

logo

NTDB id 1141990 DQL12 RS08895 WP 000614538.1
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 384 SMU RS02325 WP 002263039.1
NTDB id 467 HSISS4 RS06915 WP 021143821.1
consensus



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NTDB id 1141990 DQL12 RS08895 WP 000614538.1
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NTDB id 1141990 DQL12 RS08895 WP 000614538.1
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NTDB id 1141990 DQL12 RS08895 WP 000614538.1	KNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVL	VAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEK	398
NTDB id 146 SP RS08570 WP 000614538.1	KNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVL	VAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEK	398
NTDB id 216 SPD RS08205 WP 000614552.1	KNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVL	VAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEK	398
NTDB id 182 SPR RS07820 WP 000614552.1	KNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVL	VAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEK	398
NTDB id 257 KZH43 RS07655 WP 220041236.1	KNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVL	VAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEK	398
NTDB id 384 SMU RS02325 WP 002263039.1	KVDHKSQPKTKPQPKPKKKRHLARHFKIFFILVAIGIAVLT	TVLVLTKEPSTVSVPDVSGDKLSTAKMIIRKSGLVKGVQ	398
NTDB id 467 HSISS4 RS06915 WP 021143821.1	PAPSKQPRKKTPAKKKKKKNEFFSTLLKVFGLVFIGIIFAYLV	FTNPDNVQVPMVVGQELSTAQTKLEGAGEKVGEVK	398
consensus	*****!!!!!!!!!!!!*****!!!!!!!!!!!!*****!!!!!!!!!!!!*****!!!!!!!!!!!!*****		

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NTDB id 1141990 DQL12 RS08895 WP 000614538.1	TEASEKVEEGRIIRTDPGAGTGRKEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK	KVPDNLKIEEEESNESEA	477
NTDB id 146 SP RS08570 WP 000614538.1	TEASEKVEEGRIIRTDPGAGTGRKEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK	KVPDNLKIEEEESNESEA	477
NTDB id 216 SPD RS08205 WP 000614552.1	TEASEKVEEGRIIRTDPGAGTGRKEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK	KVPDNLKIEEEESNESEA	477
NTDB id 182 SPR RS07820 WP 000614552.1	TEASEKVEEGRIIRTDPGAGTGRKEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK	KVPDNLKIEEEESNESEA	477
NTDB id 257 KZH43 RS07655 WP 220041236.1	TEASEKVEEGRIIRTDPGAGTGRKEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK	KVPDNLKIEEEESNESEA	477
NTDB id 384 SMU RS02325 WP 002263039.1	KIEDINVGACKVVRTPSAGSKKRECSVDIFVAS	AKGFKMEDYTGQDYKDAIDNLTNNYGVSRDSIVLKVSSDDYSG	477
NTDB id 467 HSISS4 RS06915 WP 021143821.1	EVEDLSVDTGKVIKTDPTAGTTRKEGSSIDVYVSSGSKGFALKDYKCKNYKDAIED	LTSNFCVSEIQIDTGHVEDLSAE	478
consensus	*****!!!!!!!!!!!!*****!!!!!!!!!!!!*****!!!!!!!!!!!!*****!!!!!!!!!!!!*****		

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NTDB id 1141990 DQL12 RS08895 WP 000614538.1	GTVLKQSLPEGTTYDLSKATQ	IVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE		EESSESE	548
NTDB id 146 SP RS08570 WP 000614538.1	GTVLKQSLPEGTTYDLSKATQ	IVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE		EESSESE	548
NTDB id 216 SPD RS08205 WP 000614552.1	GTVLKQSLPEGTTYDLSKATQ	IVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE		EESSESE	548
NTDB id 182 SPR RS07820 WP 000614552.1	GTVLKQSLPEGTTYDLSKATQ	IVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE		EESSESE	548
NTDB id 257 KZH43 RS07655 WP 220041236.1	GTVLKQSLPEGTTYDLSKATQ	IVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE		EESSESE	548
NTDB id 384 SMU RS02325 WP 002263039.1	GTVIGQSPKPKCKTYHPSSDKK	ITLKV...VKVTMPNLNKNTYEBAYSTLTAMG	SSSRIRKAY..DASDYSS	ETSSPSS	550
NTDB id 467 HSISS4 RS06915 WP 021143821.1	GEILSQSPGKNKSFNPKDKAK	IKFRVAT.PKIVTMPDVTGLTVSTAVQTINRN	SSSSIEYH	YNTGAKLDKAKIPSS	557
consensus	*****!!!!!!!!!!!!*****!!!!!!!!!!!!*****!!!!!!!!!!!!*****!!!!!!!!!!!!*****		*****!!!!!!!!!!!!*****		

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NTDB id 1141990 DQL12 RS08895 WP 000614538.1	PGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNLIQIVGIKEANIEVVEVTTAPAGS	AEGMVV	628
NTDB id 146 SP RS08570 WP 000614538.1	PGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNLIQIVGIKEANIEVVEVTTAPAGS	AEGMVV	628
NTDB id 216 SPD RS08205 WP 000614552.1	PGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNLIQIVGIKEANIEVVEVTTAPAGS	VEGMVV	628
NTDB id 182 SPR RS07820 WP 000614552.1	PGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNLIQIVGIKEANIEVVEVTTAPAGS	VEGMVV	628
NTDB id 257 KZH43 RS07655 WP 220041236.1	PGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNLIQIVGIKEANIEVVEVTTAPAGS	VEGMVV	628
NTDB id 384 SMU RS02325 WP 002263039.1	SSLVVGQSPYYCNIIVSLSSNDDIDILYVSTSGGSHSGSSSESSNS...E.....	CTTS.....SEASTDSSSATT...	614
NTDB id 467 HSISS4 RS06915 WP 021143821.1	TE..ILYQDPQ...AGTSVDGTVILYVSVATASSSLQSSSSSTH...S.....SST.....SSSTSTTS	STETS..	615
consensus	*****!!!!!!!!!!!!*****!!!!!!!!!!!!*****!!!!!!!!!!!!*****!!!!!!!!!!!!*****		

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NTDB id 1141990 DQL12 RS08895 WP 000614538.1	EQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 146 SP RS08570 WP 000614538.1	EQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 216 SPD RS08205 WP 000614552.1	EQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 182 SPR RS07820 WP 000614552.1	EQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 257 KZH43 RS07655 WP 220041236.1	EQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 384 SMU RS02325 WP 002263039.1	SH.....	616
NTDB id 467 HSISS4 RS06915 WP 021143821.1	TEA [~] TH [~] EL [~] Q.....	624
consensus	*****~*****	

- [~] non conserved
- [~] similar
- [~] ≥ 50% conserved