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logo

NTDB id 1017 ACIAD RS02540 WP 011182074.1    MSKWTINKTLGCAAIVGAFILSGCQTVPT.TKDPEKGVKVRTQLAAEYIKSGDLDSAKRALDQALKVNPKDSSANMMMGVLLQQE   84
NTDB id 81180 OCUAC RS15325 WP 180635462.1  MS.TPKLKIMLCMGVAVALLASGCQTSQTVKKDPEKAVKVRTQLAAEYIRSGDLDSAKRSLDQALSVDSRDATANMMMGILLQQE   84
NTDB id 1051 ABD1 RS02465 WP 004738086.1    MS.TPKLKIMLCMGVAVALLASGCQTSQTVKKDPEKAVKVRTQLAAEYIRSGDLDSAKRSLDQALSVDSRDATANMMMGILLQQE   84
consensus                                   !! ****!***!*****!***!!!!**!**!!!!*!!!!!!!!!!!!*!!!!!!!!!*!!!!!*!***!**!!!!!!*!!!!!

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NTDB id 1017 ACIAD RS02540 WP 011182074.1    GSRPNLEKADAYFERAVQLDPENAQAHNNYGTYLYQMKRYHDAIQQLTLAGSSLGYDQRYRALENIGRIYLHLGGVANAEKMFSQ  169
NTDB id 81180 OCUAC RS15325 WP 180635462.1  GSKSNLEKAEHYFKRAISSEPDNAQARNNYGTYLYQMERYNDAIEQFRIAGATLGYDQRYQALENLGRIYLKLGDIASAEKTFKQ  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 81180 OCUAC RS15325 WP 180635462.1  ALLANRDSYISMLELAEIFYLQQQIPAATQMYEQYVRTVGQKNQGARALWIGLRVARANTDKMGMQVLVNQLRALFPESPEYQRY  254
NTDB id 1051 ABD1 RS02465 WP 004738086.1    ALLANRDSYISMLELAEIFYLQQQIPAATQMYEQYVRTVGQKNQGARALWIGLRVARANADKMGMQVLVNQLRALFPESPEYQRY  254
consensus                                   !!*!!!!!**!!!!!!!!!*!!!!*!!!!!*!!!!!!!!!!!!
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