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logo

NTDB id 79421 IGS56 RS04915 WP 103708077.1  MKLNLITLAVLLIVADLTLLFLPQPLLLPWQVALVIALVLIFLFIFLRRNFLVSLAFFVASLGYFHYSALSLLQQAQNITAQKQV   85
NTDB id 1359 HI 0061 AAC21739.1             MKLNLITLVVLLIVADLTLLFLPQPLLLPWQVALVIALVLIFLFIFLRRNFLVSLAFFVASLGYFHYSALSLSQQAQNITAQKQV   85
consensus                                   !!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 79421 IGS56 RS04915 WP 103708077.1  VTFKIQEILHQQDYQTLIATATLANNLQEQRIFLNWKAKEVPQLSEIWQAEISLRPLSARLNFGGFDRQQWYFSKGITAVGTVKS   170
NTDB id 1359 HI 0061 AAC21739.1             VTFKIQEILHQQDYQTLIATATLENNLQEQRIFLNWKAKEVPQLSEIWQAEISLRSLSARLNFGGFDRQQWYFSKGITAVGTVKS   170
consensus                                   !!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 79421 IGS56 RS04915 WP 103708077.1  AVKIADVSSLRAEKLQQVKKQTEGLSLQGLLIALAFGERAWLDKTTWSIYQQTNTAHLIAISGLHIGLAMGIGFCLARVVQVFFP   255
NTDB id 1359 HI 0061 AAC21739.1             AVKIADVSSLRAEKLQQVKKQTEGLSLQGLLIALAFGERAWLDKTTWSIYQQTNTAHLIAISGLHIGLAMGIGFCLARVVQVFFP   255
consensus                                   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 79421 IGS56 RS04915 WP 103708077.1  TRFIHPYFPLVFGVLFALIYAYLAGFSVPTFRAISALVFVFFVQIMRRHYSPLQLFTLVVGFLLFCDPLMPLSVSFWLSCGAVGC   340
NTDB id 1359 HI 0061 AAC21739.1             TRFIHPYFPLVFGVLFALIYAYLAGFSVPTFRAISALVFVLFIQIMRRHYSPIQFFTLVVGFLLFCDPLMPLSVSFWLSCGAVGC   340
consensus                                   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !*!!!!!!!!!*! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 79421 IGS56 RS04915 WP 103708077.1  LLLWYRYVPFSLFQWKNRPFSPKVRWILSLFHLQFGLLLFFTPLQLFLFNGLSLSGFLANLMAVPIYSFLLVPLILFAVFTNGTM   425
NTDB id 1359 HI 0061 AAC21739.1             LLLWYRYVPFSLFQWKNRPFSPKVRWIFSLFHLQFGLLLFFTPLQLFLFNGLSLSGFLANFMAVPIYSFLLVPLILFAVFTNGTM   425
consensus                                   !!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 79421 IGS56 RS04915 WP 103708077.1  FSWQLANKLAEGITGLISVFQGNWFNVSFNLALVLTALCAGIFMLIIWSIYREPEVSSSTWKIKRPRFFTLNLSKPLLKTDRINV   510
NTDB id 1359 HI 0061 AAC21739.1             FSWQLANKLAEGITGLISVFQGNWLTVSFNLALGLTALCAGIFMLIIWNIYREPEISSSNWQIKRAKFFTLNLSKPLLKNERINV   510
consensus                                   !!!!!!!!!!!!!!!!!!!!!!!!   !!!!!!! !!!!!!!!!!!!!!! !!!!!!*!!!! ! !!!**!!!!!!!!!!!!! *!!!!

logo

NTDB id 79421 IGS56 RS04915 WP 103708077.1  LRCSFGIILMCFMILLFKQLSKPTWQVDTLDVGQGLATLIVKNGKGILYDTGSSWRGGSMAELEILPYLQREGIVLEKLILSHDD   595
NTDB id 1359 HI 0061 AAC21739.1             LRCSFGIILLCFTILLFKQLSKPTWQVDTLDVGQGLATLIVKNGKGILYDTGSSWRGGSMAELEILPYLQREGIVLEKLILSHDD   595
consensus                                   !!!!!!!!!*!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
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logo                                             NDHAGGASTILKAYPNVELITPSRKNYGENYRTFCTAGRDWHWQGLHFQILSPHNVVTRADNSHSCVILVDDGKNSVLLTGDAEA

NTDB id 79421 IGS56 RS04915 WP 103708077.1        NDHAGGASTILKAYPNVELITPSRKNYGENYRTFCTAGRDWHWQGLHFQILSPHNVVTRADNPHSCVILVDDGKNNVLLTGDAEA   680
NTDB id 1359 HI 0061 AAC21739.1                   NDHAGGASTILKAYPNVELITPSRKNYGENYRTFCTAGRDWHWQGLHFQILSPHNVVTRADNSHSCVILVDDGKNSVLLTGDAEA   680
consensus                                        !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!! !!!!!!!!!

logo                                             KNEQIFARTLGKIDVLQVGHHGSKTSTSEYLLSQVRPDVAIISSGRWNPWKFPHYSVMERLHRYKSAVENTAVSGQVRVNFFQDR

NTDB id 79421 IGS56 RS04915 WP 103708077.1        KNEQIFARTLGKIDVLQVGHHGSKTSTSEYLLSQVRPDVAIISSGRWNPWKFPHYSVMERLHRYKSAVENTAVSGQVRVNFFQDR   765
NTDB id 1359 HI 0061 AAC21739.1                   KNEQIFARTLGKIDVLQVGHHGSKTSTSEYLLSQVRPDVAIISSGRWNPWKFPHYSVMERLHRYKSAVENTAVSGQVRVNFFQDR   765
consensus                                        !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo                                             LEIQQARTKFSPWYARVIGLSKE

NTDB id 79421 IGS56 RS04915 WP 103708077.1        LEIQQARTKFSPWYARVIGLSKE   788
NTDB id 1359 HI 0061 AAC21739.1                   LEIQQARTKFSPWYARVIGLSKE   788
consensus                                        !!!!!!!!!!!!!!!!!!!!!!!
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X non conserved

X similar

X ≥ 50% conserved