

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 231062 BAGQ RS11825 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 231062 BAGQ RS11825 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

EKAKELKAIPFALDEEAGVVRVAFLNPLD¹TL²SLEEVEDLTGLVPEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQDKLKLGE
 EKAKELKAIPFALDEEAGVVRVAFLNPLD¹TL²SLEEVEDLTGLVPEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQDKLKLGE

170
0
0
0

logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

LLLQKGWISREALVEEALVEQKTGDLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVDPDPSAALLLLRSDALRYGAVPIGF
 LLLQKGWISREALVEEALVEQKTGDLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVDPDPSAALLLLRSDALRYGAVPIGF

255
0
0
0

logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus
```

QNGEVEVVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
QNGEVEVVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
.....

340
0
0
0

logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGPIPCDPEEDPIFDISP
VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGPIPCDPEEDPIFDISP
.....MSQYITICEQQTERIFDISPSRWQKNCEEKQLLLRYLAVP
.....* * * * *

420
39
0
0

logo

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consensus

YRLTECNRLVWLVDERNLTALCELYRMLTRKGLNKYEAPYVWISDEITYLINALNSPEQEPFYKSELSLEYEQEFAKKQAEEEVPS
 HRLEGNRIIVLLKDPRIILALDVRLLALKRKGLNYEVAPAVATEAAITTKLI.....ERFYGKAELSEIAKEFAKKQAEEEVPS
 VQETEHKLWLAVDDENNLTACEIFAFM.TH....KQLPVPVIASDELKYLLNALNSPEQQPIYEESELAFVEQE.....QE

 * * * * * * * * * * * * * * * * *

498
109
0
0

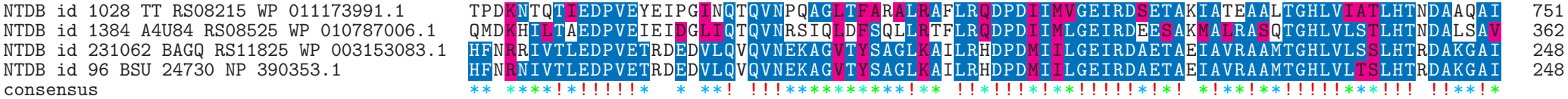
logo



logo



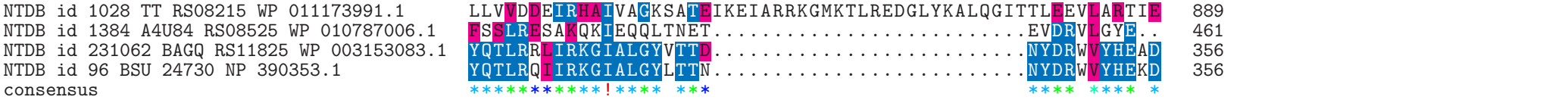
logo



logo



logo



⊠ non conserved
⊠ similar
⊠ ≥ 50% conserved