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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 SRLGLKFPITIGNVLAKASIAKFSRTLATSFAAGIPILASLKT	308
NTDB id 1402	DSB67 RS12670 WP 010643257.1	FIGLSFVSLISGKIMSQRSDS	FRLLAARSSLRVPILGPVLSKAAIAKFSRTLATSFSAGIPILTALKTT	307
NTDB id 1112	NGFG RS09215 WP 003689811.1	MVLIALGFATIGYFLKLR	KARSIKIQRMDAILLRMPIFGDIVRKGTIARWGRTTATLFAAGVPLVDVLD	310
NTDB id 1113	AAA85695.1 219..1451()	MVLIALGFATIGYFLKLR	KARSIKIQRMDAILLRMPIFGDIVRKGTIARWGRTTATLFAAGVPLVDVLD	310
NTDB id 1252	GC085 RS07730 WP 011213805.1	IIIFGALGGVVVSFFFYAKN	HSLEFAQTTIDRVMLKLPVIGPILEKAAIARFARTLSITFAAGLPLVEALKSV	307
NTDB id 1198	PSJM300 03950 AFN76868.1	VVLAGFVGAAAYAFKQAH	TRSEKFRNWFDRMLMKVPVVGDIYKSSVARFARTLSTTFAAGVPLVDAL	306
NTDB id 1016	ACIAD RS01680 WP 004920476.1	ILILVIGAATAGFLEARK	RSKRFRDQLDRLVLRPIVGDLYKAI	309
NTDB id 1053356	ABUS27 RS15980 WP 000279218.1	IMITATIGAVIAAFLEAK	KRSKKFRDGLDKLALKLPIFGDLVYKAI	309
NTDB id 1059	ABD1 RS01610 WP 000279215.1	IMITATIGAVIAAFLEAK	KRSKKFRDGLDKLALKLPIFGDLVYKAI	309
consensus		*****	*****!*****!*****!*****!*****!*****!*****!*****!*****!*****	

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NTDB id 1029	TT RS02230 WP 011228203.1	EIVEAAKLKIQQGDPLNLT	LTAQHPFVFPPMVSSMVAIGEETGALD	TMLSKVADFYEVEVDEAVASLTAAIEPLMIT	383
NTDB id 1169	A1552VC RS11075 WP 000648511.1	TAINEVYRDTAAGMPMYIAMR	.NTEAFPENVLQVMVIGEESGQLDDMLN	KVATIEYEFVDNTVDNLGKILEPLIIVFLGT	387
NTDB id 1402	DSB67 RS12670 WP 010643257.1	LAIEEVYRDTAAGMPMYIAMR	.HCHVFPELVLMQVMIGEESGRLLDMLN	KAAIEYEFVDNTVDNLGKILEPLIIVFLGV	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	KATDKIREEVATGQQMFIA	ME.NTHLFPNMVIQMV	AIGEESGALDEKMLSKVADFYEVEV	386
NTDB id 1198	PSJM300 03950 AFN76868.1	SATNKKISDVSSGMQLNFS	SMR.TTGTFPMAVQMTAIGEESGALDE	MLSKVATFYEDEVDNMVDGLTALMEPM	385
NTDB id 1016	ACIAD RS01680 WP 004920476.1	QAVLKIREDVSTGQQLNFAMR	.VSNRFPTMAIQMVAIGEESGALDEMLD	KVANYYESEVDHVDGLTSMMEPLIMAILGI	388
NTDB id 1053356	ABUS27 RS15980 WP 000279218.1	KAVMKIREDVATGQQLQFAMR	.VSNRFPTMAIQMVAIGEESGALDSMLD	KVATYYENEVDNAVDGLTSMMEPLIMAILGV	388
NTDB id 1059	ABD1 RS01610 WP 000279215.1	KAVMKIREDVATGQQLQFAMR	.ISNRFPTMAIQMVAIGEESGALDSMLD	KVATYYENEVDNAVDGLTSMMEPLIMAILGV	388
consensus		***	*****!*****	***	***!*****!*****!*****!*****!*****!*****!*****!*****!*****

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NTDB id 1029	TT RS02230 WP 011228203.1	IVGMIVAGMFLPLFKIIGT	ISVQ	406
NTDB id 1169	A1552VC RS11075 WP 000648511.1	VVGGLVIVAMYLPIFNLMS	VLG..	408
NTDB id 1402	DSB67 RS12670 WP 010643257.1	VVGGLVIVAMYLPIFNLMS	VLG..	407
NTDB id 1112	NGFG RS09215 WP 003689811.1	VIGTLLVAMYLPLFNLGN	VVA..	410
NTDB id 1113	AAA85695.1 219..1451()	VIGTLLVAMYLPLFNLGN	VVA*.	410
NTDB id 1252	GC085 RS07730 WP 011213805.1	LVGGLVIVAMYLPIFKLGS	AV...	406
NTDB id 1198	PSJM300 03950 AFN76868.1	LVGGLVIVAMYLPIFQMGS	VV...	405
NTDB id 1016	ACIAD RS01680 WP 004920476.1	LVGGLVIVAMYLPIFQMGS	VV...	408
NTDB id 1053356	ABUS27 RS15980 WP 000279218.1	LVGGLVIVAMYLPIFQMGS	VV...	408
NTDB id 1059	ABD1 RS01610 WP 000279215.1	LVGGLVIVAMYLPIFQMGS	VV...	408
consensus		***!*****!*****!*****!*****!*****		

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