

logo

NTDB id 339871 ES968 RS13005 WP 046160607.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

MRNSRLLLPMMAASATAGITAAAYFPAIFLFIILFLLIILIKTRHAFLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYPQ
MRNSRLLLPMMAASATAGITAAAYFPAIFLFIILFLLIILIKTRHAFLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYPQ
MRNSRLLLPMMAASATAGITAAAYFPAIFLFIILFLLIILIKTRHAFLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYPQ

80
80

logo

NTDB id 339871 ES968 RS13005 WP 046160607.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

KAVIDTIPKIDGRMSMMV^KTPDKEK^KWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDNEYLYRQHI
 KAVIDTIPKIDGRMSMMV^KTPDKEK^KWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDNEYLYRQHI
 KAVIDTIPKIDGRMSMMV^ETPDKEK^KWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDNEYLYRQHI

160
160

logo

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NTDB id 107 BSU 25570 NP 390435.1
consensus

HWNYSVTSIQNCSEPENFKYKVLRLKHIISFTNSLLPPDSTAGIVQALTVGDRFYVEDEVLTAYQKLGWVHLLAISGLHV
 HWNYSVTSIQNCSEPENFKYKVLRLKHIISFTNSLLPPDSTAGIVQALTVGDRFYVEDEVLTAYQKLGWVHLLAISGLHV

240
240

logo

NTDB id 339871 ES968 RS13005 WP 046160607.1
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consensus

320
320

logo

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NTDB id 107 BSU 25570 NP 390435.1
consensus

YHLFEAGFQLSFAVSFSLILSSSI^HFQVKTS^RQLTIVSLIAQLGSLPILLYYHFHQFS^IISVPMNMLMVPFYTCILPGA
 YHLFEAGFQLSFAVSFSLILSSSI^HFQVKTS^RQLTIVSLIAQLGSLPILLYYHFHQFS^IISVPMNMLMVPFYTCILPGA
 YHLFEAGFQLSFAVSFSLILSSSI^HFQVKTS^RQLTIVSLIAQLGSLPILLYYHFHQFS^IISVPMNMLMVPFYTCILPGA
 *

400
400

logo

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NTDB id 107 BSU 25570 NP 390435.1
consensus

VAGVLLLSLSASFGRLLFFSWFDLLISWNRLLITNIADV^EVFTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGQ

VAGVLLLSLSASFGRLLFFSWFDLLISWNRLLITNIADV^EVFTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGQ

VAGVLLLSLSASFGRLLFFSWFDLLISWNRLLITNIADV^EVFTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGQ

*

480
480

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consensus

560
560

logo

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consensus



640
640

logo

NTDB id 339871 ES968 RS13005 WP 046160607.1
NTDB id 107 BSU 25570 NP 390435.1
consensus



720
720

logo

NTDB id 339871 ES968 RS13005 WP 046160607.1
NTDB id 107 BSU 25570 NP 390435.1
consensus



776
776

 non conserved
 similar
 ≥ 50% conserved