

logo

NTDB id 155456 A6701 RS09795 WP 023099389.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



80
80

logo

NTDB id 155456 A6701 RS09795 WP 023099389.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



160
160

logo

NTDB id 155456 A6701 RS09795 WP 023099389.1
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consensus



240
240

logo

NTDB id 155456 A6701 RS09795 WP 023099389.1
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consensus



320
320

logo

NTDB id 155456 A6701 RS09795 WP 023099389.1
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consensus



400
400

logo

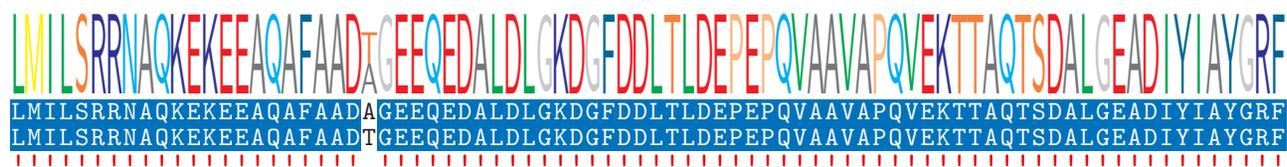
NTDB id 155456 A6701 RS09795 WP 023099389.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



480
480

logo

NTDB id 155456 A6701 RS09795 WP 023099389.1
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consensus



560
560

logo

NTDB id 155456 A6701 RS09795 WP 023099389.1
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consensus

NQAAELLQNAIYDEPQRTDLRLKLMEVYAEMGDREGFARQENELREI GGAQPQVEQLKSRYPAMVAVAAVAGLAGAKLAQ
NQAAELLQNAIYDEPQRTDLRLKLMEVYAEMGDREGFARQENELREI GGAQPQVEQLKSRYPAMVAVAAVAGLAGAKLAQ
!!

640
640

logo

NTDB id 155456 A6701 RS09795 WP 023099389.1
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consensus

DELDSFSLDDL SLDDSGHAAKPDAAAGQDLDDAFDLSLDDLGGDDVQADLKSDSGALDDLTLDSDLDLAASTAADKPVDDL
DELDSFSLDDL SLDDSGHAAKPDAAAGQDLDDAFDLSLDDLGGDDVQADLKSDSGALDDLTLDSDLDLAASTAADKPVDDL
DELDSFSLDDL SLDDSGHAAKPDAAAGQDLDDAFDLSLDDLGGDDVQADLKSDSGALDDLTLDSDLDLAASTAADKPVDDL
!!

720
720

logo

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consensus

DFGLDFAELAE TSPQPKHDDL GDFSLDL DAPEDKLSDDDFLLSLNDEVPAAPADNEFTLDTEAAEEPALSLPDDFDLSL
DFGLDFAELAE TSPQPKHDDL GDFSLDL DAPEDKLSDDDFLLSLNDEVPAAPADNEFTLDTEAAEEPALSLPDDFDLSL
DFGLDFAELAE TSPQPKHDDL GDFSLDL DAPEDKLSDDDFLLSLNDEVPAAPADNEFTLDTEAAEEPALSLPDDFDLSL
!!

800
800

logo

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ADEPTEPAAPEKGEDSFAAQLDEVSAQLDELASNLDEPKSATPSFSAEDA AVASALDGDADDDFDL SGADEAATKLDLA
ADEPTEPAAPEKGEDSFAAQLDEVSAQLDELASNLDEPKSATPSFSAEDA AVASALDGDADDDFDL SGADEAATKLDLA
ADEPTEPAAPEKGEDSFAAQLDEVSAQLDELASNLDEPKSATPSFSAEDA AVASALDGDADDDFDL SGADEAATKLDLA
!!

880
880

logo

NTDB id 155456 A6701 RS09795 WP 023099389.1
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RAYIDMGDSEGARDILDEVLAEGNDSQQAEARELLERLA
RAYIDMGDSEGARDILDEVLAEGNDSQQAEARELLERLA
RAYIDMGDSEGARDILDEVLAEGNDSQQAEARELLERLA
!!

919
919

⊠ non conserved
⊠ similar
⊠ ≥ 50% conserved