logo



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NTDB id 1015 ACIAD RS12005 WP 004928865.1  MFKIILLGWIAGIAMMGRTLECIEPFTCITGILTFLIWAIYFYKKRKHQIGSGLRALVILISASFLFGIGAQYADSALEERLYYK  85
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NTDB id 47498 H7K53 RS14600 WP 000472707.1  MFKIILLGWIGGIALMGIDFPLIMQYEKVGEALLLLAFIFYLYKRPMFVDRPFLKAVFCLLCTTSLFMVGYHYAEKALIERLEQR  85
NTDB id 1070 ABD1 RS13145 WP 000472713.1  MFKIILLGWIGGIALMGIDFPLIMQYEKVGEALLLLAFIFYLYKRPMFVDRPFLKAVFCLLCTTSLFMVGYHYAEKALIERLEQR  85
consensus                                  !!!!!!!!!!*!!!*!!*****!*********!**!****!*!!*********!*!***!*****!!**!**!!**!!*!!!***
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NTDB id 47498 H7K53 RS14600 WP 000472707.1  ETDTRNLDIIVYINRLSEEKDNKVQQTAQVLNFSKEPVNWLLYLKSNNQNLLKNNQNLELGHYYRISGKTRPAHSYATPGAFDQE  170
NTDB id 1070 ABD1 RS13145 WP 000472713.1  ETDTRNLDIIVYINRLSEEKDNKVQQTAQVLNLSKEPVNWLLYLKSNNQNLLKNNQNLELGHYYRISGKTRPAHSYATPGAFDQE  170
consensus                                  *********!*!*****!**!****!*!*!!!*****!!!**!***************!!!!*!!!!**!*!*!!!!**!*!!*!
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NTDB id 47498 H7K53 RS14600 WP 000472707.1  KWFIQRNIMSGFNVRYIEPLSLDEIYRLGYQQHLKEQQSFSNSFRLNIEKLRLTFRQILNSSSLQQKGLILALLTGDESLLSDET  255
NTDB id 1070 ABD1 RS13145 WP 000472713.1  KWFIQRNIMSGFNVRYIEPLSLDEIYRLGYQQHLKEQQSFSNSFRLNIEKLRLTFRQILNSSSLQQKGLILALLTGDESLLSDET  255
consensus                                  !!!*!*!******!**!**!!**!****!***!*!*!!*****!****!!!!!*!!*****!!!****!*!!!!!!!*!!!**!*
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NTDB id 1083 HON27 RS04190 WP 168726476.1  QLQFKQLGISHLLAISGPHVFIFAIMLSWACHQFISRYYPKIYLWKPKQILMAVPCCLGVLIYTAFVGFEIPALRTLLSTLIFIG  340
NTDB id 47498 H7K53 RS14600 WP 000472707.1  QLQFKQLGISHLLAISGPHVLIFAIMLSWACHQFISRYYPQIYLWKPKQVLMTVPCCLGVLIYTAFVGFEIPALRTLLSALIFIG  340
NTDB id 1070 ABD1 RS13145 WP 000472713.1  QLQFKQLGISHLLAISGPHVLIFAIMLSWAYHQFISRYYPQIYLWKPKQVLIAVPCCLGVLIYTAFVGFEIPALRTFLSVLIFIG  340
consensus                                  **!!***!!!!!!!!!!!!!*!!!*!*!!*****!*!**!*!!!**!*!*!***!****!**!*!!!!!!!!!!!*********
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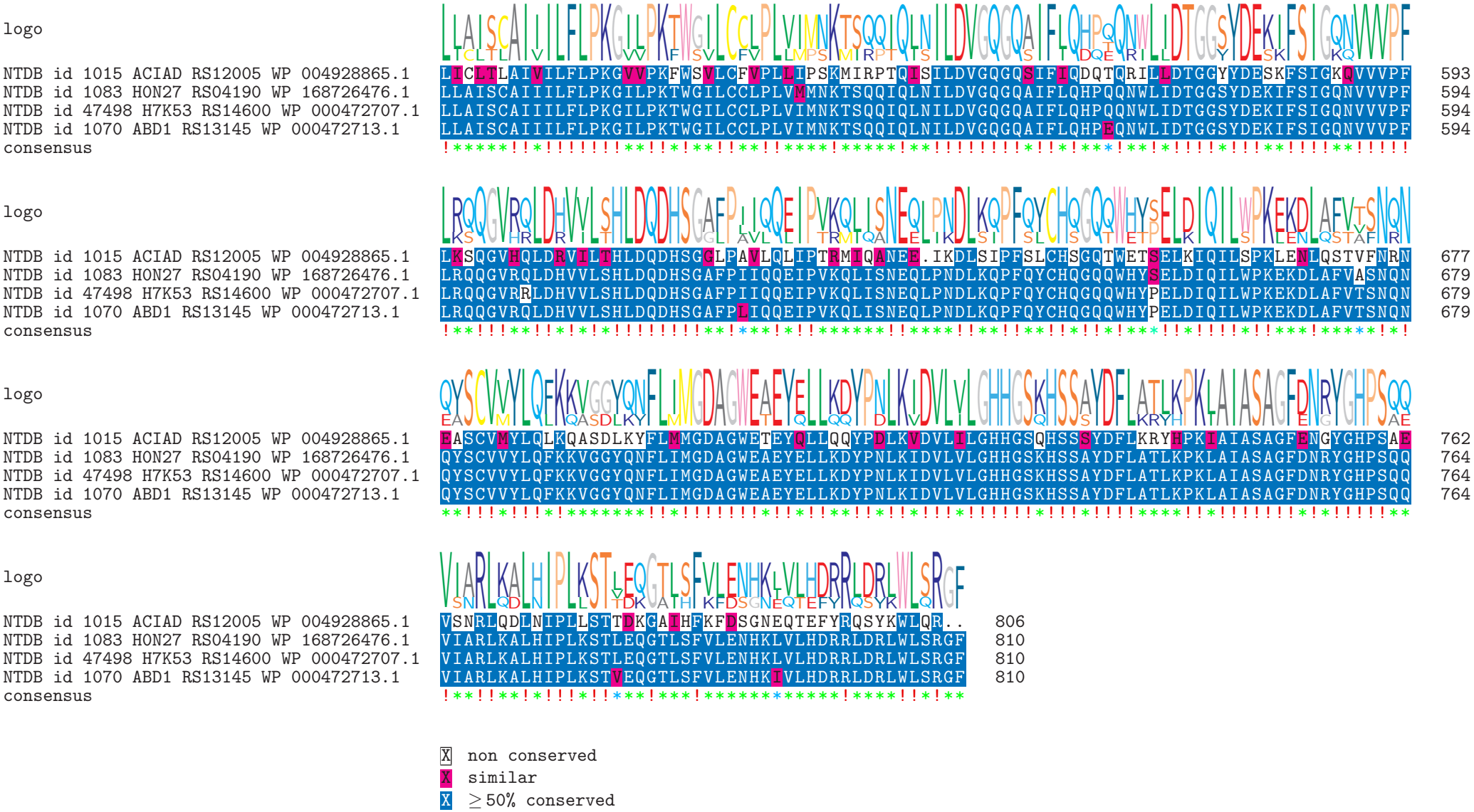


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NTDB id 1083 HON27 RS04190 WP 168726476.1  FLLLKQPIKPFTLLVYSASLLLLMDPFSVLSAGFWLSYGACFILIRIYQTIAQLPEQHFLSLSSKMIFMSKVLIESQGKIFIALS  425
NTDB id 47498 H7K53 RS14600 WP 000472707.1  FLLLKQPIKPFTLLVYSASLLLLIDPFSVLSAGFWLSYGACFILLRIYQTIAQLPEQHFLSLSSKMIFMGKVLIESQGKIFIALS  425
NTDB id 1070 ABD1 RS13145 WP 000472713.1  FLLLKQPIKPFTLLVYSASLLLLMDPFSVLSAGFWLSYGACFILLRIYQTIAQLPEQHFLSLSSKMIFMSKVLIESQGKIFIALS  425
consensus                                  *!!*******!*!!!*!!!!!!**!!!!*!!!*!!!!!!!!!!!!*!!!!***!****!*******!****!*!**!!*!!!*!!*
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consensus                                  !!!!!**!**!!!*!!!*!*!!*!!*! *!!!!*!*!*!*!******!!*!!**!**!!******!**!**!  ***!!****!!!*
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consensus
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consensus
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consensus
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NTDB id 47498 H7K53 RS14600 WP 000472707.1 VIARLKALHIPLKSTLEQGTLSFVLENHKLVLHDRRLDRLWLSRGF 810
NTDB id 1070 ABD1 RS13145 WP 000472713.1 VIARLKALHIPLKSTVEQGTLSFVLENHKIVLHDRRLDRLWLSRGF 810
consensus
X non conserved
X similar
X ≥ 50% conserved