

logo

NTDB id 183942 A9E77 RS13540 WP 011947563.1
NTDB id 1278 LPP RS13600 WP 011947563.1
NTDB id 1262 GC085 RS13310 WP 011947563.1
consensus

MIKVLIVDDHALVRMGI RRLLEDMSD VDVVADAESGEQALAYVKTHSPDWLLDMKMPGIDGWEVT RRLKKTNPNIKVIA
MIKVLIVDDHALVRMGI RRLLEDMSD VDVVADAESGEQALAYVKTHSPDVLLDMKMPGIDGWEVT RRLKKTNPNIKVIA
MIKVLIVDDHALVRMGI RRLLEDMSD VDVVADAESGEQALAYVKTHSPDVLLDMKMPGIDGWEVT RRLKKTNPNIKVIA
MIKVLIVDDHALVRMGI RRLLEDMSD VDVVADAESGEQALAYVKTHSPDVLLDMKMPGIDGWEVT RRLKKTNPNIKVIA

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NTDB id 1262 GC085 RS13310 WP 011947563.1
consensus

VTAI CSDPLPTRVLQLGAMGYLTKE SGA EEMAAAIRK VAKGEKYL SAEIAQKMAINS LQEPQDSPFDLLSEREMQVLM I
VTAI CSDPLPTRVLQLGAMGYLTKE SGA EEMAAAIRK VAKGEKYL SAEIAQKMAINS LQEPQDSPFDLLSEREMQVLM I
VTAI CSDPLPTRVLQLGAMGYLTKE SGA EEMAAAIRK VAKGEKYL SAEIAQKMAINS LQEPQDSPFDLLSEREMQVLM I
VTAI CSDPLPTRVLQLGAMGYLTKE SGA EEMAAAIRK VAKGEKYL SAEIAQKMAINS LQEPQDSPFDLLSEREMQVLM I
VTAI CSDPLPTRVLQLGAMGYLTKE SGA EEMAAAIRK VAKGEKYL SAEIAQKMAINS LQEPQDSPFDLLSEREMQVLM I

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consensus

TSGMNVQDIADRLFSSKTINGYRYRMFEKLGIKNDVELTYLAMKHRIIEHPNDLSQDD
TSGMNVQDIADRLFSSKTINGYRYRMFEKLGIKNDVELTYLAMKHRIIEHPNDLSQDD
TSGMNVQDIADRLFSSKTINGYRYRMFEKLGIKNDVELTYLAMKHRIIEHPNDLSQDD
TSGMNVQDIADRLFSSKTINGYRYRMFEKLGIKNDVELTYLAMKHRIIEHPNDLSQDD

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219

- X non conserved
- X similar
- X $\geq 50\%$ conserved