


```

NTDB id 412413 GTF75 RS02305 WP 000648511.1
NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 N6GF RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
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

```

logo

Sequence	Position
EYRDTAAGMPMYIAMRNTTEAFPEMVLQMVMIGEESQLDDMLNKVATIYEFYEVNDTVDNLGKILEPLIIIVFLGTIVVGG	392
EYRDTAAGMPMYIAMRNTTEAFPEMVLQMVMIGEESQLDDMLNKVATIYEFYEVNDTVDNLGKILEPLIIIVFLGTIVVGG	392
EYRDTAAGMPMYIAMRHCHVPELVLQMVMIGEESGRDDMLNKAATYEFYEVNDTVDNLSKILEPLIIIVFLGVVGG	391
EIRTRVIQGLSMTSGMRATELFPNMMLQMSSIGEESGLDDMLNKAAEFYDEVDNAVGRLSAMMEPTIIIVILGLVIGTL	394
EIRTRVIQGLSMTSGMRATELFPNMMLQMSSIGEESGLDDMLNKAAEFYDEVDNAVGRWSAMMEPTIIILILGLVIGTL	394
KIREEVATGQQMFIAMENTHLFPNMVIQMVAIGEESGALDKMLSKVADFYEEVNNAVDALSSILEPIIMSILGILVGG	391
KIKSDVSSGMQLNFSMRTNGTFPSMAVQMTAIGEESGALDEMLSKVATFYDEVDNMVDGLTALMEPMIMAVLGVLVGG	390
KIREVDSTGQQLNFAMRVSGRFPMTATQMVAIGEESGALDEMLDKVANFYEYSEVDHVDGLTSMMEPLIMAILGILVGG	393
KIREVDVATGQQLQFAMRISNRFPMTATQMVAIGEESGALDSMLDKVATFYENYEVNDNAVGLTSMMEPLIMAILGVLVGG	393

[illegible]

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