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	MKEILKTLTEAMLPEDSYSEQDAEFLEIFIEEIEEIFVDLQPLINKWMQSENIATLTEIRRHFTLKGSGRMI GAKSSAELAWT	
NTDB id 1419 SGL RS12550 WP 010873252.1	.....	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	MKEILKTLTEAMLPEDSYSEQDAEFLEIFIEEIEEIFVDLQPLINKWMQSENIATLTEIRRHFTLKGSGRMI GAKSSAELAWT	85
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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	VEDTLNRVINQSLQLTPTTIQSYVQLVFKFYFLKLVDFNFKQKRAHTLDFRPLILLGQQQLQQQSSLEPALEELLQLSHTLMAETVTG	
NTDB id 1419 SGL RS12550 WP 010873252.1	.....	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	VEDTLNRVINQSLQLTPTTIQSYVQLVFKFYFLKLVDFNFKQKRAHTLDFRPLILLGQQQLQQQSSLEPALEELLQLSHTLMAETVTG	170
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
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	LELDYIEQDSLAEPIIQATHTEIEINLDETTLTFMEEAEEHLATIHQFLDQELHQYDSYNALIRALHTLRGSSAMAQVETIFEAS	
NTDB id 1419 SGL RS12550 WP 010873252.1	.....	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	LELDYIEQDSLAEPIIQATHTEIEINLDETTLTFMEEAEEHLATIHQFLDQELHQYDSYNALIRALHTLRGSSAMAQVETIFEAS	255
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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	TKVEHLFKILLQEELSSHSEEILLLQYREFVDRSLELLSRYSSSEQLLEARLLQFNQSWDAYVEQHGDRTPMLPESHGLVSQLLQ	
NTDB id 1419 SGL RS12550 WP 010873252.1	.....	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	TKVEHLFKILLQEELSSHSEEILLLQYREFVDRSLELLSRYSSSEQLLEARLLQFNQSWDAYVEQHGDRTPMLPESHGLVSQLLQ	340
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
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NTDB id 1419 SGL RS12550 WP 010873252.1	LDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEVLLDKPALLQSDYIFEIYQKAHQQLI	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	LDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEVLLDKPALLQSDYIFEIYQKAHQQLI	425
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
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NTDB id 1419 SGL RS12550 WP 010873252.1	QLFDALAAAGQRVGVVKQHQSI L EELKLYTQYTPDISNDLPQQDSTS FEP I YN I E P S I M T S D P N E W A T V E N F S D S A D W A V L G Q S V Q Q D R Q Y	6
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	QLFDALAAAGQRVGVVKQHQSI L EELKLYTQYTPDISNDLPQQDSTS FEP I YN I E P S I M T S D P N E W A T V E N F S D S A D W A V L G Q S V Q Q D R Q Y	510
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	I S S T Q V N R N F D A D L L D I F L E E A E E L L E G I D T D L N I W V G E Q E N F A A L N N L M R Y L H T L K G G A N M V Q A T Y L G L I A H E L E S I Y E R L I Q K	31
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	I S S T Q V N R N F D A D L L D I F L E E A E E L L E G I D T D L N I W V G E Q E N F A A L N N L M R Y L H T L K G G A N M V Q A T Y L G L I A H E L E S I Y E R L I Q K	595
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	Q L V V T S D L I D F I R L V Q D D L A D R L Q I M R E Q Q L D Y A A P Y T I N A L K R A G Q N S N F Q P L P V V E A F D A E S E V F S E Q K V I S E I I I D E I P V E L	60
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	Q L V V T S D L I D F I R L V Q D D L A D R L Q I M R E Q Q L D Y A A P Y T I N A L K R A G Q N S N F Q P L P V V E A F D A E S E V F S E Q K V I S E I I I D E I P V E L	680
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	EPALAELETHHDQVFDTSYI <sup>EL</sup> ER <sup>P</sup> VE <sup>M</sup> IT <sup>D</sup> Y <sup>K</sup> SP <sup>E</sup> EX <sup>E</sup> Y <sup>R</sup> ANEQDI <sup>E</sup> AV <sup>V</sup> ERT <sup>F</sup> TE <sup>R</sup> ET <sup>T</sup> E <sup>T</sup> Y <sup>R</sup> E <sup>F</sup> LE <sup>R</sup> SS <sup>W</sup> LE <sup>F</sup> B <sup>I</sup> N <sup>R</sup> EADPLA	118
NTDB id 413 AAK55817.1 141..845( )	.....PSSGEEDPPVRMTLDFNSPEAELD.....ALFTDEDAPTTAIDPFESSSWLEIDQMMEADPLA	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	EPALAELETHHDQVFDTA <sup>V</sup> TELAT <sup>P</sup> VEMIT <sup>T</sup> VTSQENEVAANEQDIEAVVEQTFLEEATELLEMAESLLKQWFEQRTNR.....	759
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	S <sup>I</sup> PETNAPAYQTAGGLDDN <sup>D</sup> L <sup>L</sup> ELQDLLGVGDVEPA <sup>T</sup> LR <sup>B</sup> Y <sup>H</sup> S <sup>K</sup> G <sup>R</sup> RV <sup>M</sup> Y <sup>L</sup> EN <sup>M</sup> L <sup>T</sup> AY <sup>R</sup> ENAFEQFA <sup>T</sup> H <sup>H</sup> F <sup>N</sup> S <sup>I</sup> Y <sup>R</sup> F <sup>L</sup>	194
NTDB id 413 AAK55817.1 141..845( )	SIPETNAPAYQTAGGLDDNDLAE <sup>L</sup> QDLLGVGDVEPAITNPGLHSFAGADRVTPLTNEVLL.FA.....AIDDLGLINAPVL	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	S.....ILL.....QLQRAVHSLKGGARMVGL <sup>E</sup> AVQAIAYQLENAFEQFALH <sup>H</sup> F <sup>N</sup> S <sup>N</sup> IYD <sup>H</sup> LL	812
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	EE <sup>F</sup> LA <sup>W</sup> FLKDAIFNHN <sup>Y</sup> QH <sup>F</sup> FDGLQ <sup>S</sup> LENIQ <sup>F</sup> .FETTIQ <sup>I</sup> PTKLTRA.DLFSS.EP.....VMTFIQGDGTEPPPMGAWEQ	266
NTDB id 413 AAK55817.1 141..845( )	DEEFSF.....TELETLIDTPAPKSFATISHPPFRLPVAMEEDEFKDLALLEQADSSLG <sup>G</sup> GIAGMGGARLPARP...R	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	ESAI <sup>A</sup> WLKDAIFNHN <sup>Y</sup> QH <sup>F</sup> FDGLQ <sup>S</sup> LENIQ <sup>F</sup> .FETTIQ <sup>I</sup> PTKLTRA.DLFSS.EP.....VMTFIQGDGTEPPPMGAWEQ	885
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	QVKSRSFEQ <sup>T</sup> MRVPIKQLDNLSNLIGELVVKRNRLEQDQERLRQFLDNLLNQAQNLSDVGGRMQDLYERTLLEGALLASRSTNSR	351
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	TERLDQNNEMIRVSADLIEKMIDLSGENSINRSRIEMDLSQFSHTLVEMELAIQRLADQLRRMEGELETQI.....	956
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
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NTDB id 1419 SGL RS12550 WP 010873252.1	HHSSKTTYPDAR IAKFSSTETSRYSIBFDALVMDYISFHNLSSRMAL EAYBYKREPKSSTISKYBETESYLRISRBYGAEYQEQMR	436
NTDB id 413 AAK55817.1 141..845( )	HHSSKTTYPDARITASSSSTMNDQPLDALEMDRFTGFHLLSQEMIELIVRVRESASDIQYLVDETDQVARTLRQVTTQLQEGMTK	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	.....IAKHGIEHSRYTDFDPLEMDQYSSLNQLSKSLAESASDLVDFKNTLSDKIRDTESLLLQQSRIQAEIQEGLMR	1029
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	SRMVPFPTQTADRLPRAIREVSMKLNKKVNLEVEGGDVLIDKMLEHLYNPMTHLVNNAITHGIEPTAERQKLGKSPEGHLAVRAF	521
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	TRLVPPFSRLLPRLQRIVRQTSTALNRPaelFVNNTegELDRNILERLVTPLEHMLRNAIDHGLEDRAQRQQANKPETGRIELNIQ	1114
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	LQGNQTLITVSDDGAGINTEMVKQKAVEKGLISRMEASLSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVRTSLIDIRGAIN	606
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	RQGTDVVVVFSDDGQGIDVEKVRQKALLAGLIKP..EQDLEQQDILQLIFHPGLSTAEQVTQISGRGVGLDVVQSDIKSLGGHVS	1197
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

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NTDB id 1419 SGL RS12550 WP 010873252.1	IESTVGKGTTFITIRLPLTL SICKALCCLSDRARIAPFMDGVEDTRDYLPKDIVTD.EDGQKCVKYGKEALLLPLYPLKDLLNYNR	690
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	VESVYGQGTTFITIRVPTTVAVSDALMVKVADQQFAIPLAQIDRIIRVSPASLEQYFESPQELFEYEN...KRYPLRYLSEFVG	1277
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		



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NTDB id 1419 SGL RS12550 WP 010873252.1	MNGLELLERIQEDPNLSRIPVAMITSRGAQKMRMAAEKGAKEKGFVRRNNAEETLLEASQRLIRGEVLLDLAALSAD	1095
NTDB id 413 AAK55817.1 141..845( )	...RRLREKIEDTPSR...PEYILT.....RRGVGYMNRNNA*.....	234
NTDB id 374 SMU RS06885 WP 002262930.1	...RRLREKIEDTPSR...PEYILT.....RRGVGYMKSYD.....	235
NTDB id 1046 HON27 RS03130 WP 168726941.1	.....	1506
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	127
NTDB id 66457 LBWT RS02620 WP 017290869.1	.....	121
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	121
consensus		

- non conserved
- similar
- ≥ 50% conserved