

logo

NTDB id 30626 SPCG RS08825 WP 000614540.1
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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consensus



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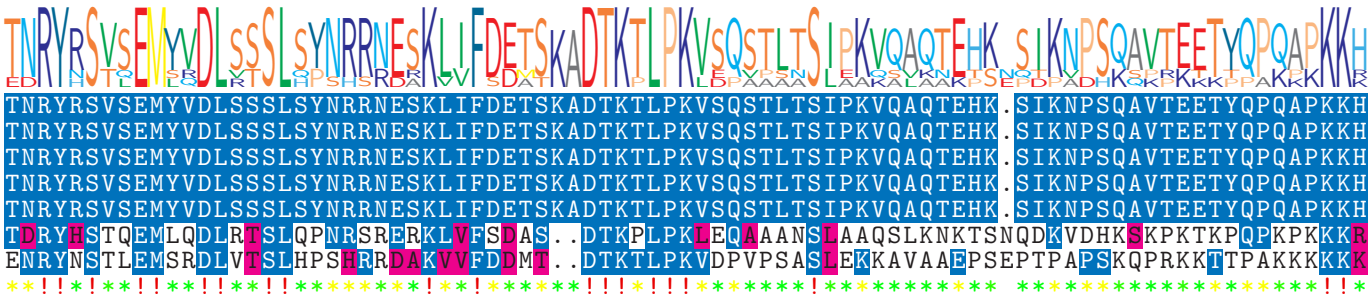
NTDB id 30626 SPCG RS08825 WP 000614540.1
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NTDB id 30626 SPCG RS08825 WP 000614540.1	RFKMRYLILLASLVLVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTPGAGTGRKE	423
NTDB id 146 SP RS08570 WP 000614538.1	RFKMRYLILLASLVLVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTPGAGTGRKE	423
NTDB id 216 SPD RS08205 WP 000614552.1	RFKMRYLILLASLVLVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTPGAGTGRKE	423
NTDB id 182 SPR RS07820 WP 000614552.1	RFKMRYLILLASLVLVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTPGAGTGRKE	423
NTDB id 257 KZH43 RS07655 WP 220041236.1	RFKMRYLILLASLVLVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTPGAGTGRKE	423
NTDB id 384 SMU RS02325 WP 002263039.1	FLARFKIFFILVAIGTAVLTVLVLTKESTVSVDPVSGDKLSTAKMIIRKSGLKVGVDVQKIEDNVGAGKVVRTNPSAGSKKRE	423
NTDB id 467 HSISS4 RS06915 WP 021143821.1	NEFSTLLKVLGLVFIGIIFAYLVFTNPDNMQPNVVGQELSTAQTKLEGAGFKVGEVKEVEDDSVDTGKVIKTDPTAGTTRKE	423
consensus	*!*****!!* *****!!*****!*!*!*****!*****!!*****!!*!*!*!*!*!*!*!*!*!*!*!	

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NTDB id 30626 SPCG RS08825 WP 000614540.1	GTKINLVSSGKQSFQISNYVGRKSSDVIAELKEKKEKKEKVPDNLKIEEEESNESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKK	506
NTDB id 146 SP RS08570 WP 000614538.1	GTKINLVSSGKQSFQISNYVGRKSSDVIAELKEKKEKKEKVPDNLKIEEEESNESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKK	506
NTDB id 216 SPD RS08205 WP 000614552.1	GTKINLVSSGKQSFQISNYVGRKSSDVIAELKEKKEKKEKVPDNLKIEEEESNESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKK	506
NTDB id 182 SPR RS07820 WP 000614552.1	GTKINLVSSGKQSFQISNYVGRKSSDVIAELKEKKEKKEKVPDNLKIEEEESNESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKK	506
NTDB id 257 KZH43 RS07655 WP 220041236.1	GTKINLVSSGKQSFQISNYVGRKSSDVIAELKEKKEKKEKVPDNLKIEEEESNESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKK	506
NTDB id 384 SMU RS02325 WP 002263039.1	CSSVDIFVASLAKGFKMEDYTGQDYKDAIDNLTNNYGVSRDSIVLKEVSSDDYSGGTVIGQSPKPGKTYHPSSDKK.ITLKV...	503
NTDB id 467 HSISS4 RS06915 WP 021143821.1	CSSIDVYVSSGSKCEALKDYKCKNYKDAIEDLTSNFVGSEQIDITQHVEDSAEEGETLSQSPGKNKSFNPKDSKAKIKFRVAT.	507
consensus	!*****!*!******!*****!*!******!*!*!*****!*!******!*****!!*****!!*****!!*****!!*	

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NTDB id 30626 SPCG RS08825 WP 000614540.1	ATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE.....EESSESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSA	583
NTDB id 146 SP RS08570 WP 000614538.1	ATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE.....EESSESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSA	583
NTDB id 216 SPD RS08205 WP 000614552.1	ATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE.....EESSESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSA	583
NTDB id 182 SPR RS07820 WP 000614552.1	ATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE.....EESSESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSA	583
NTDB id 257 KZH43 RS07655 WP 220041236.1	ATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE.....EESSESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSA	583
NTDB id 384 SMU RS02325 WP 002263039.1	.VKVTMPNLKNSTYEAIVSTLTAMGSSSRKAY..DASDYSSSTSSSSSLVVGQSPYYGNVTSSSNDDITLYVSTSGGSHS	585
NTDB id 467 HSISS4 RS06915 WP 021143821.1	PKIITMPDVTGLTVSTAVQTINRNKSSSLEYHGYNTGAKLKDCAKIPSSSTEILYQDPQ...AGTSDVGTIVLYVSVATASSS	587
consensus	*****!!*****!!*****!!*****!!*****!!*****!!*****!!*****!!*****!!*****!!*****!!*	

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NTDB id 30626 SPCG RS08825 WP 000614540.1	MPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTAPAGSVEGMVV.....EQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 146 SP RS08570 WP 000614538.1	MPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTAPAGSAEGMVV.....EQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 216 SPD RS08205 WP 000614552.1	MPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTAPAGSVEGMVV.....EQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 182 SPR RS07820 WP 000614552.1	MPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTAPAGSVEGMVV.....EQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 257 KZH43 RS07655 WP 220041236.1	MPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTAPAGSVEGMVV.....EQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 384 SMU RS02325 WP 002263039.1	GSSSSESSNS.....EGTSSSEASTSSSSATTI...SH.....	616
NTDB id 467 HSISS4 RS06915 WP 021143821.1	LQSSSSSTH.....SSST..SSSTSTTSSTETSTEATHTELQ.....	624
consensus	*!*!*!*!******!!*****!!*****!!*****!!*****!!*****!!*****!!*****!!*****!!*****!!*	

- X non conserved
- X similar
- X ≥ 50% conserved