

logo

NTDB id 150655 WX61 RS08525 WP 154236097.1
NTDB id 1128 NGFG RS01490 WP 020997305.1
consensus

85
85

logo

NTDB id 150655 WX61 RS08525 WP 154236097.1
NTDB id 1128 NGFG RS01490 WP 020997305.1
consensus

TDMPRSDGRRVQFAAKAVDSGGRTFDLLLSDYKRREWAVGSRWRITARVHPVVGELNLRGLNREAWALSNGVGGVGTVGADRVLL
 TDMPRSDGRRVQFAAKAVDSGGRTFDLLLSDYKRREWAVGSRWRITARVHPVVGELNLRGLNREAWALSNGVGGVGTVGADRVLL
 TDMPRSDGRRVQFAAKAVDSGGRTFDLLLSDYKRREWAVGSRWRITARVHPVVGELNLRGLNREAWALSNGVGGVGTVGADRVLL

170
170

logo

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consensus

HGGSGWGIAVWRSRISRNWRQADADGGLSDGIGLMRALSVGEQSALRPGLWQAFRPLGLTHLVSISGLHVTMVAVLFAWLAKRLL
HGGSGWGIAVWRSRISRNWRQADADGGLSDGIGLMRALSVGEQSALRPGLWQAFRPLGLTHLVSISGLHVTMVAVLFAWLAKRLL
HGGSGWGIAVWRSRISRNWRQADADGGLSDGIGLMRALSVGEQSALRPGLWQAFRPLGLTHLVSISGLHVTMVAVLFAWLAKRLL

255
255

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ACSPRLPARPRAWLAAGCAGALFYALLAGFSVPTQRSVLMFAFAWARRGRLSAWATWWQALAAVLLFDPLAVLVGVTWLSFG
ACSPRLPARPRAWLAAGCAGALFYALLAGFSVPTQRSVLMFAFAWARRGRLSAWATWWQALAAVLLFDPLAVLVGVTWLSFG
ACSPRLPARPRAWLAAGCAGALFYALLAGFSVPTQRSVLMFAFAWARRGRLSAWATWWQALAAVLLFDPLAVLVGVTWLSFG

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340

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LVAALIWACAGRLYEGRQTAVRGQWAASVLSVLVLLGYLFASLPLVSPVLNAVSIPIWFSWVLTPLALLGSVVPFAPLQQAGAFLA
 LVAALIWACAGRLYEGRQTAVRGQWAASVLSVLVLLGYLFASLPLVSPVLNAVSIPIWFSWVLTPLALLGSVVPFAPLQQAGAFLA
 LVAALIWACAGRLYEGRQTAVRGQWAASVLSVLVLLGYLFASLPLVSPVLNAVSIPIWFSWVLTPLALLGSVVPFAPLQQAGAFLA

425
425

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EYTLRFLVWVLADVSPEFAVAAAAPLLVLAVCAALLLLLPRGLGLRPWAVLLLAGFVSYPPEAVPENEAAVTVMDAGQGLSVLVR
 EYTLRFLVWVLADVSPEFAVAAAAPLLVLAVCAALLLLLPRGLGLRPWAVLLLAGFVSYPPEAVPENEAAVTVMDAGQGLSVLVR
 EYTLRFLVWVLADVSPEFAVAAAAPLLVLAVCAALLLLLPRGLGLRPWAVLLLAGFVSYPPEAVPENEAAVTVMDAGQGLSVLVR

510
510

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TANRHLLFDTGTVAAAQTGIVPSLNAAGVRRLDKLVLSHHDSHDGGFQAVGKIPNGGIYAGQPEFYEGARHCAEQRWQWDGVDF
TANRHLLFDTGTVAAAQTGIVPSLNAAGVRRLDKLVLSHHDSHDGGFQAVGKIPNGGIYAGQPEFYEGARHCAEQRWQWDGVDF
TANRHLLFDTGTVAAAQTGIVPSLNAAGVRRLDKLVLSHHDSHDGGFQAVGKIPNGGIYAGQPEFYEGARHCAEQRWQWDGVDF

595
595

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consensus



680
680

logo

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consensus



739
739

-  non conserved
-  similar
-  ≥ 50% conserved