

logo

NTDB id 83 BSU 35490 NP 391429.1
NTDB id 391751 GAN20 RS03230 WP 014613482.1
NTDB id 4 SA RS09755 WP 000153535.1
consensus

MTKVNIVITLQLFIEVKRLDFESTFAADDDEARIVEHYHWTLTLENNKILVELYES
 MTKVNIVITDDHQLFREGVKRILDFEPTFEVVAEGDDGDEAARIVEHYHPDVIIMDINMPNVNGVEATKQLVELYPESKV
 MTIKVLVDDHEMVRIGISSYLSTQPDIEVVGEGASGKEATAKAHELKPDILMDLMTMDMGVEATMQIKKDLPHITKV

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consensus

Sequence	160	161	162
160	TILSIHDDENYVTHALKTGARGYLKEMDADTILIEAVKVVAEGGSYLHPKVTHNLVNEFRRLATSGVSAHPQHEVYPEIR		
142	VMLTSYIEDKEVYRALDAGVDSYILKTTTSANDIAEAVRKTSRGESVFEEVLVKMRNRMKQ		
142	LMLTSFIEDKEVYRALDAGVDSYILKTTTSAKDIAEAVRKTSRGESVFEEVLVKMRNRMKQ		

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ELYE^MLTEREMEILLIAKGSNQEIASASHITIKTVKTHVSNILSKLEVQDRTQAVIYAFQHGLIE

RPLHT^RRCVCVQLSDKRGESLFISEIKTVKNHVSNI^LLQKMNVDRTQAVVAIKNGWVEMR 229

ELYELLTEREMEILLISKGYSNQEIASASHITIKTVKTHVSNILSKLEVQDRTQAVIYAFQHGLIE.. 209

ELYEMLTEREMEILLIAKGSYNQEIASASHITIKTVKTHVSNILSKLEVQDRTQAVIYAFQHNLIQ.. 209

- X non conserved
- X similar
- X > 50% conserved