```
logo
NTDB id 400 FSA28 RS09375 WP 002263441.1      MLFMKKIKAGVLLYALFMSAIFSLFLQFYLNRVVATERQNQAQLSASKAYLIADLTQDLAKDNFGQLTFNHGRSSYHKQE   80
NTDB id 391 SMU RS09005 WP 002263441.1        MLFMKKIKAGVLLYALFMSAIFSLFLQFYLNRVVATERQNQAQLSASKAYLIADLTQDLAKDNFGQLTFNHGRSSYHKQE   80
NTDB id 206241 BOC72 RS10285 WP 000265630.1   MWKKKKVKAGVLLYAVTMAAIFSLLLQFYLNRQVAHYQDYALNKEKLVAFAMAKRTKDKAEQESGEQVFNLGQVSYQNKK   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..FSVANKTFTKSIEKRHFQK.......   129
NTDB id 391 SMU RS09005 WP 002263441.1        GELLIKVHLTDGQSYQYRFNCVTENKEKGK..FSVANKTFTKSIEKRHFQK.......   129
NTDB id 206241 BOC72 RS10285 WP 000265630.1   TSLVTTVRT.DKSQYEFLFPSVKIKEEKRDKKEEVATDSSEKAEKKKSEEKPEKKENS   137
NTDB id 276 KZH43 RS09235 WP 000265622.1      TGLVTRVRT.DKSQYEFLFPSVKIKEEKRDKKEEVATDSSEKVEKKKSEEKPEKKENS   137
NTDB id 235 SPD RS09865 WP 000265622.1        TGLVTRVRT.DKSQYEFLFPSVKIKEEKRDKKEEVATDSSEKVEKKKSEEKPEKKENS   137
NTDB id 201 SPR RS09400 WP 000265622.1        TGLVTRVRT.DKSQYEFLFPSVKIKEEKRDKKEEVATDSSEKVEKKKSEEKPEKKENS   137
NTDB id 166 SP RS10350 WP 000265622.1         TGLVTRVRT.DKSQYEFLFPSVKIKEEKRDKKEEVATDSSEKVEKKKSEEKPEKKENS   137
NTDB id 506 SM12261 RS00975 WP 000265657.1    TSLVTTVRT.PKSQYEFIFPSVKIKEEKTDKKEKVVTDSSNQAEKKKSEEKVEKKENF   137
NTDB id 534 SMSK321 RS0102610 WP 000265640.1  TSLVTRVRT.PKSQYEFIFPSVKLKEEKTAKKEKVVTDSSNQAEKKKSEEKVEKKENS   137
consensus                                     *  !***!** ****!***!**!****!!******!******* **!****!*******
```

X non conserved

X similar

X $\geq$ 50% conserved