

logo

NTDB id 234772 CD084 RS03675 WP 040083678.1
NTDB id 116 BSU 08520 NP 388733.1
consensus

MPFITKIS^AQKKNTERFNI^FLDDKYAF^SVDADVL^VK^FEL^KKKGKEL^DDDLDI^IEIQY^GDEV^KKKGF^NRAL^FFLSY^RMR^STKEV

MPFITKIS^AQKKNTERFNI^FLDDKYAF^SVDADVL^VK^FEL^KKKGKEL^DDDLDI^IEIQY^GDEV^KKKGF^NRAL^FFLSY^RMR^STKEV

MPFITKIS^TQKKNTERFNI^FLDDKYAF^SVDADVL^VK^FEL^KKKGKEL^DDDLDI^IEIQY^GDEV^KKKGF^NRAL^FFLSY^RMR^STKEV

!!

80
80

logo

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NTDB id 116 BSU 08520 NP 388733.1
consensus

EDHL^KKKKETS^PPPVIAE^VIH^RLNDY^KYLND^QEFAAA^YVSTH^KKTNG^KGP^DVL^FREL^RAKG^IDD^DTI^KEAL^SSFS^FED^QTRE

EDHL^KKKKETS^PPPVIAE^VIH^RLNDY^KYLND^QEFAAA^YVSTH^KKTNG^KGP^DVL^FREL^RAKG^IDD^DTI^KEAL^SSFS^FED^QTRE

EDHL^KKKKETS^PPPVIAE^VIH^RLNDY^KYLND^QEFAAA^YVSTH^KKTNG^KGP^DVL^FREL^RAKG^IDD^DTI^KEAL^SSFS^FED^QTRE

!!

160
160

logo

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NTDB id 116 BSU 08520 NP 388733.1
consensus

AV^KH^VE^KLL^KDK^KLST^KEL^KQRA^QML^QLQR^KGF^SFD^VISA^ALE^QIEY^ENDED^TEKE^ALR^LHA^EKA^LRKY^RYD^GSY^ESAM^K

AV^KH^VE^KLL^KDK^KLST^KEL^KQRA^QML^QLQR^KGF^SFD^VISA^ALE^QIEY^ENDED^TEKE^ALR^LHA^EKA^LRKY^RYD^GSY^ESAM^K

AV^KH^VE^KLL^KDK^KLST^KEL^KQRA^QML^QLQR^KGF^SFD^VISA^AMD^QIEY^ENDED^TEKE^ALR^LHA^EKA^FRKY^RYD^GSY^ESAM^K

!!

240
240

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consensus

V^KQ^FL^FR^KG^FS^LD^LI^EQ^FL^HEEEE^Y

V^KQ^FL^FR^KG^FS^LD^LI^EQ^FL^HEEEE[.] 263

V^KQ^FL^FR^KG^FS^LD^LI^EQ^LL^QEEEE^Y 264

!!

^X non conserved
^X similar
^X ≥ 50% conserved