


```

NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 168073 AL539 RS08745 WP 047102659.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus

```

Sequence	Position
EIRDVATGQMLNMRNINLFPNMLQMAIGEESGALDDMLNKVATIYEFVDNVTVDNLGKILEPLIIVFLGTVVGG	392
EYVRDTAAGMPMYIAMRNTAEAFPEMVLQMVMIGEESGRLDDMLNKVATIYEFVDNVTVDNLSKILEPLIIVFLGTVVGG	391
EYVRDTAAGMPMYIAMRHCHVFPFELVLQMVMIGEESGRLDDMLNKTAIYEFVDNVTVDNLSKILEPLIIVFLGTVVGG	391
EIRTRVIQGLSMTSGMRATELFPNMMLQMSSIGEESGSLDDMLNKAAEFYDEVDNAVGNLSAMMEPIIILGLVIGTL	394
EIRTRVIQGLSMTSGMRATELFPNMMLQMSSIGEESGSLDDMLNKAAEFYDEVDNAVGRLSAMMEPIIIVILGLVIGTL	394
KIREEVATGQQMFIAMENTHLFPNMVIQMVAIGEESGALDKMLSKVADFYEEVNNAVDALSSILEPIIMSILGILVGG	391
KIKSDVSSGQMLNFSMRTVNGTFPSMAVQMTAIGEESGALDEMLSKVATFYDEVDNMVDGLTALMEPMIMAVLGVLVGG	390
KIREDDVSTGQQLNFAMRVTSNRFPSTMAIQMTAIGEESGALDEMLDKVANFYEYSEVDHVDGLTSMMEPLIMAILGILVGG	393
KIREDDVATGQQLQFAMRISNRFPSTMAIQMTAIGEESGALDSMLDKVATYYENVEVDNAVGLTSMMEPLIMAILGVLVGG	393

logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1	VVAMYLPPIFNLM SV LG.	408
NTDB id 168073 AL539 RS08745 WP 047102659.1	VTAMYLPPIFNLM SV LG.	407
NTDB id 1402 DSB67 RS12670 WP 010643257.1	VTAMYLPPIFNLM SV LG.	407
NTDB id 1113 AAA85695.1 219..1451()	LVAMYLPPIFN LGN VVA*	410
NTDB id 1112 NFGF RS09215 WP 003689811.1	LVAMYLPPIFN LGN VVA.	410
NTDB id 1252 GC085 RS07730 WP 011213805.1	VVAMYLPPIF KL GS AV ..	406
NTDB id 1198 PSJM300 03950 AFN76868.1	II AMYLPPIF QM GSVV..	405
NTDB id 1016 ACIAD RS01680 WP 004920476.1	VIAMYLPPIF QM GSVV..	408
NTDB id 1059 ABD1 RS01610 WP 000279215.1	VIAMYLPPIF QM GSVV..	408
consensus	* * ! ! ! ! ! * ! * * * * * * *	

- non conserved
- similar
- $\geq 50\%$ conserved