

logo

NTDB id 1121441 EL130 RS08315 WP 002272311.1
NTDB id 405 SMU RS06370 WP 002262721.1
consensus

MGRGKLTPODLEAMKAI SANLQRLLAESGMKQSHLATILNIPTSSLNEYVKGKSLPKIGNIQKIADYFGLQKSDIDPRFA
MGRGKLTPODLEAMKAI SANLQRLLAESGMKQSHLATILNIPTSSLNEYVKGKSLPKIGNIQKIADYFGLQKSDIDPRFA
MGRGKLTPODLEAMKAI SANLQRLLAESGMKQSHLATILNIPTSSLNEYVKGKSLPKIGNIQKIADYFGLQKSDIDPRFA
!!

80
80

logo

NTDB id 1121441 EL130 RS08315 WP 002272311.1
NTDB id 405 SMU RS06370 WP 002262721.1
consensus

SKKDSSTLDKINHIAKQLEMKRQIKVLDFCYQSQSKEQDETASLSASDETTEDKVVADKSNHTQYQKVVDSFIAKELNDQTI
SKKDSSTLDKINHIAKQLEMKRQIKVLDFCYQSQSKEQDETASLSASDETTEDKVVADKSNHTQYQKVVDSFIAKELNDQTI
SKKDSSTLDKINHIAKQLEMKRQIKVLDFCYQSQSKEQDETASLSASDETTEDKVVADKSNHTQYQKVVDSFIAKELNDQTI
!!*

160
160

logo

NTDB id 1121441 EL130 RS08315 WP 002272311.1
NTDB id 405 SMU RS06370 WP 002262721.1
consensus

IHWDNVIKAKDLYQTFGRQYENIQIYGEVSAGTGIWVGHEKQETIKYPIPIPEHDIALTVNGNSMEPLFYNGDVI FVKKT
IHWDNVIKAKDLYQTFGRQYENIQIYGEVSAGTGIWVGHEKQETIKYPIPIPEHDIALTVNGNSMEPLFYNGDVI FVKKT
IHWDNVIKAKDLYQTFGRQYENIQIYGEVSAGTGIWVGHEKQETIKYPIPIPEHDIALTVNGNSMEPLFYNGDVI FVKKT
!!

240
240

logo

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consensus

KAIHHGQIIVVIVNNEAYVKKLYRKNKEIRLISLNPDYDDIILKEDDTIEVIGTVIT
KAIHHGQIIVVIVNNEAYVKKLYRKNKEIRLISLNPDYDDIILKEDDTIEVIGTVIT
KAIHHGQIIVVIVNNEAYVKKLYRKNKEIRLISLNPDYDDIILKEDDTIEVIGTVIT
!!

297
297

- non conserved
- similar
- ≥ 50% conserved