

logo

NTDB id 358345 SB21 RS11650 WP 032874060.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

MLAELSIKNFAII EELTY SFERGLTVLTGETGAGKSIIDAY SLLVGGRSSEFVRYGETAKAELEGLFLLDSGHPVFDVCREQGI
MLAELSIKNFAII EELTISFERGLTVLTGETGAGKSIIDAVSLLVGGRSSEFVRYGETAKAELEGLFLLDSGHPVFDVCREQGI
MLAELSIKNFAII EELTVSFERGLTVLTGETGAGKSIIDAVSLLVGGRSSEFVRYGETAKAELEGLFLLDSGHPVLDVCAEQGI

85
85

logo

NTDB id 358345 SB21 RS11650 WP 032874060.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

170
170

logo

NTDB id 358345 SB21 RS11650 WP 032874060.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

KLLKKYKQLSESEQEMAHRLDLIQFQLEEIESAKLELNEDELQEEERKQISNFEKIYESLQNAVNALRSEQGGLDIVGMASQAQL
 KLLKKYKQLSESEQEMAHRLDLIQFQLEEIESAKLELNEDELQEEERKQISNFEKIYESLQNAVNALRSEQGGLDIVGMASQAQL
 *

255
255

logo

NTDB id 358345 SB21 RS11650 WP 032874060.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

DISDINEPLQKMSSESVS_NSYLLEDATFQMR_NLD_ELE_FYDPERLN_FETRLNEIKQLKRKYGATVEDILEY_ASKIEEEIDQIENR

340
340

logo

NTDB id 358345 SB21 RS11650 WP 032874060.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

DSHL^{QS}_{EA}KKEL^{ES}_{DS}VGKDVAVEAAN^{YS}_{LS}Q^{SR}_{KR}AWAKKLA^{EE}_{DA}IR^{HQ}_{EL}KSLYMG^{KE}_{ST}FDTEFL^{VR}_{KV}T^{DSSNS}_{APRAE}EAP^{YV}_{LV}NGQPVL^{TQ}_{EQ}GID
 DSHLEALKKEL^{ES}_{DS}VGKDVAVEAAN^{LS}_{SK}IR^{KA}WAKKLA^{EA}_{TH}QELKSLYMG^{KE}_{ST}FDTEFL^{VKT}_{DPSAS}EAP^{YV}_{LV}NGQPVL^{TQK}_{GID}
 DSHL^{QS}_{LS}KKEL^{ES}_{DS}VGKDVAVEAAN^{YS}_{LS}Q^{SR}_{KR}AWAKKLA^{EE}_{DA}IR^{HQ}_{EL}KSLYME^{KE}_{ST}FDTEFL^{VR}_{KV}T^{DSSNS}_{APRAE}EAP^{YV}_{LV}NGQPVL^{TQ}_{EQ}GID
 * * * * *

425
425

logo

NTDB id 358345 SB21 RS11650 WP 032874060.1
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consensus

LVKFLISTNTGEPLKSLKVASGGELSRVMLAKSIFSSQQDVTSLIFDEVDTGVSGRVAQAIAEKIHKVSTGSQVLCITHLPQV
 LVKFLISTNTGEPLKPLSKVASGGELSRVMLAKSIFSSQQDVTSLIFDEVDTGVSGRVAQAIAEKIHKVSTGSQVLCITHLPQV
 LVKFLISTNTGEPLKSLKVASGGELSRVMLAKSIFSSQQDVTSLIFDEVDTGVSGRVAQAIAEKIHKVSTGSQVLCITHLPQV

510
510

logo

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NTDB id 352 BSU 24240 NP 390304.2
consensus

AAMADTHLYIAKEFKDGRITTV^RVTPLSKQ^DKVAEIGRMAGVEVTDLTKRHA^KELLKQAGQVKT^DTG
 AAMADTHLLIAKEFKDGRITTV^RVTPLSKQ^DKVAEIGRMAGVEVTDLTKRHA^KELLKQAGQVKT^DTG
 AAMADTHLYIAKE^LKDGRITTV^RVK^{*}PLSKQ^DKVAEIGRMAGVEVTDLTKRHA^KELLKQADQVKT^DTG

576
576

 non conserved
 similar
 $\geq 50\%$ conserved