

logo

NTDB id 123717 BSUB RS13865 WP 009967776.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

MRNSRLLLPMAAASATAGITAAAYFPAIFLFI¹L²LIKTRHAF³LI⁴VCFFS⁵FI⁶FFV⁷LYAVTDSQNVSSYRQGT⁸YQFKAV⁹ID

85
85

logo

NTDB id 123717 BSUB RS13865 WP 009967776.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

TIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDYNEYLYRQHIHWNYSVTSIQ
TIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDYNEYLYRQHIHWNYSVTSIQ
TIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDYNEYLYRQHIHWNYSVTSIQ

170
170

logo

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consensus

NCSEPE¹NFKYK²VL³SL⁴RK⁵HI⁶IS⁷FT⁸NS⁹LL¹⁰PP¹¹DST¹²G¹³IV¹⁴QAL¹⁵TV¹⁶GDR¹⁷FY¹⁸VE¹⁹DE²⁰VL²¹TAY²²QKL²³GW²⁴HLL²⁵AI²⁶SG²⁷LHV²⁸GIL²⁹TAG³⁰LFY³¹IM³²IRL³³G³⁴

255
255

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consensus

ITREKASILLLLFLPLYVMLTGAAPSVLRAALMSGVYLAGSLVKWRVRSATAICLSYIVLLLFNPYHLFEAGFQLSFAVSFSLIL
ITREKASILLLLFLPLYVMLTGAAPSVLRAALMSGVYLAGSLVKWRVRSATAICLSYIVLLLFNPYHLFEAGFQLSFAVSFSLIL
ITREKASILLLLFLPLYVMLTGAAPSVLRAALMSGVYLAGSLVKWRVRSATAICLSYIVLLLFNPYHLFEAGFQLSFAVSFSLIL

340
340

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SSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSISVPMNMLMVPFYTFCILPGAVAGVLLLSLSASFGRLLFFSWFDLLI
SSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSISVPMNMLMVPFYTFCILPGAVAGVLLLSLSASFGRLLFFSWFDLLI
SSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSISVPMNMLMVPFYTFCILPGAVAGVLLLSLSASFGRLLFFSWFDLLI

425
425

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SWINRLITNIADVDFVTIMIAHPAPVLLFLFTVTIILLLMAIEKRSLSQLMVTGGICCTVMFLLFIYPCLSSSEGEVDMIDIGGD
SWINRLITNIADVDFVTIMIAHPAPVLLFLFTVTIILLLMAIEKRSLSQLMVTGGICCTVMFLLFIYPCLSSSEGEVDMIDIGGD
SWINRLITNIADVDFVTIMIAHPAPVLLFLFTVTIILLLMAIEKRSLSQLMVTGGICCTVMFLLFIYPCLSSSEGEVDMIDIGGD

510
510

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SMFVGAPHQGRVLI DTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKGIKQLDALILTHADQDHIGEAEILLKHHKVKRLVIP
SMFVGAPHQGRVLI DTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKGIKQLDALILTHADQDHIGEAEILLKHHKVKRLVIP
SMFVGAPHQGRVLI DTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKGIKQLDALILTHADQDHIGEAEILLKHHKVKRLVIP

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595

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consensus



680
680

logo

NTDB id 123717 BSUB RS13865 WP 009967776.1
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consensus



765
765

logo

NTDB id 123717 BSUB RS13865 WP 009967776.1
NTDB id 107 BSU 25570 NP 390435.1
consensus



776
776

☐ non conserved
☒ similar
☒ ≥ 50% conserved