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logo

NTDB id 139695 TE32 RS16945 WP 002027454.1  MS.TPKLKIMLCMGVAVALLTSGCQTSQTVKKDPEKAVKVRTQLAAEYIRSGDLDSAKRSLDQALSVDSRDATANMMMGILLQQE   84
NTDB id 1051 ABD1 RS02465 WP 004738086.1    MS.TPKLKIMLCMGVAVALLASGCQTSQTVKKDPEKAVKVRTQLAAEYIRSGDLDSAKRSLDQALSVDSRDATANMMMGILLQQE   84
NTDB id 1017 ACIAD RS02540 WP 011182074.1   MSKWTINKTLGCAAIVGAFILSGCQTVPT.TKDPEKGVKVRTQLAAEYIKSGDLDSAKRALDQALKVNPKDSSANMMMGVLLQQE   84
consensus                                   !! ****!***!*****!** !!!!!**!**!!!!*!!!!!!!!!!!!*!!!!!!!!!*!!!!!*!***!**!!!!!!*!!!!!

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NTDB id 139695 TE32 RS16945 WP 002027454.1  GSKPNLEKAEHYFKRAISSEPDNAQARNNYGTYLYQMERYNDAIEQFRIAGATLGYDQRYQALENLGRIYLKLGDIASAEKTFKQ  169
NTDB id 1051 ABD1 RS02465 WP 004738086.1    GSKPNLEKAEHYFKRAISSEPDNAQARNNYGTYLYQMERYNDAIEQFRIAGATLGYDQRYQALENLGRIYLKLGDIASAEKTFKQ  169
NTDB id 1017 ACIAD RS02540 WP 011182074.1   GSRPNLEKADAYFERAVQLDPENAQAHNNYGTYLYQMKRYHDAIQQLTLAGSSLGYDQRYRALENIGRIYLHLGGVANAEKMFSQ  169
consensus                                   !!*!!!!!!**!!*!!****!*!!!!*!!!!!!!!!!*!!*!!!!*!***!!**!!!!!!!*!!!!*!!!!!*!!**!*!!!!*!*!

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NTDB id 139695 TE32 RS16945 WP 002027454.1  ALLANRDSYISMLELAEIFYLQQQIPAATQMYEQYVRTVGQKNQGARALWIGLRVARANADKMGMQVLVNQLRALFPESPEYQRY  254
NTDB id 1051 ABD1 RS02465 WP 004738086.1    ALLANRDSYISMLELAEIFYLQQQIPAATQMYEQYVRTVGQKNQGARALWIGLRVARANADKMGMQVLVNQLRALFPESPEYQRY  254
NTDB id 1017 ACIAD RS02540 WP 011182074.1   ALQANRDSSVSMLELAEIYYLQQRIPAATGMYEQYVRTVGQKNQGARALWIGIRVARANADRTGVQVLVNQLRIFYPDSSEYQRY  254
consensus                                   !!*!!!!!**!!!!!!!!!*!!!!*!!!!!*!!!!!!!!!!!!
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