

logo

NTDB id 50055 HT085 RS01785 WP 154236097.1
NTDB id 1128 NGFG RS01490 WP 020997305.1
consensus

85
85

logo

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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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ACSPRLPARPRAWLAAGCAGALFYALLAGFSVPTQRSVLMFAAWRRGRLSAWATWWQALAAVLLFDPLAVLGVGTWLSFG
ACSPRLPARPRAWLAAGCAGALFYALLAGFSVPTQRSVLMFAAWRRGRLSAWATWWQALAAVLLFDPLAVLGVGTWLSFG
ACSPRLPARPRAWLAAGCAGALFYALLAGFSVPTQRSVLMFAAWRRGRLSAWATWWQALAAVLLFDPLAVLGVGTWLSFG

340
340

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LVAALIWACAGRLYEGKRQTAVRGQWAASVLSVLVLLGYLFASLPLVSPLVNAVSIPIWFSWVLTPLALLGSVVPFAPLQQAGAFLA
LVAALIWACAGRLYEGKRQTAVRGQWAASVLSVLVLLGYLFASLPLVSPLVNAVSIPIWFSWVLTPLALLGSVVPFAPLQQAGAFLA
LVAALIWACAGRLYEGKRQTAVRGQWAASVLSVLVLLGYLFASLPLVSPLVNAVSIPIWFSWVLTPLALLGSVVPFAPLQQAGAFLA

425
425

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EYTLRFLVWLADVSPFEFAVAAPLPLLVLAACAALLLLLPRGLGLRPWAVLLLAGFVSYPPEAVPENEAAVTVMDAGQGLSVLVR
 EYTLRFLVWLADVSPFEFAVAAPLPLLVLAACAALLLLLPRGLGLRPWAVLLLAGFVSYPPEAVPENEAAVTVMDAGQGLSVLVR
 EYTLRFLVWLADVSPFEFAVAAPLPLLVLAACAALLLLLPRGLGLRPWAVLLLAGFVSYPPEAVPENEAAVTVMDAGQGLSVLVR

510
510

logo

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TANRHLLFDTGTVAAAQTGIVPSLNAAGVRRLDKLVLSHHDSDHDGGFQAVGKIPNGGIYAGQPEFYEGARHCAEQRWQWDGVDF
TANRHLLFDTGTVAAAQTGIVPSLNAAGVRRLDKLVLSHHDSDHDGGFQAVGKIPNGGIYAGQPEFYEGARHCAEQRWQWDGVDF
TANRHLLFDTGTVAAAQTGIVPSLNAAGVRRLDKLVLSHHDSDHDGGFQAVGKIPNGGIYAGQPEFYEGARHCAEQRWQWDGVDF

595
595

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consensus



680
680

logo

NTDB id 50055 HT085 RS01785 WP 154236097.1
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consensus



739
739

-  non conserved
-  similar
-  ≥ 50% conserved