1   .MSTPKLKIMLCMGVAVALLASGCQTSQTVKKDPEKAVKVRTQLAAEYIRSGDLDSAKRSLDQALSVDSRDATANMMMGILLQQE   84
consensus                                   ******!***!*****!***!!!!!**!**!!!!*!!!!!!!!!!!!*!!!!!!!!!*!!!!!*!***!**!!!!!!*!!!!!

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NTDB id 1017 ACIAD RS02540 WP 011182074.1   GSRPNLEKADAYFERAVQLDPENAQAHNNYGTYLYQMKRYHDAIQQLTLAGSSLGYDQRYRALENIGRIYLHLGGVANAEKMFSQ   169
NTDB id 127671 IX88 RS04740 WP 187406181.1  GSKPNLEKAEHYFKRAISSEPDNAQARNNYGTYLYQMERYNDAIEQLRIAGATLGYDQRYQALVNLGRIYLKLGDIASAEKTFKQ   169
NTDB id 1051 ABD1 RS02465 WP 004738086.1   GSKPNLEKAEHYFKRAISSEPDNAQARNNYGTYLYQMERYNDAIEQFRIAGATLGYDQRYQALENLGRIYLKLGDIASAEKTFKQ   169
consensus                                   !!*!!!!!!**!!*!!****!*!!!!*!!!!!!!!!!*!!*!!!*!***!!**!!!!!!!*!!*!*!!!!!*!!**!*!!!*!*!

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NTDB id 1017 ACIAD RS02540 WP 011182074.1   ALQANRDSSVSMLELAEIYYLQQRIPAATGMYEQYVRTVGQKNQGARALWIGIRVARANADRTGVQVLVNQLRIFYPDSSEYQRY   254
NTDB id 127671 IX88 RS04740 WP 187406181.1  ALLANRDSYISMLELAEIFYLQQQIPAATQMYEQYVRTVGQKNQGARALWIGLRVARANADKMGMQVLVNQLRALFPESPEYQRY   254
NTDB id 1051 ABD1 RS02465 WP 004738086.1   ALLANRDSYISMLELAEIFYLQQQIPAATQMYEQYVRTVGQKNQGARALWIGLRVARANADKMGMQVLVNQLRALFPESPEYQRY   254
consensus                                   !!*!!!!!**!!!!!!!!!*!!!!*!!!!!*!!!!!!!!!!!!
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