```
logo
NTDB id 122817 RR41 RS16455 WP 004738086.1  MS.TPKLKIMLCMGVAVALLASGCQTSQTVKKDPEKAVKVRTQLAAEYIRSGDLDSAKRSLDQALSVDSRDATANMMMGILLQQE   84
NTDB id 1051 ABD1 RS02465 WP 004738086.1    MS.TPKLKIMLCMGVAVALLASGCQTSQTVKKDPEKAVKVRTQLAAEYIRSGDLDSAKRSLDQALSVDSRDATANMMMGILLQQE   84
NTDB id 1017 ACIAD RS02540 WP 011182074.1   MSKWTINKTLGCAAIVGAFILSGCQTVPT.TKDPEKGVKVRTQLAAEYIKSGDLDSAKRALDQALKVNPKDSSANMMMGVLLQQE   84
consensus                                   !! ****!***!*****!***!!!!!**!**!!!!*!!!!!!!!!!!!*!!!!!!!!!*!!!!!*!***!**!!!!!!*!!!!!

logo
NTDB id 122817 RR41 RS16455 WP 004738086.1  GSKPNLEKAEHYFKRAISSEPDNAQARNNYGTYLYQMERYNDAIEQFRIAGATLGYDQRYQALENLGRIYLKLGDIASAEKTFKQ  169
NTDB id 1051 ABD1 RS02465 WP 004738086.1    GSKPNLEKAEHYFKRAISSEPDNAQARNNYGTYLYQMERYNDAIEQFRIAGATLGYDQRYQALENLGRIYLKLGDIASAEKTFKQ  169
NTDB id 1017 ACIAD RS02540 WP 011182074.1   GSRPNLEKADAYFERAVQLDPENAQAHNNYGTYLYQMKRYHDAIQQLTLAGSSLGYDQRYRALENIGRIYLHLGGVANAEKMFSQ  169
consensus                                   !!*!!!!!!**!!*!!****!*!!!!*!!!!!!!!!!*!!*!!!!*!***!!**!!!!!!!*!!!!*!!!!!*!!**!*!!!*!*!

logo
NTDB id 122817 RR41 RS16455 WP 004738086.1  ALLANRDSYISMLELAEIFYLQQQIPAATQMYEQYVRTVGQKNQGARALWIGLRVARANADKMGMQVLVNQLRALFPESPEYQRY  254
NTDB id 1051 ABD1 RS02465 WP 004738086.1    ALLANRDSYISMLELAEIFYLQQQIPAATQMYEQYVRTVGQKNQGARALWIGLRVARANADKMGMQVLVNQLRALFPESPEYQRY  254
NTDB id 1017 ACIAD RS02540 WP 011182074.1   ALQANRDSSVSMLELAEIYYLQQRIPAATGMYEQYVRTVGQKNQGARALWIGIRVARANADRTGVQVLVNQLRIFYPDSSEYQRY  254
consensus                                   !!*!!!!!**!!!!!!!!!*!!!!*!!!!!*!!!!!!!!!!!!
```