

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 221914 BAJT RS11720 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTI¹GDKRLGAALLDAGLLTDEELQRALERHREVGGS²⁰LAEVLVDMGLLSERRIAQT³⁰IEDRFGIPLVELHRVEIPPKVKALLPA⁴⁰
MSVLTI¹GDKRLGAALLDAGLLTDEELQRALERHREVGGS²⁰LAEVLVDMGLLSERRIAQT³⁰IEDRFGIPLVELHRVEIPPKVKALLPA⁴⁰
.....
.....

85
0
0
0

logo

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NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 221914 BAJT RS11720 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

EKAKELKAIPFALDEEAGVVRVAFLNPLD¹TL²SLEEV³EDLTGLV⁴EPYQT⁵TKSAFLYALAKHY⁶PELGLPVP⁷PPPPSGEGQ⁸KDLKLGE
 EKAKELKAIPFALDEEAGVVRVAFLNPLD¹TL²SLEEV³EDLTGLV⁴VEPYQT⁵TKSAFLYALAKHY⁶PELGLPVP⁷PPPPSGEGQ⁸KDLKLGE

170
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logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

LLLQKGWISREALVEEALVEQKGTGDLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVPDPSAALLLLRSDALRYGAVPIGF
 LLLQKGWISREALVEEALVEQKGTGDLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVPDPSAALLLLRSDALRYGAVPIGF

255
0
0
0

logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

QNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
QNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
.....

340
0
0
0

logo

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consensus

VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGPIPCDPEEDPIFDISP
VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGPIPCDPEEDPIFDISP
.....MSQYTICETQTERIFDISPSRWQKNCEEKQLLLRYLAVP
.....* * * * *

420
39
0
0

logo

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NTDB id 221914 BAJT RS11720 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

[illegible]

498
109
0
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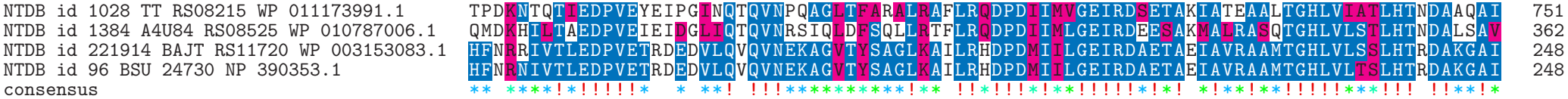
logo



logo



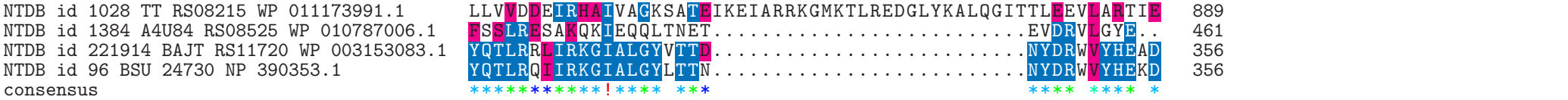
logo



logo



logo



⊠ non conserved
⊠ similar
⊠ ≥ 50% conserved