```
logo
NTDB id 400 FSA28 RS09375 WP 002263441.1      MLFMKKIKAGVLLYALFMSAIFSLFLQFYLNRVVATERQNQAQLSASKAYLIADLTQDLAKDNFGQLTFNHGRSSYHKQE   80
NTDB id 391 SMU RS09005 WP 002263441.1        MLFMKKIKAGVLLYALFMSAIFSLFLQFYLNRVVATERQNQAQLSASKAYLIADLTQDLAKDNFGQLTFNHGRSSYHKQE   80
NTDB id 1128356 FJH52 RS09805 WP 000265617.1  MWKKKKVKAGVLLYAVTIAAIFSLLLQFYLNRQIAHYQDYALNKEKLVAFAMAKRTKDKAEQESGEQVFNLGQVSYQNKK   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
NTDB id 506 SM12261 RS00975 WP 000265657.1    MWKKQKVKAGVLLYAVTMAAIFSLLLQFYLNRQVAHYQDYTLNKEKLVAFAMAKRTKDKVEQESGEQAFNLGQVSYQNKK   80
NTDB id 534 SMSK321 RS0102610 WP 000265640.1  MWKKQKVKAGVLIYAVTIAAIFSLLLQFYLNRQVAHYQDYALNKEKLIAFAMAKRTKGKVEQESGEQAFNLGQVSYQNKK   80
consensus                                     !****!*!!!!!*!!****!!!!!*!!!!!!!!**!*************!***!**!********!**  !!*!**!!****

logo
NTDB id 400 FSA28 RS09375 WP 002263441.1      GELLIKVHLTDGQSYQYRFNCVTENKEKGK..FSVANKTFTKSIEKRHFQ.K.......   129
NTDB id 391 SMU RS09005 WP 002263441.1        GELLIKVHLTDGQSYQYRFNCVTENKEKGK..FSVANKTFTKSIEKRHFQ.K.......   129
NTDB id 1128356 FJH52 RS09805 WP 000265617.1  TSLVTTVRTS.KSQYEFLFPSVKIKEEKRDKKEEVATDSSE.KVEKKKSEEKPEKKENS   137
NTDB id 276 KZH43 RS09235 WP 000265622.1      TGLVTRVRTD.KSQYEFLFPSVKIKEEKRDKKEEVATDSSE.KVEKKKSEEKPEKKENS   137
NTDB id 235 SPD RS09865 WP 000265622.1        TGLVTRVRTD.KSQYEFLFPSVKIKEEKRDKKEEVATDSSE.KVEKKKSEEKPEKKENS   137
NTDB id 201 SPR RS09400 WP 000265622.1        TGLVTRVRTD.KSQYEFLFPSVKIKEEKRDKKEEVATDSSE.KVEKKKSEEKPEKKENS   137
NTDB id 166 SP RS10350 WP 000265622.1         TGLVTRVRTD.KSQYEFLFPSVKIKEEKRDKKEEVATDSSE.KVEKKKSEEKPEKKENS   137
NTDB id 506 SM12261 RS00975 WP 000265657.1    TSLVTTVRTP.KSQYEFIFPSVKIKEEKTDKKEKVVTDSSN.QAEKKKSEEKVEKKENF   137
NTDB id 534 SMSK321 RS0102610 WP 000265640.1  TSLVTRVRTP.KSQYEFIFPSVKLKEEKTAKKEKVVTDSSN.QAEKKKSEEKVEKKENS   137
consensus                                     *  !***!**  ***!***!**!****!!******!******  **!!*****!*******
```

X non conserved

X similar

X $\geq$ 50% conserved