

logo

NTDB id 615 LCA RS02545 WP 011374200.1
NTDB id 593 KW2 RS05130 WP 021037147.1
NTDB id 95322 QML92 RS10275 WP 000867588.1
NTDB id 277 KZH43 RS10090 WP 000867601.1
NTDB id 236 SPD RS10765 WP 000867601.1
NTDB id 202 SPR RS10250 WP 000867601.1
NTDB id 167 SP RS11275 WP 000867616.1
NTDB id 507 SM12261 RS09240 WP 000867722.1
NTDB id 535 SMSK321 RS10515 WP 000867726.1
consensus



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logo

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consensus

PNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHD

SVRMETLAAFDKRNQTKTEIKRQYSLMVTISLRNIALTSEIKNI NEAXLVMAIALBRKYAGNLVRIYYG

PAVRRIETLHNDPDQIKKITAFAQQIQLMVTITILERGVTFKKNVAVLVLGAEHMFVNEAVLVQIAGRACGRHKDYAQNPVHFYYQ

PKEKMAFVASTTKSRKTIVEAFRKKQVSILITTSILERGVTFSSIDVFVINSEHPNFTKSSALIQMAGRVRGDSKRPTGLVSFFHS

PNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHD

PNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHD

PNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHD

PNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHD

PNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHD

PNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHD

PHEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCVDVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHD

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consensus

GLNASIKKAIKEIQMNMKEAGL

DKSTAMCSLRQIKRQISGRFLQSTV

DYTRAIKGACRQIKTQNRGRRLQSTV

GKSLAMCQAQKEIKKMNRLLGGF.....

GLNASIKKAIKEIQMNMKEAGL.....

GLNASIKKAIKEIQMNMKEAGL.....

GLNASIKKAIKEIQMNMKEAGL.....

GLNASIKKAIKEIQMNMKEAGL.....

GLNASIKKAIKEIQMNMKEAGL.....

GLNTSIKKAIKEIQMNMKEAGL.....

GLNTSIKKAIKEIQMNMKEAGL.....

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445
430
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- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved