

logo

NTDB id 315764 TFGB2 RS02195 WP 134797923.1
NTDB id 1132 OK783 RS01905 WP 010951035.1
consensus



80
80

logo

NTDB id 315764 TFGB2 RS02195 WP 134797923.1
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consensus



160
160

logo

NTDB id 315764 TFGB2 RS02195 WP 134797923.1
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consensus



240
240

logo

NTDB id 315764 TFGB2 RS02195 WP 134797923.1
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consensus



320
320

logo

NTDB id 315764 TFGB2 RS02195 WP 134797923.1
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consensus



400
400

logo

NTDB id 315764 TFGB2 RS02195 WP 134797923.1
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consensus



480
480

logo

NTDB id 315764 TFGB2 RS02195 WP 134797923.1
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consensus



560
560

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QQLHDFNQNESEILHWAESARTALDIWQKYGIFAAMQQFSQTHGIETRLLSRNNGRSLTNYFQELLELLAAEDAQNRPAA
 QQLHDFNQNESEILHWAESARTALDIWQKYGIFAAMQQFSQTHGIETRLLSRNNGRSLTNYFQELLELLAAEDAQNRPAA
 QQLHDFNQNESEILHWAESARTALDIWQKYGIFAAMQQFSQTHGIETRLLSRNNGRSLTNYFQELLELLAAEDAQNRPAA

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LHKWLRDQISLAGNNGGDNRAIRLESDDELVKIVTMHASKGLQYPLVYCPFAWDAQDTGPSDWQILHQQGANRAELLAQAQ 800
LHKWLRDQISLAGNNGGDNRAIRLESDDELVKIVTMHASKGLQYPLVYCPFAWDAQDTGPSDWQILHQQGANRAELLAQAQ 800

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LSESEKQYADEEMAERLRLLYVALTRAEQNLNIAAYSSDTADNPLAYLIEGLPQDSRETVRRTYACEKDGITMLKRNW
 LSESEKQYADEEMAERLRLLYVALTRAEQNLNIAAYSSDTADNPLAYLIEGLPQDSRETVRRTYACEKDGITMLKRNW 880
 LSESEKQYADEEMAERLRLLYVALTRAEQNLNIAAYSSDTADNPLAYLIEGLPQDSRETVRRTYACEKDGITMLKRNW 880

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RRVADVAPAGTDFATFTEDAPPPAAAYRGNADQAAEFAANSIPERGFRFVRHTSFTALSRQTQTPDGGEEDACPSLDAAETS
 RRVADNAPAGTDFATFTEDAPPPAAAYRGNADQAAEFAANSIPERGFRFVRHTSFTALSRQTQTPDGGEEDACPSLDAAETS 960
 RRVADNAPAGTDFATFTEDAPPPAAAYRGNADQAAEFAANSIPERGFRFVRHTSFTALSRHTQTPDGGEEDACPSLDAAETS 960

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VPAMPSETPTASDGLSIHDFPKGTQAGLCLHEILEDFKFGQAAAEQETLIADKLKKYGFEEIWLPAVAEMA EACRKTPLT
 VPAMPSETPTASDGLSIHDFPKGTQAGLCLHEILEDFKFGQAAAEQETLIADKLKKYGFEEIWLPAVAEMA EACRKTPLT 1040
 VPAMPSETPTASDGLSIHDFPKGTQAGLCLHEILEDFKFGQAAAEQETLIADKLKKYGFEEIWLPAVAEMA EACRKTPLT 1040

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GAYGLSDIPPGCRRPEMGFTLHTEDFGLKRLRDWLARDDIRLPEVCRAAAETLDFHTVNGFLNGFIDMVCQDPDGNICII 1120
 GAYGLSDIPPGCRRPEMGFTLHTEDFGLKRLRDWLARDDIRLPEVCRAAAETLDFHTVNGFLNGFIDMVCQDPDGNICII 1120

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consensus



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