

logo

NTDB id 326381 EGY29 RS21925 WP 023098502.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



80
80

logo

NTDB id 326381 EGY29 RS21925 WP 023098502.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



160
160

logo

NTDB id 326381 EGY29 RS21925 WP 023098502.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



240
240

logo

NTDB id 326381 EGY29 RS21925 WP 023098502.1
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consensus



320
320

logo

NTDB id 326381 EGY29 RS21925 WP 023098502.1
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consensus



400
400

logo

NTDB id 326381 EGY29 RS21925 WP 023098502.1
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consensus



480
480

logo

NTDB id 326381 EGY29 RS21925 WP 023098502.1
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consensus



560
560

logo

NTDB id 326381 EGY29 RS21925 WP 023098502.1
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consensus

NQAAELLQNAIYDEPQRTDLRLKLMVYAEMGDREGFARQENELREI GGAQPQVEQLKSRYPAMVAVAAVAGLAGAKLAQ
NQAAELLQNAIYDEPQRTDLRLKLMVYAEMGDREGFARQENELREI GGAQPQVEQLKSRYPAMVAVAAVAGLAGAKLAQ
!!

640
640

logo

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consensus

DELDSFSLDDL SLDDSGHAAK PDAAGQDLDDAFDLSLDDLGGGDLGSDDVQADLKSDSGALDDLTLDSDLDLAASTAADK
DELDSFSLDDL SLDDSGHAAK PDAAGQDLDDAFDLSLDDLGGGDLGSDDVQADLKSDSGALDDLTLDSDLDLAASTAADK
DELDSFSLDDL SLDDSGHAAK PDAAGQDLDDAFDLSLDDLGG.....DDVQADLKSDSGALDDLTLDSDLDLAASTAADK
!!

720
715

logo

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PVDDLDFGLDFAELAE TSPQPKHDDL GDFSLNLDAPEDKLSDDDFLLSLNDEVPAAPADNEFTLDTEAAE EEPALSLPDD
PVDDLDFGLDFAELAE TSPQPKHDDL GDFSLNLDAPEDKLSDDDFLLSLNDEVPAAPADNEFTLDTEAAE EEPALSLPDD
PVDDLDFGLDFAELAE TSPQPKHDDL GDFSLNLDAPEDKLSDDDFLLSLNDEVPAAPADNEFTLDTEAAE EEPALSLPDD
!!

800
795

logo

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consensus

FDLSLADEPT E PAAPEK GEDSFAAQLDEVSAQLDELASNLDEPKSATPSFSAEDA AVASALDGDADDDFDL SGADEAAT
FDLSLADEPT E PAAPEK GEDSFAAQLDEVSAQLDELASNLDEPKSATPSFSAEDA AVASALDGDADDDFDL SGADEAAT
FDLSLADEPT E PAAPEK GEDSFAAQLDEVSAQLDELASNLDEPKSATPSFSAEDA AVASALDGDADDDFDL SGADEAAT
!!

880
875

logo

NTDB id 326381 EGY29 RS21925 WP 023098502.1
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consensus

KLDLARAYIDMGDSEGARDILDEVLAEGNDSQQAEARELLERLA
KLDLARAYIDMGDSEGARDILDEVLAEGNDSQQAEARELLERLA
KLDLARAYIDMGDSEGARDILDEVLAEGNDSQQAEARELLERLA
!!

924
919

⊠ non conserved
⊠ similar
⊠ ≥ 50% conserved