

logo

NTDB id 329626 BVMH RS12330 WP 007408359.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

MLAELSIKNFAII EELTY SFERGLTVLTGETGAGKSIIDAY SLLVGGRSSEFVRYGE_A KAELEGLFLL_D SGHPV_F EVC_A REQGI

85
85

logo

NTDB id 329626 BVMH RS12330 WP 007408359.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

170
170

logo

NTDB id 329626 BVMH RS12330 WP 007408359.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

KLLKKYKQLSESEQEMAHRLDLQFQLEEIESAKLELNEDEQLQEERKQISNFEEKIYESLQNAVNALRSEQGGLDVGVMASQAQL
 KLLKKYKQLSESEQEMAHRLDLQFQLEEIESAKLELNEDEQLQEERKQISNFEEKIYESLQNAVNALRSEQGGLDVGVMASQAQL
 *

255
255

logo

NTDB id 329626 BVMH RS12330 WP 007408359.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

DISDINEPLQKMSSEVS_NSYLLEDATFQMR_NLD_LE_FYDPERL_NY_FETRLNEIKQLKRKYGATVEDILEY_ASKIEEEIDQIENR

340
340

logo

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425
425

logo

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consensus

LVKFLISTNTGEPLK_SLSKVASGGELSRVMLA_MKSIFSSQQDVTSIFDEVDTGVSGRVAQAIAEKIHKVSTGSQVLCITHLPQV
 LVKFLISTNTGEPLK_PLSKVASGGELSRVMLA_MKSIFSSQQDVTSIFDEVDTGVSGRVAQAIAEKIHKVSTGSQVLCITHLPQV
 LVKFLISTNTGEPLK_SLSKVASGGELSRVMLA_IKSIFSSQQDVTSIFDEVDTGVSGRVAQAIAEKIHKVSTGSQVLCITHLPQV

510
510

logo

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consensus

AAMADTHLYIAKEFKDGRITTH^RVTPLSKQ^EKVAEIGRMAGVEVTDLTKRHA^QELLKQAGQVKTTG
 AAMADTHLLIAKEFKDGRITTH^VTPLSKQ^DKVAEIGRMAGVEVTDLTKRHA^QELLKQAGQVKTTG
 AAMADTHLYIAKELKDGRITTH^RVKPLSKQ^EKVAEIGRMAGVEVTDLTKRHA^KELLKQADQVKTTG

576
576

 non conserved
 similar
 $\geq 50\%$ conserved