

logo

NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1468 ASU2 RS05090 WP 014991734.1
NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 52784 BVAD3 RS12110 WP 014418378.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA
.....
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA
.....

0
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85
0
0

logo

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NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 52784 BVAD3 RS12110 WP 014418378.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

EKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQKDLKLGE
 EKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQKDLKLGE

0
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170
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logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

.....
 LLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLLLRSDALRYGAVPIGF

 LLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLLLRSDALRYGAVPIGF

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0
255
0
0

logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

QNGEVEVVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
.....
QNGEVEVVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
.....

0
0
340
0
0

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NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 52784 BVAD3 RS12110 WP 014418378.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGQYTYCDSTORYFDISPSWQKNCKLEKSLRLRYLAVP
MSQYTTICEQQTERIFDISPSRWQKNCEEKQLLLRYLAVP
MQYSVCDMNSQQVFDISESLWQKNCKDEKHILLRYLAIP
 VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEEDPPDPGAPL....LLPEDLCRRYGVFP

*

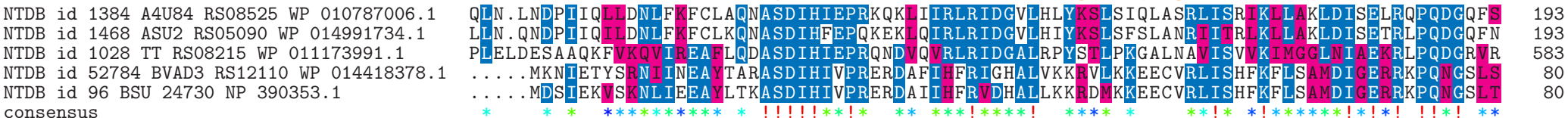
39
38
420
0
0

* * *

logo



logo



logo



logo

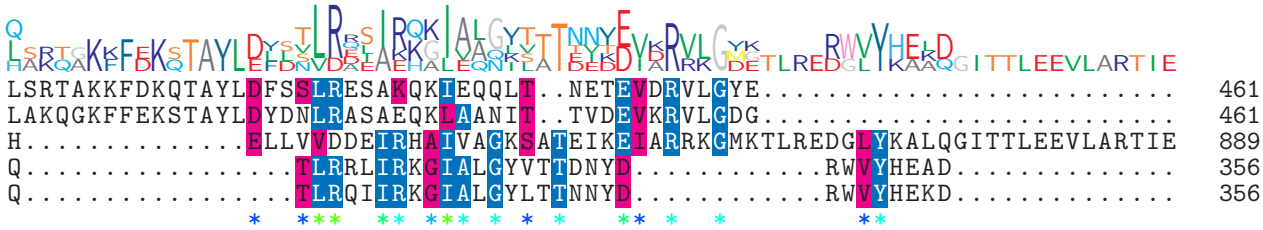


logo



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consensus



ⓧ non conserved
ⓧ similar
ⓧ ≥ 50% conserved