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logo
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```
NTDB id 1148440 MFPB19 RS06895 WP 056947129.1  ........MKKKRKAFTLIEMVVVLAIIALLILLVAPNLGAQKQRAEKKTDQALVTTLQTQVELASDEIGHEIKSLDELT   72
NTDB id 619 LCA RS06470 WP 011374999.1         ........MKKKRKAFTLIEMVVVLAIIALLILLVAPNLGAQKQRAEKKTDQALVTTLQTQVELASDEIGHQIKSLDELT   72
NTDB id 460 GPW51 RS00790 WP 011921720.1       ....MKKLVEMKVKGFTLVEMLVVLGIISLLLLLFVPNLSQQKDAIQKKGDAAVVKVVESQMELYELEHDEEATVADLQA   76
NTDB id 502 SM12261 RS00955 WP 000738641.1     MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNEDASLSKLQA   80
NTDB id 162 SP RS10370 WP 000738627.1          MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNEDASLRKLQA   80
NTDB id 272 KZH43 RS09255 WP 000738626.1       MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNEDASLRKLQA   80
NTDB id 231 SPD RS09885 WP 000738626.1         MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNEDASLRKLQA   80
NTDB id 197 SPR RS09420 WP 000738626.1         MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNEDASLRKLQA   80
NTDB id 530 SMSK321 RS09230 WP 000738643.1     MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNEDASLSKLQA   80
consensus                                      **********!*!*!!!*!!*!!!*!!**!*!!**!!!**!!*****!***!*!*****!*!!***********  ****
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logo
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NTDB id 1148440 MFPB19 RS06895 WP 056947129.1  DKYISKDQLKHAKEKGITIEGGTVKQK.   99
NTDB id 619 LCA RS06470 WP 011374999.1         DKYISKDQLKHAKEKGITIEGGTVKQK.   99
NTDB id 460 GPW51 RS00790 WP 011921720.1       KGYITEKQAKQYATAKK...........   93
NTDB id 502 SM12261 RS00955 WP 000738641.1     DGRITEEQAKAYKEYHAKQNTSQTVAD.  107
NTDB id 162 SP RS10370 WP 000738627.1          DGRITEEQAKAYKEYNDKNGGANRKVND  108
NTDB id 272 KZH43 RS09255 WP 000738626.1       DGRITEEQAKAYKEYHDKNGGANRKVND  108
NTDB id 231 SPD RS09885 WP 000738626.1         DGRITEEQAKAYKEYHDKNGGANRKVND  108
NTDB id 197 SPR RS09420 WP 000738626.1         DGRITEEQAKAYKEYHDKNGGANRKVND  108
NTDB id 530 SMSK321 RS09230 WP 000738643.1     DGRITEEQAKAYNEYYAKNGGANRKVND  108
consensus                                      ***!***!*!*****  ***********
```

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