

logo

NTDB id 362 SMU RS05200 WP 002262214.1	..MKKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDDLMLPEKNGFQVLKELREKGITTPVLIIMTAK	82
NTDB id 258 KZH43 RS03475 WP 000590640.1	..MKKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDDLMLPEKNGFQVLKELREKGITTPVLIIMTAK	82
NTDB id 217 SPD RS03775 WP 000590640.1	..MKKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDDLMLPEKNGFQVLKELREKGITTPVLIIMTAK	82
NTDB id 183 SPR RS03560 WP 000590640.1	..MKKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDDLMLPEKNGFQVLKELREKGITTPVLIIMTAK	82
NTDB id 147 SP RS03905 WP 000590640.1	..MKKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEEGLYEAESGVYDLILLDDLMLPEKNGFQVLKELREKGITTPVLIIMTAK	82
NTDB id 374 SMU RS06885 WP 002262930.1	..MKKILLVDDDKPISDIKFNLAKEGYDITITAFDGREALSKYEENPDLIILDDLMLPELDGLEVAKEVRKN.SHVPIIMLSAKD	82
NTDB id 413 AAK55817.1 141..845()	..MKKILLVDDDKPISDIKFNMTKEGYEVVTAENGREALEQFEAEQPDIIILDDLMLPEIDGLEVAKTIRKT.SSVPIIMLSAKD	82
NTDB id 26562 LACR RS08195 WP 011676631.1	MASKKILIIIEDEKNLARFVSLLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDDLMLPELDGFEVARRLRKE.KDTPIMMTARD	84
NTDB id 606 V4T04 RS01910 WP 003130756.1	MTSKKILIIIEDEKNLARFVSLLELEHEGYATEIKDNGRSGLEEATSKDYDLILLDDLMLPELDGFEVARRLRKE.KDTPIMMTARD	84
NTDB id 471 HSISS4 RS01430 WP 002883757.1	.MSKRILIVDEEKNLARFVSLLELQHEGYDVVTADNGREGLEMALEKDFDLILLDDLMLPEMDGFEVTRRLQKE.KDTYIMMTARD	83
consensus	* **!* **!* ***** **** **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!	

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NTDB id 362 SMU RS05200 WP 002262214.1	GLDDDKGHGFELGADDYLTTPFYLEELKMRIQALLKRSQKFNEN.....NLTYGELTVDATNTTMVNGKEVELLGKEFDL	157
NTDB id 258 KZH43 RS03475 WP 000590640.1	SLDDDKGHGFELGADDYLTTPFYLEELKMRIQALLKRSQKFNEN.....TLTYGNIVVNLSTNTVKVEDTPVELLGKEFDL	157
NTDB id 217 SPD RS03775 WP 000590640.1	SLDDDKGHGFELGADDYLTTPFYLEELKMRIQALLKRSQKFNEN.....TLTYGNIVVNLSTNTVKVEDTPVELLGKEFDL	157
NTDB id 183 SPR RS03560 WP 000590640.1	SLDDDKGHGFELGADDYLTTPFYLEELKMRIQALLKRSQKFNEN.....TLTYGNIVVNLSTNTVKVEDTPVELLGKEFDL	157
NTDB id 147 SP RS03905 WP 000590640.1	SLDDDKGHGFELGADDYLTTPFYLEELKMRIQALLKRSQKFNEN.....TLTYGNIVVNLSTNTVKVEDTPVELLGKEFDL	157
NTDB id 374 SMU RS06885 WP 002262930.1	SEFDKVICLELGADDYVTKPFNSRELARVKAHLRRTENIESAVAEENASGIPEIIGDLQILPDAFVAKKRGTEVELTHREFEL	167
NTDB id 413 AAK55817.1 141..845()	SEFDKVICLELGADDYVTKPFNSRELQARVKAHLRRSQMPVVGQ.EADSKPQPIQIGDLIVPDAYVAKKYGEELDLTHREFEL	166
NTDB id 26562 LACR RS08195 WP 011676631.1	STMDRVAGLDIGADDYITKPFATEELARVRAFFRREEHGHAV.ERAENTS....FRDLVIDKTNRVHRGKKVIDLTRREYDL	163
NTDB id 606 V4T04 RS01910 WP 003130756.1	STMDRVAGLDIGADDYITKPFATEELARVRAFFRREEHGHAV.ERAENTS....FRDLVIDKTNRVHRGKKVIDLTRREYDL	163
NTDB id 471 HSISS4 RS01430 WP 002883757.1	STMDRVAGLDIGADDYITKPFATEELARIRATFRQDIEAAKNAPAKAST....YRDLKLDVQNRVTVRGDEAIPLTKEFDL	163
consensus	***!* *! **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!	

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NTDB id 362 SMU RS05200 WP 002262214.1	LVYFLQNNQNVLPKTSQIFDRWGFSDTTISVVEVYVSKVRKKLKDTT.FCKNLQTLRSVGYLLKDVQ.	224
NTDB id 258 KZH43 RS03475 WP 000590640.1	LVYFLQNNQNVLPKTSQIFDRWGFSDTTISVVEVYVSKVRKKLKDTT.FAENLQTLRSVGYLLKDVQ.	224
NTDB id 217 SPD RS03775 WP 000590640.1	LVYFLQNNQNVLPKTSQIFDRWGFSDTTISVVEVYVSKVRKKLKDTT.FAENLQTLRSVGYLLKDVQ.	224
NTDB id 183 SPR RS03560 WP 000590640.1	LVYFLQNNQNVLPKTSQIFDRWGFSDTTISVVEVYVSKVRKKLKDTT.FAENLQTLRSVGYLLKDVQ.	224
NTDB id 147 SP RS03905 WP 000590640.1	LVYFLQNNQNVLPKTSQIFDRWGFSDTTISVVEVYVSKVRKKLKDTT.FAENLQTLRSVGYLLKDVQ.	224
NTDB id 374 SMU RS06885 WP 002262930.1	LHHLATHTGQVMTREHLLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYVMKSYD.	235
NTDB id 413 AAK55817.1 141..845()	LYHLASHTGQVMTREHLLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYVMRINA*	234
NTDB id 26562 LACR RS08195 WP 011676631.1	LLTLMQNQVGDVVTREHLLVSQVWGYEEGTETNVVDVYTRYLRNKI.DVEGQDSYIQTVRGLGYVMRERK.	230
NTDB id 606 V4T04 RS01910 WP 003130756.1	LLTLMQNQVGDVVTREHLLVSQVWGYEEGTETNVVDVYTRYLRNKI.DVEGQDSYIQTVRGLGYVMRERK.	230
NTDB id 471 HSISS4 RS01430 WP 002883757.1	LNTLLSNMNMQVMTREHLLQVWVYDDAETNVVDVYTRYLRNKI.DVPGKESYIQTVRGMGYVIREK..	229
consensus	! **! ***** **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!* **!	

-  non conserved
-  similar
-  ≥ 50% conserved