

logo

NTDB id 1015 ACIAD RS12005 WP 004928865.1	MFKIILLGWVAGIALMGIDFPLIMQYKVGELLLAFIFLYLKRPMEVDRPFLKAVFCLLCITSLFMVGYHYAEKALIE	80
NTDB id 313007 AhaeAN59 RS04990 WP 160126531.1	MFKIILLGWVAGIALMGIDFPLIMQYKVGELLLAFIFLYLKRPMEVDRPFLKAVFCLLCITSLFMVGYHYAEKALIE	80
NTDB id 1070 ABD1 RS13145 WP 000472713.1	MFKIILLGWVAGIALMGIDFPLIMQYKVGELLLAFIFLYLKRPMEVDRPFLKAVFCLLCITSLFMVGYHYAEKALIE	80
NTDB id 1083 HON27 RS04190 WP 168726476.1	MFKIILLGWVAGIALMGIDFPLIMQYKVGELLLAFIFLYLKRPMEVDRPFLKAVFCLLCITSLFMVGYHYAEKALIE	80
consensus	*****	

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NTDB id 1015 ACIAD RS12005 WP 004928865.1	RLLEQRETDRTLEIVYNRLSEKDNKVGQVQLVNLISKEPVMVLYLKSNNQNLKNNQNLGLGHYYRISCKTRPAHS	159
NTDB id 313007 AhaeAN59 RS04990 WP 160126531.1	RLLEQRETDRTLEIVYNRLSEKDNKVGQVQLVNLISKEPVMVLYLKSNNQNLKNNQNLGLGHYYRISCKTRPAHS	159
NTDB id 1070 ABD1 RS13145 WP 000472713.1	RLLEQRETDRTLEIVYNRLSEKDNKVGQVQLVNLISKEPVMVLYLKSNNQNLKNNQNLGLGHYYRISCKTRPAHS	160
NTDB id 1083 HON27 RS04190 WP 168726476.1	RLLEQRETDRTLEIVYNRLSEKDNKVGQVQLVNLISKEPVMVLYLKSNNQNLKNNQNLGLGHYYRISCKTRPAHS	160
consensus	*****	

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NTDB id 1015 ACIAD RS12005 WP 004928865.1	YATPGAFDEKNFQRLQSGFNVRYLPLSDEFYRLGYQQLKQOQSFASFRNVYKRLRLFRQLNSSLQKGLL	239
NTDB id 313007 AhaeAN59 RS04990 WP 160126531.1	YATPGAFDEKNFQRLQSGFNVRYLPLSDEFYRLGYQQLKQOQSFASFRNVYKRLRLFRQLNSSLQKGLL	239
NTDB id 1070 ABD1 RS13145 WP 000472713.1	YATPGAFDEKNFQRLQSGFNVRYLPLSDEFYRLGYQQLKQOQSFASFRNVYKRLRLFRQLNSSLQKGLL	240
NTDB id 1083 HON27 RS04190 WP 168726476.1	YATPGAFDEKNFQRLQSGFNVRYLPLSDEFYRLGYQQLKQOQSFASFRNVYKRLRLFRQLNSSLQKGLL	240
consensus	*****	

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NTDB id 1015 ACIAD RS12005 WP 004928865.1	LALLTGDESLLSDETCQFQQLGSHLLAISGPHVLIFAAMLSWYHQHLSRYYPVLYLWKPRQYLYLPFCLGVLLYTA	319
NTDB id 313007 AhaeAN59 RS04990 WP 160126531.1	LALLTGDESLLSDETCQFQQLGSHLLAISGPHVLIFAAMLSWYHQHLSRYYPVLYLWKPRQYLYLPFCLGVLLYTA	319
NTDB id 1070 ABD1 RS13145 WP 000472713.1	LALLTGDESLLSDETCQFQQLGSHLLAISGPHVLIFAAMLSWYHQHLSRYYPVLYLWKPRQYLYLPFCLGVLLYTA	320
NTDB id 1083 HON27 RS04190 WP 168726476.1	LALLTGDESLLSDETCQFQQLGSHLLAISGPHVLIFAAMLSWYHQHLSRYYPVLYLWKPRQYLYLPFCLGVLLYTA	320
consensus	*****	

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NTDB id 1015 ACIAD RS12005 WP 004928865.1	FVGFEIPALRTITVLIESGELLRLQPKPFTLLVYSASLLLMDPFSYLSAGFWLSYGACFILIRIYQDTIAQLPQSHL	399
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NTDB id 1070 ABD1 RS13145 WP 000472713.1	FVGFEIPALRTITVLIESGELLRLQPKPFTLLVYSASLLLMDPFSYLSAGFWLSYGACFILIRIYQDTIAQLPQSHL	400
NTDB id 1083 HON27 RS04190 WP 168726476.1	FVGFEIPALRTITVLIESGELLRLQPKPFTLLVYSASLLLMDPFSYLSAGFWLSYGACFILIRIYQDTIAQLPQSHL	400
consensus	*****	

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NTDB id 1015 ACIAD RS12005 WP 004928865.1	SLSSKMIFMSKVLIESQWKIFYALSPLTLFFQQLSWYAPLTNLIAPIYGSYVPLNMAACAWLYIPPLGNLFEHND	478
NTDB id 313007 AhaeAN59 RS04990 WP 160126531.1	SLSSKMIFMSKVLIESQWKIFYALSPLTLFFQQLSWYAPLTNLIAPIYGSYVPLNMAACAWLYIPPLGNLFEHND	479
NTDB id 1070 ABD1 RS13145 WP 000472713.1	SLSSKMIFMSKVLIESQWKIFYALSPLTLFFQQLSWYAPLTNLIAPIYGSYVPLNMAACAWLYIPPLGNLFEHND	480
NTDB id 1083 HON27 RS04190 WP 168726476.1	SLSSKMIFMSKVLIESQWKIFYALSPLTLFFQQLSWYAPLTNLIAPIYGSYVPLNMAACAWLYIPPLGNLFEHND	480
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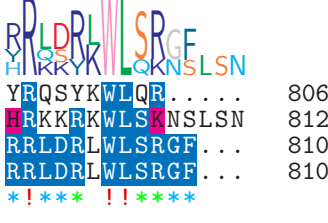
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consensus



☒ non conserved
☒ similar
☒ ≥ 50% conserved