

logo

NTDB id 385532 F1C79 RS14115 WP 151187796.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



63
80

logo

NTDB id 385532 F1C79 RS14115 WP 151187796.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



137
160

logo

NTDB id 385532 F1C79 RS14115 WP 151187796.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



217
239

logo

NTDB id 385532 F1C79 RS14115 WP 151187796.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



289
319

logo

NTDB id 385532 F1C79 RS14115 WP 151187796.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



363
399

logo

NTDB id 385532 F1C79 RS14115 WP 151187796.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



380
479

logo

NTDB id 385532 F1C79 RS14115 WP 151187796.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus



380
559

logo

NTDB id 385532 F1C79 RS14115 WP 151187796.1
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consensus

FNQAAELLQNAIYDEPQRDLRLKLMEVYAEMGDREGFARQENELREI GGAQPQVEQLKSRY PAMVAVAAVAGLAGAKLA
.....
FNQAAELLQNAIYDEPQRDLRLKLMEVYAEMGDREGFARQENELREI GGAQPQVEQLKSRY PAMVAVAAVAGLAGAKLA

380
639

logo

NTDB id 385532 F1C79 RS14115 WP 151187796.1
NTDB id 1196 PAKAF RS09365 WP 016253171.1
consensus

QDELDSFSLDDL SLDLDSGHAAK PDAAGQDLDDAFDLSLDDLGGDDVQADLKSDSGALDDLTLDSDLDLAASTAADKPVDD
.....
QDELDSFSLDDL SLDLDSGHAAK PDAAGQDLDDAFDLSLDDLGGDDVQADLKSDSGALDDLTLDSDLDLAASTAADKPVDD

380
719

logo

NTDB id 385532 F1C79 RS14115 WP 151187796.1
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consensus

LDFGLDFAEL AETPSQPKHDDL GDFSLDL DAPEDKLSDDDFLLSLNDEV PAAAPADNEFTLDTEAAEEPALSLPDDFDLS
.....
LDFGLDFAEL AETPSQPKHDDL GDFSLDL DAPEDKLSDDDFLLSLNDEV PAAAPADNEFTLDTEAAEEPALSLPDDFDLS

380
799

logo

NTDB id 385532 F1C79 RS14115 WP 151187796.1
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consensus

LADEPTEPAAPEKGEDSFAAQLDEVSAQLDELASNLDEPKSATPSFSAEDA AVASALDGDADDDFDLSGADEAATKLDL
.....
LADEPTEPAAPEKGEDSFAAQLDEVSAQLDELASNLDEPKSATPSFSAEDA AVASALDGDADDDFDLSGADEAATKLDL

380
879

logo

NTDB id 385532 F1C79 RS14115 WP 151187796.1
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consensus

ARAYIDMGDSEGARDILDEVLAEGNDSQQAEARELLERLA
.....
ARAYIDMGDSEGARDILDEVLAEGNDSQQAEARELLERLA

380
919

- X non conserved
- X similar
- X ≥ 50% conserved