

logo

NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 216489 BZG08 RS22795 WP 001013201.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MLTNLSTVLRQAGILTFSQEESLLEQVKASGISMPEALLSSGFFQSHELTEHLSSVFGLSCTSLNQYEYASLCQTLGLRD
MLTNLSTVLRQAGILTFSQEESLLEQVKASGISMPEALLSSGFFQSHELTEHLSSVFGLSCTSLNQYEYASLCQTLGLRD
.....

80
0
0

logo

NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 216489 BZG08 RS22795 WP 001013201.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

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

160
0
0

logo

NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 216489 BZG08 RS22795 WP 001013201.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

240
62
62

logo

NTDB id 1401 DSB67 RS12665 WP 010643259.1
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NTDB id 96 BSU 24730 NP 390353.1
consensus

[illegible]

320
140
140

logo

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NTDB id 216489 BZG08 RS22795 WP 001013201.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

YFTGPTGSGKTTTLVLLSYLRPTKRNIVTLEDPVERRNDVQLQVQVNEKAGVTYSEAGLKAILRHDPDVMVGEIRDLE
 TMISSMAGVEYAKIKHFEIRSIALLKYDDGINIQFIIFGFAEARSFRLQIMVTLAD
 LMTGPTGSGKTVSLYIGLSIINK.PEINISTAEDPVEIYLPGINQVQVQPKIGFGFAEALRSFLRQDPDVMVGEIRDLD
 VFTGPTGSGKTTTMYALLEVIRKKKTRIVTLEDPVEKRNDVQLQIQINEKAGITYEAGLKAILRHDPDVLVGEIRDDE
 IFTGPTGSGKTTTLVSLVQYAKKHFNRIIVTLEDPVETRDNDVQLQVQVNEKAGVTYSAGLKAILRHDPDMILGEIRDAE

399
220
220

logo

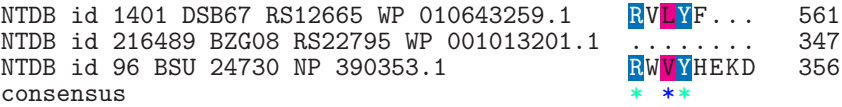
NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 216489 BZG08 RS22795 WP 001013201.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

476
299
300

logo



logo



non conserved
 similar
 ≥ 50% conserved