logo

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NTDB id 1085586 ACLBNB RS09975 WP 124310675.1  MRTGMLALAAGLLSLRFLPALPALGWLAVLPVVALMLLPFRTYALALFLLGLTWACVSAQWALDDRLPEPLDGQTRWLEG  80
NTDB id 1199 PSJM300 12650 AFN78592.1          MRAGMLALAAGLLLLRFLPQLPSFGWLLALAPVGLLLLPFRWRPLAFFLFGLVWACLNAQWALDDRLPVDLDGRTFWLEG  80
consensus                                      !! !!!!!!!!!! !!!!! !!  !!!  !**!*!*!!!!!  *!! !! !! !!!* !!!!!!!!!! *!!! ! !!!!
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logo

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NTDB id 1085586 ACLBNB RS09975 WP 124310675.1  RVVGLPQQAEGVLRFELEGGRARRGRIPQRLRLAWYGGPPLHSGERWRLAVKLKRPAGLVNLQAFDHEAWLLAQRIGATG  160
NTDB id 1199 PSJM300 12650 AFN78592.1          QVTGLPDRRGDVVRFELEDIHSRHAGLPSRIRLAWYGGPEIRSGERWRLAARLKRPSGMVNPSAFDYEAWLLARRIGATG  160
consensus                                       ! !!!      !*!!!!!  * !** *! !*!!!!!!!!***!!!!!!!! *!!!! !*!!* !!!*!!!!!! !!!!!!
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logo

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NTDB id 1085586 ACLBNB RS09975 WP 124310675.1  TVKDGQLLV..AAEGGWRARIRQRLLQVDAHGRAGGLAALVLGDGSGLSREDWRVLQDTGTVHLMVISGQHISLLAAVMY  238
NTDB id 1199 PSJM300 12650 AFN78592.1          TIKAGERIAEAASSGAWRDRLRQRLLTVEAHGRAGAIAALVVGDGSGLSTADWRVLQDTGTVHLMVISGQHVGMLAGLLY  240
consensus                                      !*!  !   * **!  !*!! !*!!!!! !*!!!!!!**!!!!*!!!!!!! !!!!!!!!!!!!!!!!!!!!* *!!***!
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NTDB id 1085586 ACLBNB RS09975 WP 124310675.1  VLVAGLARYGLWPGRWPWLPWACGLAFASALGYGLLAGFDVPVRRACVMVALVLIWRLRFRHLGAWWPFLLALNTVLLFE  318
NTDB id 1199 PSJM300 12650 AFN78592.1          AVVLLLARWGIWPQRLPWLPWACGLAFAGALGYGWLAGFEVPVRRACLMVAIVLLWRLRFRHLGVWLPLLIALVGVLMVD  320
consensus                                       *!  !!!*!*!! ! !!!!!!!!!!!! !!!!! !!!!*!!!!!!!*!!!*!!*!!!!!!!!! ! ! !*!!  !!* *
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NTDB id 1085586 ACLBNB RS09975 WP 124310675.1  PLASLQPGFWLSFTAVAVLILIFGGRLGGWRWWQSWTRIQWLIAIGLCPVLLALGLPISLSGPLANLLAEPWISLLVLPP  398
NTDB id 1199 PSJM300 12650 AFN78592.1          PLASLQPGFWLSFVAVALLIWIFRGRLGVPSWWRALLYAQWAMALGLLPAMLVLGLPVSLSGPLANLLAVPWISVLVVPL  400
consensus                                      !!!!!!!!!!!!!! !!!*!! !! !!!! * !!          !! *!*!!*! *! !!!!*!!!!!!!!!! !!!!*!!*!*
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NTDB id 1085586 ACLBNB RS09975 WP 124310675.1  ALLGTLLLAVPYLGEGLLWLAGGLLDVLFRGLALLAAQWPAWVAPAIPLWAWSISALGAFLLLLPAGVPLRPLGWPMLLL  478
NTDB id 1199 PSJM300 12650 AFN78592.1          ALLGSFLLWVPWLGEALLWLAGGLLAVLFELLALIAGWQPAWLASALPLWAWSLVALGTLVLLLPAGVPLRVLGTLLLAP  480
consensus                                      !!!!* !! !!*!!!*!!!!!!!!! !!!  !!!*!* !!!*!*!*!!!!!!* !!!  *!!!!!!!!!!*!! **! *
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logo

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NTDB id 1085586 ACLBNB RS09975 WP 124310675.1  MVFPPKNTLPYGEAEVWQLDVGQGLAVLVRTREHAMLYDAGPRLGDSDLGARVVLPSLRKLGVERLDLMLLSHADADHAG  558
NTDB id 1199 PSJM300 12650 AFN78592.1          LLFAPDSRPEQGRAEVWVFDVGQGLSVLVRTRDHALLYDAGPRYGDFDIGERVVFPSLRQLGLERLDLMMLSHADSDHAG  560
consensus                                      **!*!    ** ! !!!!  !!!!!! !!!!!!*!!*!!!!!!! !! !*! !!! !!!! !!*!!!!!!*!!!!! !!!!
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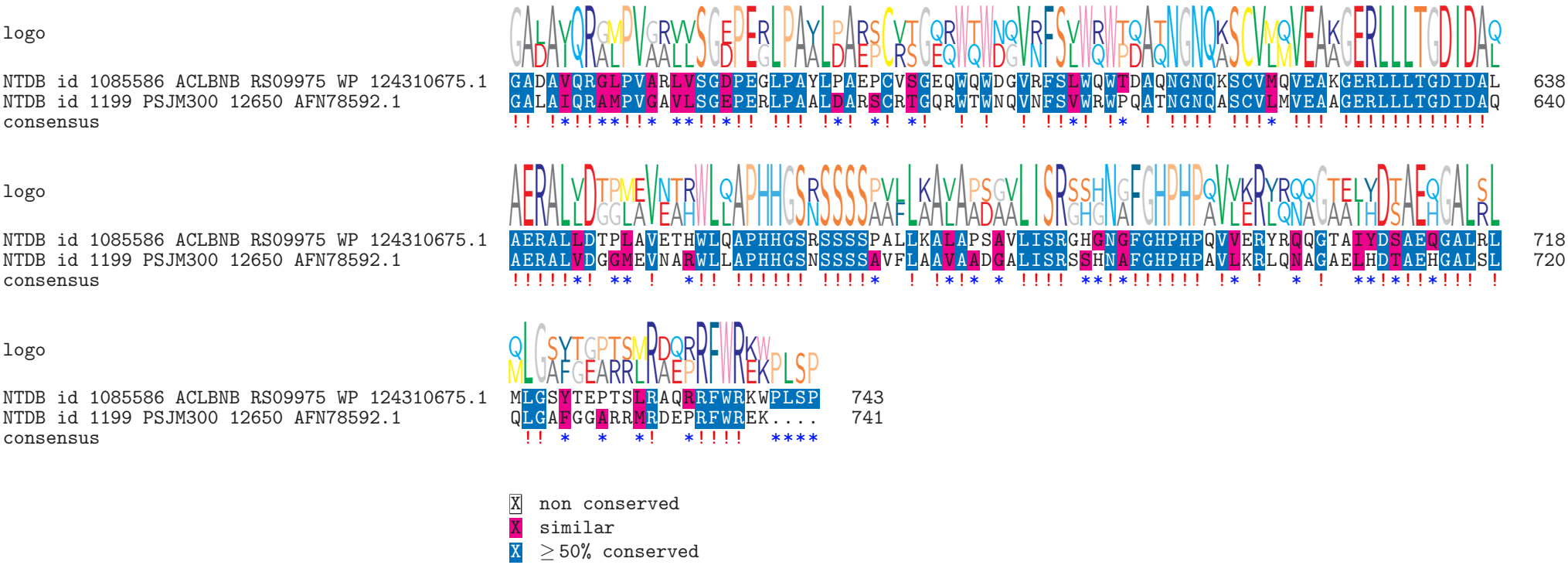
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NTDB id 1199 PSJM300 12650 AFN78592.1 GALAIQRAMPVGAVLSGEPERLPAALDARSCRTGQRWTWNQVNFSVWRWPQATNGNQASCVLMVEAAGERLLLTGDIDAQ 640
consensus
logo
NTDB id 1085586 ACLBNB RS09975 WP 124310675.1 AERALLDTPLAVETHWLQAPHHGSRSSSSPALLKALAPSAVLISRGHGNGFGHPHPQVVERYRQQGTAIYDSAEQGALRL 718
NTDB id 1199 PSJM300 12650 AFN78592.1 AERALVDGGMEVNARWLLAPHHGSNSSSSAVFLAAVAADGALISRSSHNAFGHPHPAVLKRLQNAGAELHDTAEHGALSL 720
consensus
logo
NTDB id 1085586 ACLBNB RS09975 WP 124310675.1 MLGSYTEPTSLRAQRRFWRKWPLSP 743
NTDB id 1199 PSJM300 12650 AFN78592.1 QLGAFGGARRMRDEPRFWREK.... 741
consensus
X non conserved
X similar
X ≥ 50% conserved