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NTDB id 1029	TT RS02230 WP 011228203.1	PLLLLAVALFFAYRW.YYGTIPQGRRV	IDRLKLRPLVFGNINRKTAVARFSRTLALLSSSGVNIVEALDITKGTAGNSVVE	303
NTDB id 1169	A1552VC RS11075 WP 000648511.1	WAFIAIAAAIFGLKALRKN	SFQIRLKT	308
NTDB id 1402	DSB67 RS12670 WP 010643257.1	FIGLSFVSL	LISGKIMSQRSDSFRLLAARSSLRVPILGPVLSKAAIAKFSRTLATSFSAGIPILTALKTTSKTSGNMHYQ	307
NTDB id 1112	NGFG RS09215 WP 003689811.1	MVLIALGF	FAIYRFLKLKARSIKIQRRMDAILLRMPIFGDIVRKGTIARWGRTTATLFAAGVPLVDVLDSTAGAAGNLIYE	310
NTDB id 1113	AAA85695.1 219..1451()	MVLIALGF	FAIYRFLKLKARSIKIQRRMDAILLRMPIFGDIVRKGTIARWGRTTATLFAAGVPLVDVLDSTAGAAGNLIYE	310
NTDB id 1252	GC085 RS07730 WP 011213805.1	IIIFGALGCVVYSFFYAKN	HSLEFAQTTIDRVMLKLPVIGPILEKAAIARFARTLSITFAAGLPLVEALKSVAGATGNIIYA	307
NTDB id 1198	PSJM300 03950 AFN76868.1	VVLAGFVGAAYAFKQAHT	RSEKFRNWFDRMLMKVPVVGDIYKSSVARFARTLSTTFAAGVPLVDALDSVAGATGNVVF	306
NTDB id 1016	ACIAD RS01680 WP 004920476.1	ILILVIGAAT	AGFLEARKRSKRFRDQLDRLVLRPIVGDVLYKAI	309
NTDB id 1053737	ABUS41 RS17115 WP 000279216.1	IMIATIGAVIAAFLEAKKRSK	KFRDGLDKLALKLPIFGDLVYKAI	309
NTDB id 1059	ABD1 RS01610 WP 000279215.1	IMIATIGAVIAAFLEAKKRSK	KFRDGLDKLALKLPIFGDLVYKAI	309
consensus		*****	*****!*****!*	309

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NTDB id 1029	TT RS02230 WP 011228203.1	EIVEAAKLKIQQGDPLNLT	LQAHPFVFPPMVSSMVAIGEETGALD	TMLSKVADFYER	EVDEAVASLTAAIEPLMITFLGV	383	
NTDB id 1169	A1552VC RS11075 WP 000648511.1	TAINEVYRD	TAAGMPMYIAMR.NTEAFPEMVLQ	VMIGEESGQLDDMLNKVAT	IYEFVDNTVDNLGKILEPLIIVFLGT	387	
NTDB id 1402	DSB67 RS12670 WP 010643257.1	LAIEEVYRD	TAAGMPMYIAMR.HCHVFPELVLQ	VMIGEESGRLDDMLNKIAAI	IYEFVDNTVDNLGKILEPLIIVFLGV	386	
NTDB id 1112	NGFG RS09215 WP 003689811.1	EATREIRTR	VIQGLSMTSGMR.ATELFPNMMLQ	MSSIGEESGLDDMLNKAAEFY	EVDNAVGRLSAMMEPIIIVILGL	389	
NTDB id 1113	AAA85695.1 219..1451()	EATREIRTR	VIQGLSMTSGMR.ATELFPNMMLQ	MSSIGEESGLDDMLNKAAEFY	EVDNAVGRLSAMMEPIIIVILGL	389	
NTDB id 1252	GC085 RS07730 WP 011213805.1	KATDKIRE	EVATGQQMFIAME.NTHLFPNMVIQ	MVAIGEESGALDEKMLS	KVADFYEEVNNAVDALSSLEPIIMSILGI	386	
NTDB id 1198	PSJM300 03950 AFN76868.1	SATNKIKSDVSSGMQLNFSMR	.TTGTFPMSMAVQMTAIGEESGALDEMLS	KVATFYED	EVDNMVDGLTALMEPMIMAVLGV	385	
NTDB id 1016	ACIAD RS01680 WP 004920476.1	QAVLKIRE	DVSTGQQLNFAMR.VSNRFPTMAIQ	MVAIGEESGALDEMLDKVAN	YYESEVDH	AVDGLTSMMEPLIMAILGI	388
NTDB id 1053737	ABUS41 RS17115 WP 000279216.1	KAVMKIRE	DVATGQQLQFAMR.VSNRFPTMAIQ	MVAIGEESGALDSMLDKVAT	YYENEVDNAVDGLTSMMEPLIMAILGV	388	
NTDB id 1059	ABD1 RS01610 WP 000279215.1	KAVMKIRE	DVATGQQLQFAMR.VSNRFPTMAIQ	MVAIGEESGALDSMLDKVAT	YYENEVDNAVDGLTSMMEPLIMAILGV	388	
consensus		***	*****!*****	***	*!!!!	*****!*****!*	388

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NTDB id 1029	TT RS02230 WP 011228203.1	IVGMIVAGMFLPLFKIIGT	ISVQ	406
NTDB id 1169	A1552VC RS11075 WP 000648511.1	VVGGLV	VAMYLPIFNLMSVLG..	408
NTDB id 1402	DSB67 RS12670 WP 010643257.1	VVGGLV	TAMYLPIFNLMSVLG..	407
NTDB id 1112	NGFG RS09215 WP 003689811.1	VIGTLL	VAMYLPLFNLGNVVA..	410
NTDB id 1113	AAA85695.1 219..1451()	VIGTLL	VAMYLPLFNLGNVVA*.	410
NTDB id 1252	GC085 RS07730 WP 011213805.1	LVGGLV	VAMYLPIFKLGS	AV... 406
NTDB id 1198	PSJM300 03950 AFN76868.1	LVGGLI	IAMYLPIFQMGSV...	405
NTDB id 1016	ACIAD RS01680 WP 004920476.1	LVGGLV	IAMYLPIFQMGSV...	408
NTDB id 1053737	ABUS41 RS17115 WP 000279216.1	LVGGLV	IAMYLPIFQMGSV...	408
NTDB id 1059	ABD1 RS01610 WP 000279215.1	LVGGLV	IAMYLPIFQMGSV...	408
consensus		***!*****!*	*****	

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved