

logo

NTDB id 85758 STYK RS08175 WP 261804861.1
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 85758 STYK RS08175 WP 261804861.1	RFKMRYLILLASLVLV.AASLIWILSRTPATIAIPDVAGQTVAEAKETLKKANFEIGDEKSEASEKVEEGRIIRTDPDAGTGRKE	423
NTDB id 146 SP RS08570 WP 000614538.1	RFKMRYLILLASLVLV.AASLIWILSRTPATIAIPDVAGQTVAEAKETLKKANFEIGDEKSEASEKVEEGRIIRTDPDAGTGRKE	423
NTDB id 216 SPD RS08205 WP 000614552.1	RFKMRYLILLASLVLV.AASLIWILSRTPATIAIPDVAGQTVAEAKETLKKANFEIGDEKSEASEKVEEGRIIRTDPDAGTGRKE	423
NTDB id 182 SPR RS07820 WP 000614552.1	RFKMRYLILLASLVLV.AASLIWILSRTPATIAIPDVAGQTVAEAKETLKKANFEIGDEKSEASEKVEEGRIIRTDPDAGTGRKE	423
NTDB id 257 KZH43 RS07655 WP 220041236.1	RFKMRYLILLASLVLV.AASLIWILSRTPATIAIPDVAGQTVAEAKETLKKANFEIGDEKSEASEKVEEGRIIRTDPDAGTGRKE	423
NTDB id 384 SMU RS02325 WP 002263039.1	RFKMRYLILLASLVLV.AASLIWILSRTPATIAIPDVAGQTVAEAKETLKKANFEIGDEKSEASEKVEEGRIIRTDPDAGTGRKE	423
NTDB id 467 HSISS4 RS06915 WP 021143821.1	RFKMRYLILLASLVLV.AASLIWILSRTPATIAIPDVAGQTVAEAKETLKKANFEIGDEKSEASEKVEEGRIIRTDPDAGTGRKE	423
consensus	RFKMRYLILLASLVLV.AASLIWILSRTPATIAIPDVAGQTVAEAKETLKKANFEIGDEKSEASEKVEEGRIIRTDPDAGTGRKE	

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NTDB id 85758 STYK RS08175 WP 261804861.1	GTKINLVVSSGKQSFQISNYVGRKSSDVIAELKEKQKVPENLIK.....EEEESENESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKKA	466
NTDB id 146 SP RS08570 WP 000614538.1	GTKINLVVSSGKQSFQISNYVGRKSSDVIAELKEKQKVPENLIK.....EEEESENESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKKA	507
NTDB id 216 SPD RS08205 WP 000614552.1	GTKINLVVSSGKQSFQISNYVGRKSSDVIAELKEKQKVPENLIK.....EEEESENESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKKA	507
NTDB id 182 SPR RS07820 WP 000614552.1	GTKINLVVSSGKQSFQISNYVGRKSSDVIAELKEKQKVPENLIK.....EEEESENESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKKA	507
NTDB id 257 KZH43 RS07655 WP 220041236.1	GTKINLVVSSGKQSFQISNYVGRKSSDVIAELKEKQKVPENLIK.....EEEESENESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKKA	507
NTDB id 384 SMU RS02325 WP 002263039.1	GTKINLVVSSGKQSFQISNYVGRKSSDVIAELKEKQKVPENLIK.....EEEESENESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKKA	466
NTDB id 467 HSISS4 RS06915 WP 021143821.1	GTKINLVVSSGKQSFQISNYVGRKSSDVIAELKEKQKVPENLIK.....EEEESENESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKKA	467
consensus	GTKINLVVSSGKQSFQISNYVGRKSSDVIAELKEKQKVPENLIK.....EEEESENESEAGTVLKQSLPEGTTYDLSKATQIVLTVAKKA	

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NTDB id 85758 STYK RS08175 WP 261804861.1	TTIQLGNYIGRNSTEVISELKQKKVPENLIK.....EEEESESEEGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSAVMPSYIGSS	519
NTDB id 146 SP RS08570 WP 000614538.1	TTIQLGNYIGRNSTEVISELKQKKVPENLIK.....EEEESESEEGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSAVMPSYIGSS	591
NTDB id 216 SPD RS08205 WP 000614552.1	TTIQLGNYIGRNSTEVISELKQKKVPENLIK.....EEEESESEEGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSAVMPSYIGSS	591
NTDB id 182 SPR RS07820 WP 000614552.1	TTIQLGNYIGRNSTEVISELKQKKVPENLIK.....EEEESESEEGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSAVMPSYIGSS	591
NTDB id 257 KZH43 RS07655 WP 220041236.1	TTIQLGNYIGRNSTEVISELKQKKVPENLIK.....EEEESESEEGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSAVMPSYIGSS	591
NTDB id 384 SMU RS02325 WP 002263039.1	TTIQLGNYIGRNSTEVISELKQKKVPENLIK.....EEEESESEEGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSAVMPSYIGSS	515
NTDB id 467 HSISS4 RS06915 WP 021143821.1	TTIQLGNYIGRNSTEVISELKQKKVPENLIK.....EEEESESEEGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSAVMPSYIGSS	520
consensus	TTIQLGNYIGRNSTEVISELKQKKVPENLIK.....EEEESESEEGTIMKQSPGAGTTYDVSKPTQIVLTVAKKVTSAVMPSYIGSS	

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NTDB id 85758 STYK RS08175 WP 261804861.1	LEFTKNNLIQIVGIKEANIEVVEVT.....TAPAGSVEGMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP.....SSAPAGRGNQ	597
NTDB id 146 SP RS08570 WP 000614538.1	LEFTKNNLIQIVGIKEANIEVVEVT.....TAPAGSVEGMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP.....SSAPAGRGNQ	659
NTDB id 216 SPD RS08205 WP 000614552.1	LEFTKNNLIQIVGIKEANIEVVEVT.....TAPAGSVEGMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP.....SSAPAGRGNQ	659
NTDB id 182 SPR RS07820 WP 000614552.1	LEFTKNNLIQIVGIKEANIEVVEVT.....TAPAGSVEGMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP.....SSAPAGRGNQ	659
NTDB id 257 KZH43 RS07655 WP 220041236.1	LEFTKNNLIQIVGIKEANIEVVEVT.....TAPAGSVEGMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP.....SSAPAGRGNQ	659
NTDB id 384 SMU RS02325 WP 002263039.1	LEFTKNNLIQIVGIKEANIEVVEVT.....TAPAGSVEGMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP.....SSAPAGRGNQ	596
NTDB id 467 HSISS4 RS06915 WP 021143821.1	LEFTKNNLIQIVGIKEANIEVVEVT.....TAPAGSVEGMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP.....SSAPAGRGNQ	598
consensus	LEFTKNNLIQIVGIKEANIEVVEVT.....TAPAGSVEGMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP.....SSAPAGRGNQ	

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NTDB id 85758 STYK RS08175 WP 261804861.1	GSTTTTPSQGNQQGNQRGDTHSTNQSSSEGSHE NSRN	633
NTDB id 146 SP RS08570 WP 000614538.1		659
NTDB id 216 SPD RS08205 WP 000614552.1		659
NTDB id 182 SPR RS07820 WP 000614552.1		659
NTDB id 257 KZH43 RS07655 WP 220041236.1		659
NTDB id 384 SMU RS02325 WP 002263039.1	GTTSSSEAS.....TDSSSSATTT....SH.....	616
NTDB id 467 HSISS4 RS06915 WP 021143821.1	SST..SSS.....TDSTTSSTETSTEATHTELQ.	624
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved