

logo

NTDB id 275447 CSB90 RS25110 WP 003130552.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



80
80

logo

NTDB id 275447 CSB90 RS25110 WP 003130552.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



160
160

logo

NTDB id 275447 CSB90 RS25110 WP 003130552.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



240
240

logo

NTDB id 275447 CSB90 RS25110 WP 003130552.1
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consensus



320
320

logo

NTDB id 275447 CSB90 RS25110 WP 003130552.1
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consensus



400
400

logo

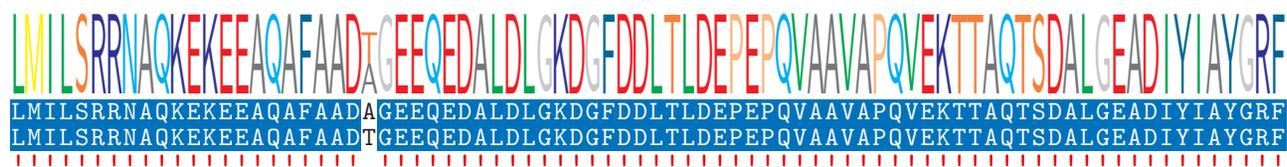
NTDB id 275447 CSB90 RS25110 WP 003130552.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



480
480

logo

NTDB id 275447 CSB90 RS25110 WP 003130552.1
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consensus



560
560

logo

NTDB id 275447 CSB90 RS25110 WP 003130552.1
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consensus

NQAAELLQNAIYDEPQRTDLRLKLMEVYAEMGDREGFARQENELREI GGAQPQVEQLKSRYPAMVAVAAVAGLAGAKLAQ
NQAAELLQNAIYDEPQRTDLRLKLMEVYAEMGDREGFARQENELREI GGAQPQVEQLKSRYPAMVAVAAVAGLAGAKLAQ
!!

640
640

logo

NTDB id 275447 CSB90 RS25110 WP 003130552.1
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DELDSFSLDDL SLDDSGHAAK PDAAGQDLDDAFDLSLDDLGGGDLGSDDVQADLKSDSGALDDLTLDSLDLDAASTAADK
DELDSFSLDDL SLDDSGHAAK PDAAGQDLDDAFDLSLDDLGGGDLGSDDVQADLKSDSGALDDLTLDSLDLDAASTAADK
DELDSFSLDDL SLDDSGHAAK PDAAGQDLDDAFDLSLDDLGG.....DDVQADLKSDSGALDDLTLDSLDLDAASTAADK
!!

720
715

logo

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PVDDLDFGLDFAELAE TSPQPKHDDL GDFSLDL DAPEDKLSDDDFLLSLNDEVPAAPADNEFTLDTEAAEEPALSLPDD
PVDDLDFGLDFAELAE TSPQPKHDDL GDFSLDL DAPEDKLSDDDFLLSLNDEVPAAPADNEFTLDTEAAEEPALSLPDD
PVDDLDFGLDFAELAE TSPQPKHDDL GDFSLDL DAPEDKLSDDDFLLSLNDEVPAAPADNEFTLDTEAAEEPALSLPDD
!!

800
795

logo

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FDLSLADEPTPEAAPEKGEDSF AAQLDEVSAQLDELASNLD EPKSAATPSFSAEDA AVASALDGDADDDDFDLSGADEAAT
FDLSLADEPTPEAAPEKGEDSF AAQLDEVSAQLDELASNLD EPKSAATPSFSAEDA AVASALDGDADDDDFDLSGADEAAT
FDLSLADEPTPEAAPEKGEDSF AAQLDEVSAQLDELASNLD EPKSAATPSFSAEDA AVASALDGDADDDDFDLSGADEAAT
!!

880
875

logo

NTDB id 275447 CSB90 RS25110 WP 003130552.1
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consensus

KLDLARAYIDMGDSEGARDILDEVL AEGNDSQQAEARELLERLA
KLDLARAYIDMGDSEGARDILDEVL AEGNDSQQAEARELLERLA
KLDLARAYIDMGDSEGARDILDEVL AEGNDSQQAEARELLERLA
!!

924
919

⊠ non conserved
⊠ similar
⊠ ≥ 50% conserved