

logo

NTDB id 283239 C7M28 RS03720 WP 069704087.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

MRNSRLLLPMAAASATAGITAAAYFPAIFLFI LLLIILIKTRHAFIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYP
MRNSRLLLPMAAASATAGITAAAYFPAIFLFI LLLIILIKTRHAFIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYP
MRNSRLLLPMAAASATAGITAAAYFPAIFLFI LLLIILIKTRHAFIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYP

80
80

logo

NTDB id 283239 C7M28 RS03720 WP 069704087.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

KAVIDTIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPKHATVPGAFDNEYLYRQHI
 KAVIDTIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPKHATVPGAFDNEYLYRQHI
 KAVIDTIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDNEYLYRQHI

160
160

logo

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consensus

HNNYSVTSIQNCSEPENFKYKVLRLKHIISFTNSLLPPDSTGIVQALTVGDRFYVEDEVLTAYQKLGWHLAISGLHV
 HWNYSVTSIQNCSEPENFKYKVLRLKHIISFTNSLLPPDSTGIVQALTVGDRFYVEDEVLTAYQKLGWVHLLAISGLHV
 HWNYSVTSIQNCSEPENFKYKVLRLKHIISFTNSLLPPDSTGIVQALTVGDRFYVEDEVLTAYQKLGWVHLLAISGLHV

240
240

logo

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consensus

320
320

logo

NTDB id 283239 C7M28 RS03720 WP 069704087.1
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consensus

YHLFEAGFQLSFAVSFSLILSSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYYHFHQFSISVPMNMLMVPFYTCILPGA
YHLFEAGFQLSFAVSFSLILSSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYYHFHQFSISVPMNMLMVPFYTCILPGA
YHLFEAGFQLSFAVSFSLILSSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYYHFHQFSISVPMNMLMVPFYTCILPGA

400
400

logo

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consensus

VAGVLLLSLSASFGRFFSWFDLLISHTNRLIT_KIADVDVFTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGG
VAGVLLLSLSASFGRFFSWFDLLISHTNRLIT_KIADVDVFTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGG
VAGVLLLSLSASFGRFFSWFDLLISHTNRLIT_NIADVDVFTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGG

480
480

logo

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consensus

ICCTV^MFLFLFIYPCLSSEGEVDMIDIGQGDSMFVGAPHQRGRVLI DTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKG
 ICCTV^LFLFLFIYPCLSSEGEVDMIDIGQGDSMFVGAPHQRGRVLI DTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKG
 ICCTV^LFLFLFIYPCLSSEGEVDMIDIGQGDSMFVGAPHQRGRVLI DTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKG
 *

560
560

logo

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consensus



640
640

logo

NTDB id 283239 C7M28 RS03720 WP 069704087.1
NTDB id 107 BSU 25570 NP 390435.1
consensus



720
720

logo

NTDB id 283239 C7M28 RS03720 WP 069704087.1
NTDB id 107 BSU 25570 NP 390435.1
consensus



776
776

 non conserved
 similar
 ≥ 50% conserved