

logo

NTDB id 340727 ETK71 RS12920 WP 041850028.1  
NTDB id 107 BSU 25570 NP 390435.1  
consensus

MRNSRLLLPMAAASATxGITAAAYFPAIFLFI LLLIILIKTRHAFIIVCFFSFI LFFVLYAVTDSQNVSSYRQGTYP  
MRNSRLLLPMAAASATV GITAAAYFPAIFLFI LLLIILIKTRHAFIIVCFFSFI LFFVLYAVTDSQNVSSYRQGTYP  
MRNSRLLLPMAAASATAGITAAAYFPAIFLFI LLLIILIKTRHAFIIVCFFSFI LFFVLYAVTDSQNVSSYRQGTYP

80  
80

logo

NTDB id 340727 ETK71 RS12920 WP 041850028.1  
NTDB id 107 BSU 25570 NP 390435.1  
consensus

KAVIATIPKIDGDRMSMMVKTPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPKHATVPGAFDNEYLYRQHI  
 KAVIATIPKIDGDRMSMMVKTPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPKHATVPGAFDNEYLYRQHI  
 KAVIDTIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDNEYLYRQHI

160  
160

logo

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consensus

HNNYSVTSIQNCSEPENFKYKVLRLKHIIISFTNSLLPPDSTGIVQALTVGDRFYVEDEVLTAYQKLGWHLAISGLHV  
 HWNYSVTSIQNCSEPENFKYKVLRLKHIIISFTNSLLPPDSTGIVQALTVGDRFYVEDEVLTAYQKLGWVHLLAISGLHV  
 HWNYSVTSIQNCSEPENFKYKVLRLKHIIISFTNSLLPPDSTGIVQALTVGDRFYVEDEVLTAYQKLGWVHLLAISGLHV

240  
240

logo

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consensus

320  
320

logo

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consensus

YHLFEAGFQLSFAVSFSLILSSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSISVPMNMLMVPFYTCILPGA  
YHLFEAGFQLSFAVSFSLILSSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSISVPMNMLMVPFYTCILPGA  
YHLFEAGFQLSFAVSFSLILSSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSISVPMNMLMVPFYTCILPGA

400  
400

logo

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VAGVLLLSLSASFGRFFSWFDLLISHTNRLIT<sub>K</sub>IADVDVFTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGG  
VAGVLLLSLSASFGRFFSWFDLLISHTNRLIT<sub>K</sub>IADVDVFTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGG  
VAGVLLLSLSASFGRFFSWFDLLISHTNRLIT<sub>N</sub>IADVDVFTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGG

480  
480

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consensus

560  
560

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consensus

640  
640

NTDB id 340727 ETK71 RS12920 WP 041850028.1  
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consensus

720  
720

NTDB id 340727 ETK71 RS12920 WP 041850028.1  
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consensus

776  
776

- X non conserved
- X similar
- X  $\geq 50\%$  conserved