

logo

NTDB id 362 SMU RS05200 WP 002262214.1	..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE	82
NTDB id 42897 ST01 RS05845 WP 000590630.1	..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE	82
NTDB id 258 KZH43 RS03475 WP 000590640.1	..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE	82
NTDB id 217 SPD RS03775 WP 000590640.1	..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE	82
NTDB id 183 SPR RS03560 WP 000590640.1	..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE	82
NTDB id 147 SP RS03905 WP 000590640.1	..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDLMLPEKNGFQVLKELREKGITTPVLIMTAKE	82
NTDB id 606 V4T04 RS01910 WP 003130756.1	MTSKKILITEDKNLARFVSLLELHGYATEIKDNGRSGLEEATSKDYDLILLDLMLPELDGFEVARRLRKE.KDTPITIMMTARD	84
NTDB id 374 SMU RS06885 WP 002262930.1	..MKILLVDDKPTSDIKFNLAKGYDITITAFDGRREALSKYEENPDILILLDLMLPELDGLEVAKEVRKN.SHVPITIMLSAKD	82
consensus	***!***!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!	

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NTDB id 362 SMU RS05200 WP 002262214.1	SLDDKKGHGFEELGADDYLTkPFYleELKMRlQALLKRSGKFN.....ENN.....LTYGELTVDTATNTTMVNGKEVELLGKEFDL	157
NTDB id 42897 ST01 RS05845 WP 000590630.1	SLDDKKGHGFEELGADDYLTkPFYleELKMRlQALLKRSGKFN.....ENT.....LTYGDIVVNLSTNEVKVEDTPVELLGKEFEL	157
NTDB id 258 KZH43 RS03475 WP 000590640.1	SLDDKKGHGFEELGADDYLTkPFYleELKMRlQALLKRSGKFN.....ENT.....LTYGNIVVNLSTNTVKVEDTPVELLGKEFDL	157
NTDB id 217 SPD RS03775 WP 000590640.1	SLDDKKGHGFEELGADDYLTkPFYleELKMRlQALLKRSGKFN.....ENT.....LTYGNIVVNLSTNTVKVEDTPVELLGKEFDL	157
NTDB id 183 SPR RS03560 WP 000590640.1	SLDDKKGHGFEELGADDYLTkPFYleELKMRlQALLKRSGKFN.....ENT.....LTYGNIVVNLSTNTVKVEDTPVELLGKEFDL	157
NTDB id 147 SP RS03905 WP 000590640.1	SLDDKKGHGFEELGADDYLTkPFYleELKMRlQALLKRSGKFN.....ENT.....LTYGNIVVNLSTNTVKVEDTPVELLGKEFDL	157
NTDB id 606 V4T04 RS01910 WP 003130756.1	STMDRVAGLDIGADDYITKPFATEELARVRAFFRREEGHAVERAENT.SFRDLVI.....DKTNRTVHRGKKVIDLTRREYDL	163
NTDB id 374 SMU RS06885 WP 002262930.1	SEFDKVICLETIGADDYVTKPFNSRELARVKAHRTTENIESAVAEENASGIPEITIGDQILPDAFVAKKRGTLEVELTHREFEL	167
consensus	***!***!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!	

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NTDB id 362 SMU RS05200 WP 002262214.1	LVYFLQNNVILPKSQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGT.TFCKNLQTLRSVGYLKDDAN	224
NTDB id 42897 ST01 RS05845 WP 000590630.1	LVYFLQNNVILPKSQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGT.AFAENLQTLRSVGYLKDDVQ	224
NTDB id 258 KZH43 RS03475 WP 000590640.1	LVYFLQNNVILPKSQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGT.TFAENLQTLRSVGYLKDDVQ	224
NTDB id 217 SPD RS03775 WP 000590640.1	LVYFLQNNVILPKSQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGT.TFAENLQTLRSVGYLKDDVQ	224
NTDB id 183 SPR RS03560 WP 000590640.1	LVYFLQNNVILPKSQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGT.TFAENLQTLRSVGYLKDDVQ	224
NTDB id 147 SP RS03905 WP 000590640.1	LVYFLQNNVILPKSQIFDRLWGFDSDTTISVVEVYVSKVRKKLKGT.TFAENLQTLRSVGYLKDDVQ	224
NTDB id 606 V4T04 RS01910 WP 003130756.1	LTLMQNVGDVVTREHVSQVWGVEEGTETNVVDVYTRYLRNKIDV.EGQDSYIQTVRGLGYVMRERK	230
NTDB id 374 SMU RS06885 WP 002262930.1	LHHLATHGTQVMTREHLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYVMKSYD	235
consensus	!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!	

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