

logo

NTDB id 146 SP RS08570 WP 000614538.1
NTDB id 216 SPD RS08205 WP 000614552.1
NTDB id 182 SPR RS07820 WP 000614552.1
NTDB id 257 KZH43 RS07655 WP 220041236.1
NTDB id 467 HSISS4 RS06915 WP 021143821.1
NTDB id 1138061 DQL37 RS07040 WP 063812253.1
NTDB id 384 SMU RS02325 WP 002263039.1
consensus



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consensus



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NTDB id 146 SP RS08570 WP 000614538.1
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NTDB id 146 SP RS08570 WP 000614538.1	SIKNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVLV.	AASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGE	396					
NTDB id 216 SPD RS08205 WP 000614552.1	SIKNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVLV.	AASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGE	396					
NTDB id 182 SPR RS07820 WP 000614552.1	SIKNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVLV.	AASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGE	396					
NTDB id 257 KZH43 RS07655 WP 220041236.1	SIKNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVLV.	AASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGE	396					
NTDB id 467 HSISS4 RS06915 WP 021143821.1	.TPAPSKQPRKKKTPAKKKKKK	NFFSTLLKVF	FLGLVFI	GIIIFAYLVFTNPDNVQVPNVVGGQELSTAQTK	EGAGFKVGE	396		
NTDB id 1138061 DQL37 RS07040 WP 063812253.1	SRLDQTNQTDTLQPPTKKKSGRFLGTLFKIL	SFFIVGV	ALFTYILITKPTSVKVPNVAGTSLKVAQ	ELYDVG	GLKV	GK	398	
NTDB id 384 SMU RS02325 WP 002263039.1	DKVDHKS.KPKTKPQPKPKRHLAR	FFKIFFILVAL	GI	AVLTYLVLT	KPSTVSV	PDVSGDKLSTAKMI	IRKSLKLVGD	396
consensus	*****!*****!*****!*****!*****!*****!*****!*****!							

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NTDB id 146 SP RS08570 WP 000614538.1	EKTEASEKVEEGRIIRTDPGAGTGRKEG	TKINLVVSSGKQSFQISNYVGRKSSDVIAELKE	KKVPDNLKIEEEEEESNES	475	
NTDB id 216 SPD RS08205 WP 000614552.1	EKTEASEKVEEGRIIRTDPGAGTGRKEG	TKINLVVSSGKQSFQISNYVGRKSSDVIAELKE	KKVPDNLKIEEEEEESNES	475	
NTDB id 182 SPR RS07820 WP 000614552.1	EKTEASEKVEEGRIIRTDPGAGTGRKEG	TKINLVVSSGKQSFQISNYVGRKSSDVIAELKE	KKVPDNLKIEEEEEESNES	475	
NTDB id 257 KZH43 RS07655 WP 220041236.1	EKTEASEKVEEGRIIRTDPGAGTGRKEG	TKINLVVSSGKQSFQISNYVGRKSSDVIAELKE	KKVPDNLKIEEEEEESNES	475	
NTDB id 467 HSISS4 RS06915 WP 021143821.1	VKEVEDPSVDTCKVILKTDPTACTT	RKEGSSIDVYVSSGSKGFALKDYK	CKNYKDAIEDLTSNFGVSEDDQIDIQHVED	DSA	476
NTDB id 1138061 DQL37 RS07040 WP 063812253.1	IRQIESDTVAEGNVVRTDPKAGTAKRQC	SSITLYVSTGNKGFDMENYKGLDYQEAMNS	LIEITYGVPKSKIKIERIVTNEY		478
NTDB id 384 SMU RS02325 WP 002263039.1	VQKIEDLVNGACKVVRTNPSACSKKREGSS	VDIFVA.SAKGFKMEDYTGQDYKDAIDN	LTTNNYCVSRDSIVLKEVSSDDY		475
consensus	*****!***!*****!***!*****!*****!*****!*****!*****!*****!*****!				

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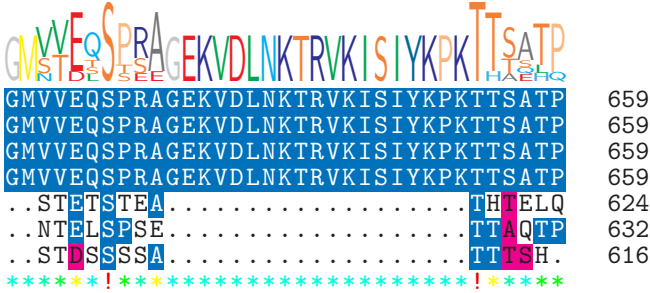
NTDB id 146 SP RS08570 WP 000614538.1	EAGTVLKQSLPEGTTYDLSKA	TQIVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE	EES	544
NTDB id 216 SPD RS08205 WP 000614552.1	EAGTVLKQSLPEGTTYDLSKA	TQIVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE	EES	544
NTDB id 182 SPR RS07820 WP 000614552.1	EAGTVLKQSLPEGTTYDLSKA	TQIVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE	EES	544
NTDB id 257 KZH43 RS07655 WP 220041236.1	EAGTVLKQSLPEGTTYDLSKA	TQIVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE	EES	544
NTDB id 467 HSISS4 RS06915 WP 021143821.1	EEGEILSQSPGKNKSFNPKDKAKIKFRVATP	KITMPDVTGLTVSTAVQTLNRKN	SSSSIEYHDYNTGAK..LDKAK	553
NTDB id 1138061 DQL37 RS07040 WP 063812253.1	PENTVTSQSPSAGDKFNPNG.KSKITLS	VAVS.DTITMPMVTEYSYADAVNTLTALG	DASRIKAYVPSSSSATGFVPIH	556
NTDB id 384 SMU RS02325 WP 002263039.1	SGGTVIGQSPKPKGKTYHPS	SS.DKKITLKV...KVITMPNLKNSTYEA	VSTLTAMGSSSRKAYDASDYS...SEIS	546
consensus	*****!***!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!			

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NTDB id 146 SP RS08570 WP 000614538.1	SESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNNL	LIQIVGIKEANIEVVEVTTAPAGSVE	624
NTDB id 216 SPD RS08205 WP 000614552.1	SESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNNL	LIQIVGIKEANIEVVEVTTAPAGSVE	624
NTDB id 182 SPR RS07820 WP 000614552.1	SESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNNL	LIQIVGIKEANIEVVEVTTAPAGSVE	624
NTDB id 257 KZH43 RS07655 WP 220041236.1	SESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNNL	LIQIVGIKEANIEVVEVTTAPAGSVE	624
NTDB id 467 HSISS4 RS06915 WP 021143821.1	IPSSSTEI.LYQDPQAC...	TSVDGTVLYSVATASSLQSSSSST...H.SSS.....TSSSTDS	TTS...610
NTDB id 1138061 DQL37 RS07040 WP 063812253.1	SPSSKAIYSGSPYYCTLSLSDKGEISLYLYPEETHSSSSSSST	SS...SNSSS.....TNSTAPCS...	618
NTDB id 384 SMU RS02325 WP 002263039.1	SPSSSSLVVGQSPYYCNTVLS	SSNDITLYVSTSGGSHGSSSSESSN.....SEG.....TTSSE.A.....	603
consensus	***!*****!***!*****!*****!*****!*****!*****!*****!*****!*****!*****!		

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NTDB id 384 SMU RS02325 WP 002263039.1
consensus



X non conserved
X similar
X ≥ 50% conserved