

logo

NTDB id 265302 C2H91 RS18050 WP 163136229.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

MRNSRLLLPMAAASATxGITAAAYFPAIFLFIILLLIILIKTRHAFLLVCFFSFILFFVLYAVTDSQNVSSYRQGTQF
MRNSRLLLPMAAASATVGITAAAYFPAIFLFIILLLIILIKTRHAFLLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTQF
MRNSRLLLPMAAASATAGITAAAYFPAIFLFIILLLIILIKTRHAFLLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTQF

80
80

logo

NTDB id 265302 C2H91 RS18050 WP 163136229.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

KAVIDTIPKIDGDRMSMMV^KTPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLLEEPNHATVPGAFDYNEYLYRQHI
 KAVIDTIPKIDGDRMSMMV^KTPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLLEEPNHATVPGAFDYNEYLYRQHI
 KAVIDTIPKIDGDRMSMMV^ETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLLEEPNHATVPGAFDYNEYLYRQHI

160
160

logo

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consensus

HNNYSVTSIQNCSEPENFKYKVLRLKHIISFTNSLLPPDSAGIVQALTVGDRFYVEDEVLTAYQKLGWHLLAISGLHV
 HWNYSVTSIQNCSEPENFKYKVLRLKHIISFTNSLLPPDSAGIVQALTVGDRFYVEDEVLTAYQKLGVVHLLAISGLHV
 HWNYSVTSIQNCSEPENFKYKVLRLKHIISFTNSLLPPDSTGIVQALTVGDRFYVEDEVLTAYQKLGVVHLLAISGLHV

240
240

logo

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320
320

logo

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consensus

YHLFEAGFQLSFAVSFSLILSSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSIISVPMNMLMVPFYTFCILPGA
YHLFEAGFQLSFAVSFSLILSSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSIISVPMNMLMVPFYTFCILPGA
YHLFEAGFQLSFAVSFSLILSSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSIISVPMNMLMVPFYTFCILPGA

400
400

logo

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consensus

VAGVLLLSLSASFGRLEFFSWFDLLYSHTNRLITNIADVDFVTIMIAHPAPALLFLFTVTIILLMAIEKRSLSQSMVTGG
VAGVLLLSLSASFGRLEFFSWFDLLYSHTNRLITNIADVDFVTIMIAHPAPALLFLFTVTIILLMAIEKRSLSQSMVTGG
VAGVLLLSLSASFGRLEFFSWFDLLYSHTNRLITNIADVDFVTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGG

480
480

logo

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consensus

560
560

logo

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consensus



640
640

logo

NTDB id 265302 C2H91 RS18050 WP 163136229.1
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consensus



720
720

logo

NTDB id 265302 C2H91 RS18050 WP 163136229.1
NTDB id 107 BSU 25570 NP 390435.1
consensus



776
776

 non conserved
 similar
 ≥ 50% conserved