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NTDB id 1029	TT RS02230 WP 011228203.1	PLLLLAVALLFFAYRW.YYGTIPQGRRV	IDRLKLRPLVFGNINRKTAVARFSRTLALLSSSGVNIVEALDITKGTAGNSVVE	303
NTDB id 1169	A1552VC RS11075 WP 000648511.1	WAFIAIAAAIFGLKALRKN	SFQIRLKT SRLGLKFPITIGNVLAKASIAKFSRTLATSFAAGIPILASLKT	308
NTDB id 1402	DSB67 RS12670 WP 010643257.1	FIGLSFVSL	LISGKIMSQRSDSFRLLAARSSLRVPILGPVLSKAAIAKFSRTLATSFSAGIPILTALKTT	307
NTDB id 1112	NGFG RS09215 WP 003689811.1	MVLIALGFAIYRFLKLKARSIKIQRRMDAILLRMP	IFGDIVRKGTIARWGRTTATLFAAGVPLVDVLDSTAGAAGNLIYE	310
NTDB id 1113	AAA85695.1 219..1451()	MVLIALGFAIYRFLKLKARSIKIQRRMDAILLRMP	IFGDIVRKGTIARWGRTTATLFAAGVPLVDVLDSTAGAAGNLIYE	310
NTDB id 1252	GC085 RS07730 WP 011213805.1	IIIFGALGGVVYSFFYAKNHSLEFAQT	IDRVMLKLPVIGPILEKAAIARFARTLSITFAAGLPLVEALKSVAGATGNIIYA	307
NTDB id 1198	PSJM300 03950 AFN76868.1	VVLAGFVGAAAYAFKQAHTRSEKFRNWF	DRLMLKVPVVGDIYKSSVARFARTLSTTFAAGVPLVDALDSVAGATGNVVF	306
NTDB id 1016	ACIAD RS01680 WP 004920476.1	ILILVIGAATAGFLEARKRSKRFRDQL	DLRLVLRPIVGDLYKAI IARYSRTLATTFAAGVPLIDALASTAGATNNTVYE	309
NTDB id 1006045	ABKP32 RS16595 WP 000279216.1	IMITAI GAVIAAFLEAKKRSKKFRDGL	DKLALKLPIFGDLVYKAI IARYSRTLATTFAAGVPLIDALESTAGATNNVIYE	309
NTDB id 1059	ABD1 RS01610 WP 000279215.1	IMITAI GAVIAAFLEAKKRSKKFRDGL	DKLALKLPIFGDLVYKAI IARYSRTLATTFAAGVPLIDALESTAGATNNVIYE	309
consensus		*****	*****!*****!*	

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NTDB id 1029	TT RS02230 WP 011228203.1	EIVEAAKLKIQGGDP	NLTLAQHPFVFPPMVSSMVAIGEETGALD	TMLSKVADFYEREVDEAVASLTAAIEPLMITIFLGV	383
NTDB id 1169	A1552VC RS11075 WP 000648511.1	TAINEVYRD	TAAGMPMYIAMR.NTEAFPEMVLQVM	MIGEESGQLDDMLNKVATIIYEF	387
NTDB id 1402	DSB67 RS12670 WP 010643257.1	LAIEEVYRD	TAAGMPMYIAMR.NTEAFPEMVLQVM	MIGEESGQLDDMLNKVATIIYEF	386
NTDB id 1112	NGFG RS09215 WP 003689811.1	EATREIRTRVIQGLSMTSGMR	.ATELFPNMMLQMSSIGEESGSLDDMLN	KAAEFYEDEVDAVGRLSAMMEPIIIVILGL	389
NTDB id 1113	AAA85695.1 219..1451()	EATREIRTRVIQGLSMTSGMR	.ATELFPNMMLQMSSIGEESGSLDDMLN	KAAEFYEDEVDAVGRLSAMMEPIIIVILGL	389
NTDB id 1252	GC085 RS07730 WP 011213805.1	KATDKIREEVATGQQMF	FIAME.NTHLFPNMVIQMV	AIGEESGALDEKMLSKVADFYE	386
NTDB id 1198	PSJM300 03950 AFN76868.1	SATNKKISDVSSGMQLNFSMR	.TTGTFPSPMAVQMTAIGEESGALDE	MLSKVATFYEDVDNMVDGLTALMEPMIMAVLGV	385
NTDB id 1016	ACIAD RS01680 WP 004920476.1	QAVLKIREDVSTGQQLNFAMR	.VSNRFPTMAIQMVAIGEESGALDE	MLDKVANYYESEVDHVDGLTSMMEPLIMAILGI	388
NTDB id 1006045	ABKP32 RS16595 WP 000279216.1	KAVMKIREDVATGQQLQFAMR	.VSNRFPTMAIQMVAIGEESGALDE	MLDKVANYYENEVDNAVDGLTSMMEPLIMAILGV	388
NTDB id 1059	ABD1 RS01610 WP 000279215.1	KAVMKIREDVATGQQLQFAMR	.ISNRFPTMAIQMVAIGEESGALDE	MLDKVANYYENEVDNAVDGLTSMMEPLIMAILGV	388
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NTDB id 1029	TT RS02230 WP 011228203.1	IVGMIVAGMFLPLFKIIGT	LSVQ	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	406
NTDB id 1198	PSJM300 03950 AFN76868.1	LVGGLI	IAMYLPIFQMGSV..	405
NTDB id 1016	ACIAD RS01680 WP 004920476.1	LVGGLV	IAMYLPIFQMGSV..	408
NTDB id 1006045	ABKP32 RS16595 WP 000279216.1	LVGGLV	IAMYLPIFQMGSV..	408
NTDB id 1059	ABD1 RS01610 WP 000279215.1	LVGGLV	IAMYLPIFQMGSV..	408
consensus		***!*****!***!*	*****	

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