

logo

NTDB id 200923 KAB01 RS17235 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus

MKLKLKLFKPNNLWYAVYSSSMIFTWLTSSVQASDLQIYASPTAGKKTIVMMLDTSGSMTNSYGENRLAMLKNGMNA
 MKLKLKLFKPNNLWYAVYSSSMIFTWLTSSVQASDLQIYASPTAGKKTIVMMLDTSGSMTNSYGENRLAMLKNGMNA
 MKLKLKLFKPNNLWYAVYSSSMIFTWLTSSVQASDLQIYASPTAGKKTIVMMLDTSGSMTNSYGENRLAMLKNGMNA

80
80

logo

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NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus

FLASNNPVLNDTRVGLGNFSAN^DDSRSGQILVAAAPLGDASTLNT^MGSQRYKLKQAVAN^LTAG^SGYTPSAHAYAEAAAYLM
FLASNNPVLNDTRVGLGNFSAN^DDSRSGQILVAAAPLGDASTLNT^VGSQRYKLKQAVAN^LTAG^{GS}TPSAHAYAEAAAYLM
FLASNNPVLNDTRVGLGNFSAD^DDSRSGQILVAAAPLGDASTLNT^MGSQRYKLKQAVAS^LTAG^{GY}TPSAHAYAEAAAYLM

160
160

logo

NTDB id 200923 KAB01 RS17235 WP 000768970.1
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consensus

235
230

logo

NTDB id 200923 KAB01 RS17235 WP 000768970.1
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consensus

314
298

logo

NTDB id 200923 KAB01 RS17235 WP 000768970.1
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consensus

394
378

logo

NTDB id 200923 KAB01 RS17235 WP 000768970.1
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consensus

474
458

logo

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consensus

KNLQEYGYLRAFEPNPANTYLTWRGNLKKYHWLSG_ANAG_AFEANTGGLVYNATGA_AFRTGT_ATKDY_ANSSTYNDGGKVFFGG
 KNLQEYGYLRAFEPNPANTYLTWRGNLKKYHVVLSG_ANAG_ATFEANTGGLVYNATGA_AFRTGT_ATKDY_ANSSTYNDGGKVFFGG
 KNLQEYGYLRAFEPNPANTYLTWRGNLKKYHVVLSG_TNAG_AFEANTGGLVYNATGA_AFRTGT_ATKDY_ANSSTYNDGGKVFFGG

554
538

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consensus



1194
1178

logo

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consensus



1274
1258

logo

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NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus



1284
1268

☒ non conserved
☒ similar
☒ ≥ 50% conserved