

logo

NTDB id 115111 N900 RS11530 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus

MRLCPRFSLDLLVKVKLLVNFVTLGFSCLSPTTHAWRSPQSFTESNTLIMFRFYRKQKFKRLQNTLMAAFLALSIIPLTIT
 MRLCPRFSLDLLVKVKLLVNFVTLGFSCLSPTTHAWRSPQSFTESNTLIMFRFYRKQKFKRLQNTLMAAFLALSIIPLTIT
 MRLCPRFSLDLLVKVKLLVNFVTLGFSCLSPTTHAWRSPQSFTESNTLIMFRFYRKQKFKRLQNTLMAAFLALSIIPLTIT

80
80

logo

NTDB id 115111 N900 RS11530 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus

ALFFFLHSHSKDLEQQSTSYLVSVRDNKQQQVIDYMMAKESEVMGFVRSELAYASGGRFYGLVNAFQRLDVSIEAAREHAQ
ALFFFLHSHSKDLEQQSTSYLVSVRDNKQQQVIDYMMAKESEVMGFVRSELAYASGGRFYGLVNAFQRLDVSIEAAREHAQ

160
160

logo

NTDB id 115111 N900 RS11530 WP 001883436.1
NTDB id 1143 A1552VC RS01965 WP 001883436.1
consensus

QRYITGSGDQIKTSVLPQSSSYVGSEYRLLHKRYHWAYLELLKRSDFDDILLVDIDGNVVYSIYKYDNFGTNLLTGKYQ
 QRYITGSGDQIKTSVLPQSSSYVGSEYRLLHKRYHWAYLELLKRSDFDDILLVDIDGNVVYSIYKYDNFGTNLLTGKYQ

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[illegible]

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Q T I L V G Q D H R S R T L T S A E I A V E K S K A W D L A L S G K R A V G T Y T N T D G E Q I I A A Y A P I N L K N I H W A L V V E L P E K E A F A R V R Q
 Q T I L V G Q D H R S R T L T S A E I A V E K S K A V V D L A L S G K R A V G T Y T N T D G E Q I I A A Y A P I N L K N I H W A L V V E L P E K E A F A R V R Q 400
 Q T I L V G Q D H R S R T L T S A E I A V E K S K A V V D L A L S G K R A V G T Y T N T D G E Q I I A A Y A P I N L K N I H W A L V V E L P E K E A F A R V R Q 400

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LEKLFVFA¹MLTAIVLV²VIASHYLSNFI³TSPLLKLTWAA⁴ERVSAGDL⁵DEAMINTERKDEIGRLAVSF⁶ERMQRSIREKISLI⁷

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consensus



640
640

logo

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consensus



720
720

logo

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consensus



800
800

logo

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consensus



880
880

logo

NTDB id 115111 N900 RS11530 WP 001883436.1
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consensus



960
960

logo

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consensus



1040
1040

logo

NTDB id 115111 N900 RS11530 WP 001883436.1
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consensus



1120
1120

logo

NTDB id 115111 N900 RS11530 WP 001883436.1
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consensus



1177
1177

- ☐ non conserved
- ☒ similar
- ☒ ≥ 50% conserved