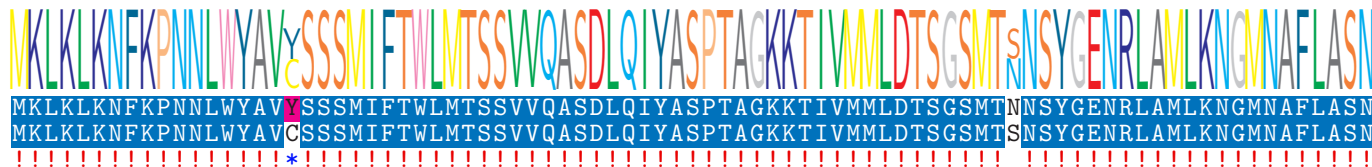


logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus

85
85

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus



170
170

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus



249
245

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus



334
318

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus



419
403

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus



504
488

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus



589
573

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus

LFTDVSAVAADGSLTKISTSGTNLLKIPAAPPEGTNPFDTVANTASYVLGKFDPSTGQNILKAFPISLKCLKILNYLGYSTDI₅AT

674
658

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus

TLPSSLVTSNEPYLSMGGSIHSLPVQLTYNGTLDDNGNLTSAREQSILYGTMEGGLHIVDASSGIEQMVFPADILNDSVASKAL
 TLPSSLVTSNEPYLSMGGSIHSLPVQLTYNGTLDDNGNLTSAREQSILYGTMEGGLHIVDASSGIEQMVFPADILNDSVASKAL
 TLPSSLVTSNEPYLSMGGSIHSLPVQLTYNGTLDDNGNLTSAREQSILYGTMEGGLHIVDASSGIEQMVFPADILNDSVASKAL

759
743

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus

844
828

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus

QSSKPVLANIRYNGSIRRVLI VGGGYDQCYEKPNI TLTDACFTNGKAKGNAVYIIDAKTGQRLWWTSDTGSNTDANMKHSIVS
QSSKPVLANIRYNGSIRRVLI VGGGYDQCYEKPNI TLTDACFTNGKAKGNAVYIIDAKTGQRLWWTSDTGSNTDANMKHSIVS
QSSKPVLANIRYNGSIRRVLI VGGGYDQCYEKPNI TLTDACFTNGKAKGNAVYIIDAKTGQRLWWTSDTGSNTDANMKHSIVS

929
913

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus

R I S T L D R D A D G L V D H L Y F G D L G G Q I F R V D L N N N Q T K T N S T Y S S F G V R V R L A N L A T N D S T Y D G T N D Y T G G N A P R F Y E P P T V T I H D
R I S T L D R D A D G L V D H L Y F G D L G G Q I F R V D L N N N Q T K T N S T Y S S F G V R V R L A N L A T N D S T Y D G T N D Y T G G N A P R F Y E P P T V T I H D
R I S T L D R D A D G L V D H L Y F G D L G G Q I F R V D L N N N Q T K T N S T Y S S F G V R V R L A N L A T N D S T Y D G T N D Y T G G N A P R F Y E P P T V T I H D

1014
998

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus

YGIHTFITLGIASGDRSTPLDVYPLTGREGMTPASALSGRPVNNVYGIIDRDFVKKNLMSLTDNQLETKDITRTGLRKNPQILRT
 YGIHTFITLGIASGDRSTPLDVYPLTGREGMTPASALSGRPVNNVYGIIDRDFVKKNLMSLTDNQLETKDITRTGLRKNPQILRT
 YGIHTFITLGIASGDRSTPLDVYPLTGREGMTPASALSGRPVNNVYGIIDRDFVKKNLMSLTDNQLETKDITRTGLRKNPQILRT
 *

1099
1083

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus

GETRVAQIFFPTTGVGKGGMYRSLSSTSDGTEKANNSFR|KGG|KAFEFPM|ITGNL|IPVYDPQGTG|VAADPCLPRVGETDR
 GETRVAQIFFPTTGVGKGGWYRSLSSTSDGTEKANNSFR|KGG|KAFEFPM|ITGNL|IPVYDPQGTG|VAADPCLPRVVGETDR
 GETRVAQIFFPTTGVGKGGWYRSLSSTSDGTEKANNSFR|KGG|KAFEFPM|ITGNL|IPVYDPQGTG|VAADPCLPRVVGETDR

1184
1168

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus



1269
1253

logo

NTDB id 120573 BL01 RS10215 WP 000768970.1
NTDB id 1063 ABD1 RS15730 WP 000768961.1
consensus



1284
1268

- X non conserved
- X similar
- X ≥ 50% conserved