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logo
NTDB id 252328 AYP RS02645 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 252328 AYP RS02645 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 252328 AYP RS02645 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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