

logo

NTDB id 243368 CGP88 RS08310 WP 001566051.1
NTDB id 22 SA RS07995 WP 001557362.1
NTDB id 38 MW RS08220 WP 001566670.1
consensus



80
80
80

logo

NTDB id 243368 CGP88 RS08310 WP 001566051.1
NTDB id 22 SA RS07995 WP 001557362.1
NTDB id 38 MW RS08220 WP 001566670.1
consensus



160
160
160

logo

NTDB id 243368 CGP88 RS08310 WP 001566051.1
NTDB id 22 SA RS07995 WP 001557362.1
NTDB id 38 MW RS08220 WP 001566670.1
consensus



240
240
240

logo

NTDB id 243368 CGP88 RS08310 WP 001566051.1
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consensus



320
320
320

logo

NTDB id 243368 CGP88 RS08310 WP 001566051.1
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consensus



400
400
400

logo

NTDB id 243368 CGP88 RS08310 WP 001566051.1
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consensus



480
480
480

logo

NTDB id 243368 CGP88 RS08310 WP 001566051.1
NTDB id 22 SA RS07995 WP 001557362.1
NTDB id 38 MW RS08220 WP 001566670.1
consensus

LYEGGKNQNLIDTGGKVIDDTKQPSYSISKYHILPTLNERGINELEYLILTHPHNDHIGEVEYIISHIKIKHI^EIYNKG

560
560
560

logo

NTDB id 243368 CGP88 RS08310 WP 001566051.1
NTDB id 22 SA RS07995 WP 001557362.1
NTDB id 38 MW RS08220 WP 001566670.1
consensus

YSSNTLMLLSKLSHKYNIKLM¹DVRQVSSFKLGDSSFLFFDSFIPNSRDKNEYSIITMITYQNKVKVLLMGDASKNNESLLL
YSSNTLMLLSKLSHKYNIKLM²DVRQVSSFKLGDSSFLFFDSFIPNSRDKNEYSIITMITYQNKVKVLLMGDASKNNESLLL 640
YSSNTLMLLSKLSHKYNIKLM³DVRQVSSFKLGDSSFLFFDSFIPNSRDKNEYSIITMITYQNKVKVLLMGDASKNNESLLL 640
YSSNTLMLLSKLSHKYNIKLM⁴DVRQVSSFKLGDSSFLFFDSFIPNSRDKNEYSIITMITYQNKVKVLLMGDASKNNESLLL 640

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consensus

KKYNLPEIDILKVGHHGSKTSSSKEFIEMIKPKISLISSGKNMYHLPNIEVKRLQRIRSRINYNSQQNGQVTIDLDDNL
720
KKYNLPEIDILKVGHHGSKTSSSKEFIEMIKPKISLISSGKNMYHLPNIEVKRLQRIRSRINYNSQQNGQVTIDLDDNL
720
KKYNLPEIDILKVGHHGSKTSSSKEFIEMIKPKISLISSGKNMYHLPNIEVKRLQRIRSRINYNSQQNGQVTIDLDDNL
720

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consensus



-  non conserved
-  similar
-  $\geq 50\%$ conserved