

logo

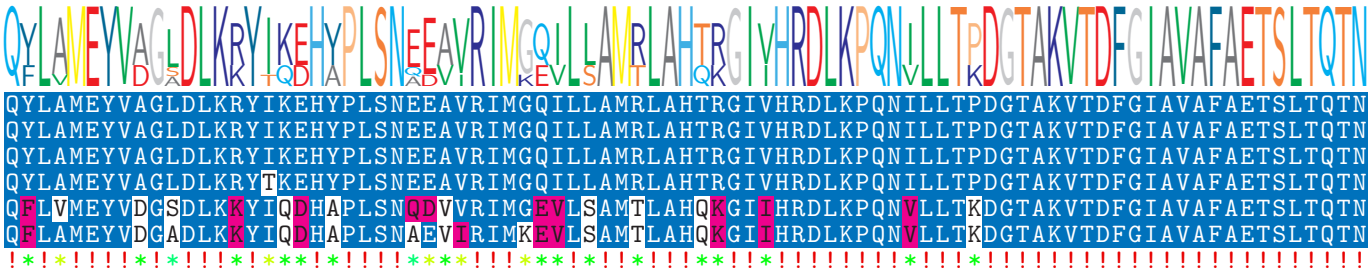
NTDB id 146 SP RS08570 WP 000614538.1
NTDB id 182 SPR RS07820 WP 000614552.1
NTDB id 216 SPD RS08205 WP 000614552.1
NTDB id 257 KZH43 RS07655 WP 220041236.1
NTDB id 467 HSISS4 RS06915 WP 021143821.1
NTDB id 384 SMU RS02325 WP 002263039.1
consensus



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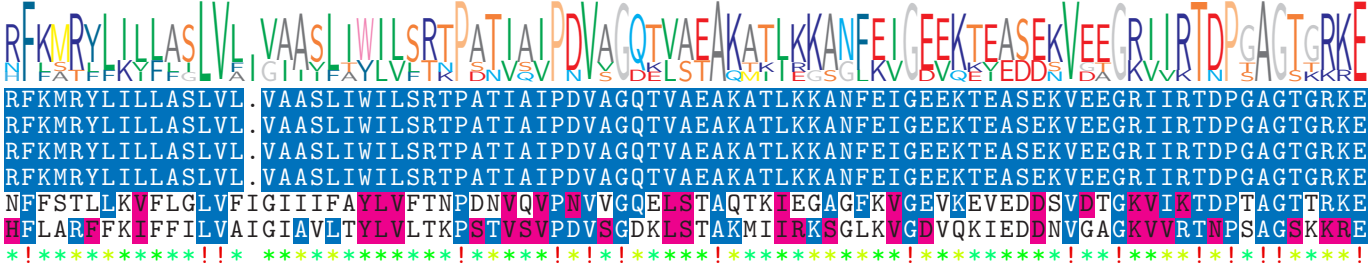
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NTDB id 146	SP RS08570 WP 000614538.1	GTKINLVVSSGKQSFQISNYVGRKSSDVIAELKEK...KVPDNLKIEEEEEESNESEAGTVLKQSLPEGTTYDLSKATQ...IVLTVAKK	506
NTDB id 182	SPR RS07820 WP 000614552.1	GTKINLVVSSGKQSFQISNYVGRKSSDVIAELKEK...KVPDNLKIEEEEEESNESEAGTVLKQSLPEGTTYDLSKATQ...IVLTVAKK	506
NTDB id 216	SPD RS08205 WP 000614552.1	GTKINLVVSSGKQSFQISNYVGRKSSDVIAELKEK...KVPDNLKIEEEEEESNESEAGTVLKQSLPEGTTYDLSKATQ...IVLTVAKK	506
NTDB id 257	KZH43 RS07655 WP 220041236.1	GTKINLVVSSGKQSFQISNYVGRKSSDVIAELKEK...KVPDNLKIEEEEEESNESEAGTVLKQSLPEGTTYDLSKATQ...IVLTVAKK	506
NTDB id 467	HSISS4 RS06915 WP 021143821.1	SSSIDVYVSSGSKGFALKDYKCKNYKDAIEDLTSNFVGVSQDIDIQHVEDDSAEGETILSQSPGKNKSFNPKDKSAKIKFRVAT.	507
NTDB id 384	SMU RS02325 WP 002263039.1	SSVDIFVAS.AKGFKMEDYTGQDYKDAIDNLTNNYGVSRDSIVLKEVSSDDYS CGTVIGQSPKPKGKTYHPSSDKK.ITLKV...	503
consensus		!*****!*!******!*****!*!******!*!******!*****!*****!*****!*****!*****!*****!*****!*****	

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NTDB id 146	SP RS08570 WP 000614538.1	ATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE.....EESSESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSA	583
NTDB id 182	SPR RS07820 WP 000614552.1	ATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE.....EESSESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSA	583
NTDB id 216	SPD RS08205 WP 000614552.1	ATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE.....EESSESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSA	583
NTDB id 257	KZH43 RS07655 WP 220041236.1	ATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEE.....EESSESEPGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSA	583
NTDB id 467	HSISS4 RS06915 WP 021143821.1	PKIVTMPDVTCLTVSTAVQTLNRKNSSSSIEYHDYNTGAKLDKAKIPSSSTE.ILYQDPQ...AGTSVDGTVILYVSVATASSS	587
NTDB id 384	SMU RS02325 WP 002263039.1	.VKVTMPNLKNSTYEEAVSTLTAMGISSSRIKAY..DASDYSS EISSPSSSSLVGQSPYYGNTVSLSSNDDIILYVSTSGGSHS	585
consensus		*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****	

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NTDB id 146	SP RS08570 WP 000614538.1	MPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTAPAGSVEGMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 182	SPR RS07820 WP 000614552.1	MPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTAPAGSVEGMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 216	SPD RS08205 WP 000614552.1	MPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTAPAGSVEGMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 257	KZH43 RS07655 WP 220041236.1	MPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTAPAGSVEGMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP	659
NTDB id 467	HSISS4 RS06915 WP 021143821.1	LQSSSSSTH...S.....SST.....SSSTDSTTSSTETS.....TEATHTELQ.....	624
NTDB id 384	SMU RS02325 WP 002263039.1	GSSSSESSNS...E.....GTTS.....SEASTDSSSATTT.....SH.....	616
consensus		**!*!!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****	

-  non conserved
-  similar
-  ≥ 50% conserved