

logo

NTDB id 44493 SYNPCCN RS10725 WP 010873252.1
NTDB id 1419 SGL RS12550 WP 010873252.1
consensus



80
80

logo

NTDB id 44493 SYNPCCN RS10725 WP 010873252.1
NTDB id 1419 SGL RS12550 WP 010873252.1
consensus



160
160

logo

NTDB id 44493 SYNPCCN RS10725 WP 010873252.1
NTDB id 1419 SGL RS12550 WP 010873252.1
consensus



240
240

logo

NTDB id 44493 SYNPCCN RS10725 WP 010873252.1
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consensus



320
320

logo

NTDB id 44493 SYNPCCN RS10725 WP 010873252.1
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consensus



400
400

logo

NTDB id 44493 SYNPCCN RS10725 WP 010873252.1
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consensus



480
480

logo

NTDB id 44493 SYNPCCN RS10725 WP 010873252.1
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consensus



560
560

logo

NTDB id 44493 SYNPCCN RS10725 WP 010873252.1
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consensus

LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVRTSLIDIRGAINIESTVKGTTFTIRLPLTLSICKALCCLSDRARI
LSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVRTSLIDIRGAINIESTVKGTTFTIRLPLTLSICKALCCLSDRARI
!!

640
640

logo

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AFPMDGVEDTRDYLPKDIVTDEDGQKCVKYGKEALLPLYPKDLLNMRQLSRGSVYGGKQEEEDVVRVVIARGSSGNLL
AFPMDGVEDTRDYLPKDIVTDEDGQKCVKYGKEALLPLYPKDLLNMRQLSRGSVYGGKQEEEDVVRVVIARGSSGNLL
!!

720
720

logo

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ALQIDQVLGEEEIVIKQIEGPIPKPPGIAGATVRGDGSIPIADVLELIEIAQGRRLTDSSGGLWRKTLTPTVNEVHMDP
ALQIDQVLGEEEIVIKQIEGPIPKPPGIAGATVRGDGSIPIADVLELIEIAQGRRLTDSSGGLWRKTLTPTVNEVHMDP
!!

800
800

logo

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TVLIVDDSI TVRELLSLSFKKAGFRVEQARDGQEAWDQLKSGLPDILVFCDIEMPRKNGLELLSELQADEELSSIPVALL
TVLIVDDSI TVRELLSLSFKKAGFRVEQARDGQEAWDQLKSGLPDILVFCDIEMPRKNGLELLSELQADEELSSIPVALL
!!

880
880

logo

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TSRGADRHRKVAAKLGATGYFTKPYTEKDILDAANRMLNGEVLLPGSTRKTNSILSRFKKARGNFSEEAPTIVRSNPCTL
TSRGADRHRKVAAKLGATGYFTKPYTEKDILDAANRMLNGEVLLPGSTRKTNSILSRFKKARGNFSEEAPTIVRSNPCTL
!!

960
960

logo

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RSTVLIIDDSVMVREMLSMTFKKADYHTETARDGQDAWEKLSGLPCDLILCDIEMPRMNGLELLERIQEDPNLSRIPVA
RSTVLIIDDSVMVREMLSMTFKKADYHTETARDGQDAWEKLSGLPCDLILCDIEMPRMNGLELLERIQEDPNLSRIPVA
!!

1040
1040

logo

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MITSRGAQMKRMAAEKGAQGYFVKPYIEETLLEASQRLIRGEVLLDLAALSAD
MITSRGAQMKRMAAEKGAQGYFVKPYIEETLLEASQRLIRGEVLLDLAALSAD
!!

1095
1095

 non conserved
 similar
 $\geq 50\%$ conserved