

logo

NTDB id 59777 BJAB0715 RS17610 WP 020753491.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus



80
80

logo

NTDB id 59777 BJAB0715 RS17610 WP 020753491.1
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consensus



160
160

logo

NTDB id 59777 BJAB0715 RS17610 WP 020753491.1
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consensus



235
230

logo

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314
298

logo

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394
378

logo

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474
458

logo

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554
538

logo

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consensus

SYANVPLPIAGQPETRDAEGNITKYYYAVQSKIRNLFTDVSAVAADGSLTKISTSGTNLLKIPAAPPEGTNPFDTVANTA
SYANVPLPIAGQPETRDAEGNITKYYYAVQSKIRNLFTDVSAVAADGSLTKISTSGTNLLKIPAAPPEGTNPFDTVANTA
SYAKVPLPIAGQNETRDAEGNITKYYYAVQSKIRNLFTDVSAVAADGSLTKISTSGTNLLKIPAAPPEGTNPFDTVANTA
!!!!!!!!!!!!!!!!!!!!*!!

634
618

logo

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SYVLGKFDPTGQNILKAFPI SLKLLKILNLYGYSTDI SATTLPSSLVTSNEPYLSMGGSIHSLPVQLTYNGTLDDNGNLT
SYVLGKFDPTGQNILKAFPI SLKLLKILNLYGYSTDI SATTLPSSLVTSNEPYLSMGGSIHSLPVQLTYNGTLDDNGNLT
SYVLGKFDPTGQNILKAFPI SLKLLKILNLYGYSTDI SATTLPSSLVTSNEPYLSMGGSIHSLPVQLTYNGTLDDNGNLT
!!

714
698

logo

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SAREQSILYGTMEGGLHIVDASSGIEQMFVFPADILNDSVASKALVVGQSDASAPAHGMDGAWVSDPAYNITTVGSGSSA
SAREQSILYGTMEGGLHIVDASSGIEQMFVFPADILNDSVASKALVVGQSDASAPAHGMDGAWVSDPAYNITTVGSGSSA
SAREQSILYGTMEGGLHIVDASSGIEQMFVFPADILNDSVASKALVVGQSDASAPAHGMDGAWVSDPAYNITTVGSGSSA
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794
778

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VSKVTAKQMNIIYGGMRMGSSYYGLDVL SPTSPKLLFRIGADQTDYSRMGQSWSKPVLANIRYNGSIRRVLIVGGGYDQC
VSKVTAKQMNIIYGGMRMGSSYYGLDVL SPTSPKLLFRIGADQTDYSRMGQSWSKPVLANIRYNGSIRRVLIVGGGYDQC
VSKVTAKQMNIIYGGMRMGSSYYGLDVL SPTSPKLLFRIGADQTDYSRMGQSWSKPVLANIRYNGSIRRVLIVGGGYDQC
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874
858

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YEKPNITLTDACFTNGKAKGNAVYIIDAKTGQRLWWTSDTGSNTDNAMKHSIVSRISTLDRDADGLVDHLYFGDLGGQI
YEKPNITLTDACFTNGKAKGNAVYIIDAKTGQRLWWTSDTGSNTDNAMKHSIVSRISTLDRDADGLVDHLYFGDLGGQI
YEKPNITLTDACFTNGKAKGNAVYIIDAKTGQRLWWTSDTGSNTDNAMKHSIVSRISTLDRDADGLVDHLYFGDLGGQI
!!

954
938

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FRVDLNNQTKTNSTYSSFGVRVRLANLATNDSTYDGTNDYTGGNAPRFYEPPTVTIHDYGIHTFITVGIASGDRSTPL
FRVDLNNQTKTNSTYSSFGVRVRLANLATNDSTYDGTNDYTGGNAPRFYEPPTVTIHDYGIHTFITVGIASGDRSTPL
FRVDLNNQTKTNSTYSSFGVRVRLANLATNDSTYDGTNDYTGGNAPRFYEPPTVTIHDYGIHTFITVGIASGDRSTPL
!!

1034
1018

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DVYPLTGREGMTPASALSGRPVNNVYGIIDRDFVKKNLMSLTDNQLTKDITRTGLRKNPQILRTGETRVAQIFFPPTGXA
DVYPLTGREGMTPASALSGRPVNNVYGIIDRDFVKKNLMSLTDNQLTKDITRTGLRKNPQILRTGETRVAQIFFPPTGXA
DVYPLTGREGMTPASALSGRPVNNVYGIIDRDFVKKNLMSLTDNQLTKDITRTGLRKNPQILRTGETRVAQIFFPPTGXA
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1114
1098

