

logo

NTDB id 88736 AAV29 RS12325 WP 043867279.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

MLAELSIKNFAII EELTY SFERGLTVLTGETGAGKSIIDAY SLLVGGRSSEFVRYGE_A KAELEGLFLL_D SGHPV_F EVCREQGI_A

MLAELSIKNFAII EELT_I SFERGLTVLTGETGAGKSIIDAV_I SLLVGGRSSEFVRYGE_T KAELEGLFLL_D SGHPV_F EVCREQGI_A

MLAELSIKNFAII EELTV_I SFERGLTVLTGETGAGKSIIDAV_I SLLVGGRSSEFVRYGE_A KAELEGLFLL_D SGHPV_L GVC_A EQGI_I

* * *

85
85

logo

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consensus

170
170

logo

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consensus

KLLKKYKQLSESEQEMAHRLDLIQFQLEEIESAKLELNEDELQEEERKQISNFEKIYESLQNAVNALRSEQGGLDIVGMASQAQL
 KLLKKYKQLSESEQEMAHRLDLIQFQLEEIESAKLELNEDELQEEERKQISNFEKIYESLQNAVNALRSEQGGLDIVGMASQAQL
 KLLKKYKQLSESEQEMAHRLDLIQFQLEEIESAKLELNEDELQEEERKQISNFEKIYESLQNAVNALRSEQGGLDIVGMASQAQL

255
255

logo

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consensus

DISDINEPLQKLSSESVS_NSYLLEDATFQMRN_LLD_LLEYDPERLN_FIETRLNEIKQLKRKYGATVEDILEY_ASKIEEEIDQIENR

340
340

logo

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consensus

DSHL^{QS}E^{EA}KKEL^{ES}VGKDVAVEAAN^{YS}SK^QIRK^ATWAKKLA^{EE}DA^{DA}THREL^QSLYMG^EKSTFDTEFL^KVRT^{DSSNS}APRAE^EEAP^VVNGQ^{SV}QLT^{EQ}GID^{EK}
 DSHLEALKKEL^{ES}VGKDVAVEAAN^{LS}SK^QIRKA^AWAKKLA^{EA}THREL^QSLYMG^EKSTFDTEFL^KVRT^{DPSAS}EAP^VVNGQ^{SV}QLT^{QK}GID^E
 DSHL^{QS}LKKEL^{ES}VGKDVAVEAAN^{YS}SK^QIRKT^TWAKKLA^{EE}DA^{DA}THREL^QSLYMG^EKSTFDTEFL^KVRT^{DSSNS}APRAE^EEAP^VVNGQ^{SV}QLT^{EQ}GID^{EK}

425
425

logo

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LVKFLISTNTGEPLKSLKVASGGELSRVMLAKSIFSSQQDVTSLIFDEVDTGVSGRVAQAIAEKIHKVSTGSQVLCITHLPQV
 LVKFLISTNTGEPLKPLSKVASGGELSRVMLAKSIFSSQQDVTSLIFDEVDTGVSGRVAQAIAEKIHKVSTGSQVLCITHLPQV
 LVKFLISTNTGEPLKSLKVASGGELSRVMLAKSIFSSQQDVTSLIFDEVDTGVSGRVAQAIAEKIHKVSTGSQVLCITHLPQV

510
510

logo

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AAMADTHLYIAKEFKDGRITTHVTPLSKQDKVAEIGRMAGVEVTDLTKRHAQELLKQAGQVKTTG
 AAMADTHLYIAKEFKDGRITTHVTPLSKQDKVAEIGRMAGVEVTDLTKRHAQELLKQAGQVKTTG
 AAMADTHLYIAKEFKDGRITTHVTPLSKQDKVAEIGRMAGVEVTDLTKRHAQELLKQADQVKTTG

576
576

 non conserved
 similar
 $\geq 50\%$ conserved