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logo
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NTDB id 506 SM12261 RS00975 WP 000265657.1     MWKKQKVKAGVLLYAVTMAAIFSLLLQFYLNRQVAHYQDYTLNKEKLVAFAMAKRTKDKVEQESGEQAFNLGQVSYQNKK   80
NTDB id 534 SMSK321 RS0102610 WP 000265640.1   MWKKQKVKAGVLIYAVTIAAIFSLLLQFYLNRQVAHYQDYALNKEKLIAFAMAKRTKGKVEQESGEQAFNLGQVSYQNKK   80
NTDB id 49192 INV104 RS10285 WP 001818966.1    MWKKKKVKAGVLLYAVTIAAIFSLLLQFYLTRQIAHYQDYALNKEKLVAFAMAKRTKDKAEQESGEQVFNLGQVSYQNKK   80
NTDB id 276 KZH43 RS09235 WP 000265622.1       MWKKKKVKAGVLLYAVTIAAIFSLLLQFYLNRQVAHYQDYALNKEKLVAFAMAKRTKDKVEQESGEQFFNLGQVSYQNKK   80
NTDB id 235 SPD RS09865 WP 000265622.1         MWKKKKVKAGVLLYAVTIAAIFSLLLQFYLNRQVAHYQDYALNKEKLVAFAMAKRTKDKVEQESGEQFFNLGQVSYQNKK   80
NTDB id 201 SPR RS09400 WP 000265622.1         MWKKKKVKAGVLLYAVTIAAIFSLLLQFYLNRQVAHYQDYALNKEKLVAFAMAKRTKDKVEQESGEQFFNLGQVSYQNKK   80
NTDB id 166 SP RS10350 WP 000265622.1          MWKKKKVKAGVLLYAVTIAAIFSLLLQFYLNRQVAHYQDYALNKEKLVAFAMAKRTKDKVEQESGEQFFNLGQVSYQNKK   80
consensus                                      !!!!*!!!!!!!*!!!!*!!!!!!!!!!!!!*!!*!!!!!!*!!!!!!*!!!!!!!!!*!*!!!!!!!*!!!!!!!!!!!!
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logo
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NTDB id 506 SM12261 RS00975 WP 000265657.1     TSLVTTVRTPKSQYEFIFPSVKIKEEKTDKKEKVVTDSSNQAEKKKSEEKVEKKENF   137
NTDB id 534 SMSK321 RS0102610 WP 000265640.1   TSLVTRVRTPKSQYEFIFPSVKLKEEKTAKKEKVVTDSSNQAEKKKSEEKVEKKENS   137
NTDB id 49192 INV104 RS10285 WP 001818966.1    TSLVTTVRTDKSQYEFLFPSVKIKEEKTDKKEEVATDSSEKAEKKIRREA.......   130
NTDB id 276 KZH43 RS09235 WP 000265622.1       TGLVTRVRTDKSQYEFLFPSVKIKEEKRDKKEEVATDSSEKVEKKKSEEKPEKKENS   137
NTDB id 235 SPD RS09865 WP 000265622.1         TGLVTRVRTDKSQYEFLFPSVKIKEEKRDKKEEVATDSSEKVEKKKSEEKPEKKENS   137
NTDB id 201 SPR RS09400 WP 000265622.1         TGLVTRVRTDKSQYEFLFPSVKIKEEKRDKKEEVATDSSEKVEKKKSEEKPEKKENS   137
NTDB id 166 SP RS10350 WP 000265622.1          TGLVTRVRTDKSQYEFLFPSVKIKEEKRDKKEEVATDSSEKVEKKKSEEKPEKKENS   137
consensus                                      !*!!!*!!!*!!!!!!!*!!!!!*!!!!*!*!!!*!*!!!!***!!!***!*******
```

X non conserved

X similar

X ≥ 50% conserved