

logo

NTDB id 1419 SGL RS12550 WP 010873252.1	MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIVFDLQPLINKWMQSENIATLTEIRRHFTLKGSGRMIGAKSSA	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIVFDLQPLINKWMQSENIATLTEIRRHFTLKGSGRMIGAKSSA	80
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus	0

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NTDB id 1419 SGL RS12550 WP 010873252.1	ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFNFKQKRAHTLDFRPLILLGQQQLQQQSLEPALEELLQLS	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFNFKQKRAHTLDFRPLILLGQQQLQQQSLEPALEELLQLS	160
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus	0

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NTDB id 1419 SGL RS12550 WP 010873252.1	HTLMAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETTLTFMEEAEEHLATIHQFLDQELHQYDSYNALIRALHTLR	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	HTLMAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETTLTFMEEAEEHLATIHQFLDQELHQYDSYNALIRALHTLR	240
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus	0

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NTDB id 1419 SGL RS12550 WP 010873252.1	GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQEYREFVDRSLELLSRYSSSEQLLEARLLQFNQSWDAYVEQ	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQEYREFVDRSLELLSRYSSSEQLLEARLLQFNQSWDAYVEQ	320
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus	0

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NTDB id 1419 SGL RS12550 WP 010873252.1	HGDRTDPLMP SHGLVSQ LQLLDVSE LLD AELDFEKG I RNEFPDYLERLSEQAD LLLQH THSQAMLGLHE YTSQLKESYEV	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	HGDRTDPLMP SHGLVSQ LQLLDVSE LLD AELDFEKG I RNEFPDYLERLSEQAD LLLQH THSQAMLGLHE YTSQLKESYEV	400
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	LLDKPALLQSDY I FE IYQKAHQQL IQLFDAL AAGQRVGVVKQHQ S ILEELKLYTQYTPD I SNDLPQQDSTS FEP IYNT E ^{MTS}	3
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	LLDKPALLQSDY I FE IYQKAHQQL IQLFDAL AAGQRVGVVKQHQ S ILEELKLYTQYTPD I SNDLPQQDSTS FEP IYNT IE ^P	480
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	^{EPN} EWATV ENFSDSADWAVLGQSVQQDRQY I SSTQVNRNFDADLLD I FLEEAEELLE G ^{PRIRINNYE EENEDALHNSLY}	28
NTDB id 413 AAK55817.1 141..845()	DPN..... PAQAFENWFEEEEGDDAFTSLI	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	EPEVVATVENFSDSADWAVLGQSVQQDRQY I SSTQVNRNFDADLLD I FLEEAEELLE GIDTDLNIWVGEQENFAALNNLM	560
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	^{RYL} EHTLKGGANMVQATYLGL I AHELES IYERL I QKQLVVTSDL I DF I RLVQDDLADRLQ I MREQQLDYAAPYT I NALKRA	31
NTDB id 413 AAK55817.1 141..845()	AGE.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	RYLHTLKGGANMVQATYLGL I AHELES IYERL I QKQLVVTSDL I DF I RLVQDDLADRLQ I MREQQLDYAAPYT I NALKRA	640
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	ETNSGIEELDFGDLFEQEPTVIQGEFGS.....PSSGEEDPPVRMTLDFNSPEAELD	84
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	GQNSNFQPLPVVEAFDAESEVFSEQKVISEIIIDEIPVELEPALAELETHHDQVFDTAVTELATPVEMITTVTSQENEVA	720
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1ALFTDEDAPTTAIDPFESSSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEQLDQLLVGVDVEP	153
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	ANEQDIEAVVEQTFLEEATELLEMAESLLKQWFQRTNR.....S.....ILL.....	763
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	AITNPGHLSFAGADRVPTLTNEVLL.FA.....AIDDLGLINAPVLDEEFSF.....TELETLIDTPAPKS	214
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	.QLQRAVHSLKGGARMVGLAVQAIAYQLENAFEQFALHHFNSNIYDHLLESIAIWLKDAIFNHNHYQHFDGLQQSLENIQ	842
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	FATISHPFRLPVAMEEDEDKLEALLEQADSSLGGGIAGMGGARLPARP...RQVKRSRFEQTMRVPIKQLDNLSNLI	291
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	F..FETTIQIPTKLTRA.DLFSS.EP.....VMTFIQGDGTEPPPMGAWEQTERLDQNNEMIRVSADLIEKMIDLS	910
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	GELVVKRNRLREQDQERLRQFLDNLNQAQNLSDVGGRMQDLYERTLLEGALLASRSTNSRHHSSKTTYPDARITASSSST	371
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	GENSINRSRIEMDLSQFSHTLVEMELAIQRLADQLRRMEGELETQI.....IAKHGIEH	964
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	MNDQPLDALEMDRFTGFHLLSQEMIELIVRVRESASDIQYLVDETDQVARTLRQVTTQLQEGMTKSRMVPFTQTADRLPR	451
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	SRYTDFDPLEMDQYSSLNQLSKSLAESASDLVDFKNTLSDKIRDTESELLLQSQSRIQAEIQEGLMRTRLVPPSRLLPRLQR	1044
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus		

logo

		
NTDB id 1419 SGL RS12550 WP 010873252.1	AIREVSMKLNKKVNLEVEGGDVLIDKMLEHLYNPMTHLVNNAITHGIETPAERQKLGKSPGHLAVRAFLQGNQTLITV	531
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	IVRQTSTALNRPALFVNNTEGELDRNILERLVTPLEHMLRNAIDHGLEDRAQRQQANKPETGRIELNIQRQGTDDVVVVV	1124
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	SDDGAGINTEMVKQKAVEKGLISRMEAQSLSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVVRTSLIDIRGAINIESTV	611
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	SDDGQGIDVEKVRQKALLAGLIKP..EQDLEQQDILQLIFHPGLSTAEQVTQISGRGVGLDVVQSDIKSLGGHVSVESVY	1202
NTDB id 221299 BMF81 RS00865 WP 006194267.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1
 NTDB id 413 AAK55817.1 141..845()
 NTDB id 374 SMU RS06885 WP 002262930.1
 NTDB id 1046 HON27 RS03130 WP 168726941.1
 NTDB id 221299 BMF81 RS00865 WP 006194267.1
 NTDB id 1417 SGL RS04200 WP 014407071.1
 NTDB id 1047 HON27 RS03110 WP 000389061.1
 consensus

GKGGTTFTIRLPYPTYSYKALMAYRBSBFAIPYRSEYBTRFVSPKSTYQDFESEQELKYEYEREALLLRYPLRRLSEFVNS
 GKGGTTFTIRLPLTLSICKALCCLSDRARIAPFMDGVEDTRDYLPKDIVTD.EDGQKCVKYGKEALLLPLYPLKDLLNYNR 690
 0
 0
 GQGTTFTIRVPTTVAVSDALMVKVADQQFAIPLAQIDRIIRVSPASLEQYFESPQELFEYEN.....KRYPLRRLSEFVNG 1277
 0
 0
 0

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NTDB id 1419 SGL RS12550 WP 010873252.1
 NTDB id 413 AAK55817.1 141..845()
 NTDB id 374 SMU RS06885 WP 002262930.1
 NTDB id 1046 HON27 RS03130 WP 168726941.1
 NTDB id 221299 BMF81 RS00865 WP 006194267.1
 NTDB id 1417 SGL RS04200 WP 014407071.1
 NTDB id 1047 HON27 RS03110 WP 000389061.1
 consensus

QLSRGSSVYGGKQEEVDVVRVVIARGSSGNLLALQIDQVLGEE.EIVIKQIEGPIPKPPGIAGATVVRGDGSIMPIADVLELI 769
 0
 0
 NQPIPRNLG...VMYSLPVLMIKANNGQTVALLDQLIGSRAQIVVKPIGQQFSSIGAIAGATILGDGQVCLILDGQNI 1354
 0
 0
 0

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NTDB id 1419 SGL RS12550 WP 010873252.1
 NTDB id 413 AAK55817.1 141..845()
 NTDB id 374 SMU RS06885 WP 002262930.1
 NTDB id 1046 HON27 RS03130 WP 168726941.1
 NTDB id 221299 BMF81 RS00865 WP 006194267.1
 NTDB id 1417 SGL RS04200 WP 014407071.1
 NTDB id 1047 HON27 RS03110 WP 000389061.1
 consensus

VLIVDDSIITVRELLSLSFKKAGFRVEQARDGQEAWDQLKSGLPCLDYL 848
 49
 49
 RQIQSTQRHKQLSEAVY.....RQRESDELLIMIVDDSVTVRKVTSRLLERQGYDVVTAKDGVDATEQLENI.KPDL 1427
 49
 49
 55
 * *****! * ***** * * * !*****! *! *!*** * *!***

logo

NTDB id 1419 SGL RS12550 WP 010873252.1
 NTDB id 413 AAK55817.1 141..845()
 NTDB id 374 SMU RS06885 WP 002262930.1
 NTDB id 1046 HON27 RS03130 WP 168726941.1
 NTDB id 221299 BMF81 RS00865 WP 006194267.1
 NTDB id 1417 SGL RS04200 WP 014407071.1
 NTDB id 1047 HON27 RS03110 WP 000389061.1
 consensus

IPVALLTSRGADRHRKVA AKLGATGYFTKPYTEKDILDAANRMLNGEVLLPGST 928
 ILDLMLPEIDGLEVAKTIRK...TSSVPILMLSAKDSEFDKVI GLELGADDYVTKPFSNRELQARVKALLRRSQMPVDG 126
 ILDLMLPELDGLEVAKEVRK...NSHVPIIMLSAKDSEFDKVI GLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAV 126
 LLDIEMPRMDGFEVLNLRHHDMHQYMPIIMITSRTGKHRERAFSLGVSQYMGKPFQEEELLENIDALLVAFESEVKS. 1506
 VLDIVMPRMNGYEVCCRILKSNPKTQNVVVMCSSKGEFDRYWGMRQADAYIAKPFQPTLVGTIKQLLRG..... 121
 FVDIMPRLDGYQTCALIKNSQNYQNIPVIMLSSKDGFLDQAKGRVVGSDYELTKPFSKDELLNARLNVSS..... 127
 *!*****!*****!***** * * * *!***** * * * *!***** * * * *!***** * * * *

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NTDB id 1419 SGL RS12550 WP 010873252.1	RKTNSILSRFKKARGNFSEEAPTIVRSNPGTL.....R.....STVLIIDDSVMVREMLSMTFKKADYHTETARDGQDAW	998
NTDB id 413 AAK55817.1 141..845()	Q.EADSKPQ.PIQIGDLEIVPDAYVAKKYGEELDLTHREFELLYHLASHTGQVITREHLLET.....VW	188
NTDB id 374 SMU RS06885 WP 002262930.1	AEENASGIP.EIIIGDLQILPDAFVAKKRGTEVELTHREFELLHHLATHTGQVMTREHLLET.....VW	189
NTDB id 1046 HON27 RS03130 WP 168726941.1	1506
NTDB id 221299 BMF81 RS00865 WP 006194267.1	121
NTDB id 1417 SGL RS04200 WP 014407071.1	121
NTDB id 1047 HON27 RS03110 WP 000389061.1	127
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	EKLLKSGLPCLLILCDIEMPRMNGLELLERIQEDPNLSRIPVAMITSRGAQKMQRMAAEKGAQKGYFVKPYIEETLLEASQR	1078
NTDB id 413 AAK55817.1 141..845()	GYDYFG...DVRTVDVTV.....RRLREKIEDTPSR...PEYILT.....RRGVGYMKNNA*.....	234
NTDB id 374 SMU RS06885 WP 002262930.1	GYDYFG...DVRTVDVTV.....RRLREKIEDTPSR...PEYILT.....RRGVGYMKNNA*.....	235
NTDB id 1046 HON27 RS03130 WP 168726941.1	1506
NTDB id 221299 BMF81 RS00865 WP 006194267.1	121
NTDB id 1417 SGL RS04200 WP 014407071.1	121
NTDB id 1047 HON27 RS03110 WP 000389061.1	127
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	LIRGEVLLDLAALSADE	1095
NTDB id 413 AAK55817.1 141..845()	234
NTDB id 374 SMU RS06885 WP 002262930.1	235
NTDB id 1046 HON27 RS03130 WP 168726941.1	1506
NTDB id 221299 BMF81 RS00865 WP 006194267.1	121
NTDB id 1417 SGL RS04200 WP 014407071.1	121
NTDB id 1047 HON27 RS03110 WP 000389061.1	127
consensus		

- non conserved
- similar
- ≥ 50% conserved