

logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1	MAKN	72
NTDB id 407355 GPY19 RS03185 WP 158131160.1	MAK	71
NTDB id 1402 DSB67 RS12670 WP 010643257.1	MAK	71
NTDB id 1112 NGFG RS09215 WP 003689811.1	MAK	74
NTDB id 1113 AAA85695.1 219..1451()	MAK	74
NTDB id 1252 GC085 RS07730 WP 011213805.1	MAK	71
NTDB id 1198 PSJM300 03950 AFN76868.1	MAK	70
NTDB id 1016 ACIAD RS01680 WP 004920476.1	MAK	73
NTDB id 1059 ABD1 RS01610 WP 000279215.1	MAK	73
consensus	*** * * * * ! * * ! * ! * * * * * ! * * * * * ! * * *	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	FTRQLATM	152
NTDB id 407355 GPY19 RS03185 WP 158131160.1	FTRQISTMLVT	151
NTDB id 1402 DSB67 RS12670 WP 010643257.1	FTRQISTMLVT	151
NTDB id 1112 NGFG RS09215 WP 003689811.1	FTRQISTMLVT	154
NTDB id 1113 AAA85695.1 219..1451()	FTRQISTMLVT	154
NTDB id 1252 GC085 RS07730 WP 011213805.1	FTRQISTMLVT	151
NTDB id 1198 PSJM300 03950 AFN76868.1	FTRQISTMLVT	150
NTDB id 1016 ACIAD RS01680 WP 004920476.1	FTRQISTMLVT	153
NTDB id 1059 ABD1 RS01610 WP 000279215.1	FTRQISTMLVT	153
consensus	* * ! * * ! * * * * ! * * * * * * * * * * * * * * * ! * * * * * ! * * *	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	ERLATYREKSEQ	232
NTDB id 407355 GPY19 RS03185 WP 158131160.1	ERLATYREKSEQ	231
NTDB id 1402 DSB67 RS12670 WP 010643257.1	ERLATYREKSEQ	231
NTDB id 1112 NGFG RS09215 WP 003689811.1	ERLATYREKSEQ	234
NTDB id 1113 AAA85695.1 219..1451()	ERLATYREKSEQ	234
NTDB id 1252 GC085 RS07730 WP 011213805.1	ERLATYREKSEQ	231
NTDB id 1198 PSJM300 03950 AFN76868.1	ERLATYREKSEQ	230
NTDB id 1016 ACIAD RS01680 WP 004920476.1	ERLATYREKSEQ	233
NTDB id 1059 ABD1 RS01610 WP 000279215.1	ERLATYREKSEQ	233
consensus	* * * ! * * ! * * * ! * * ! * * * * * ! * * ! * * * * * ! * * *	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	AIAAAIFGLKAL	312
NTDB id 407355 GPY19 RS03185 WP 158131160.1	GSISVFISARVLS	311
NTDB id 1402 DSB67 RS12670 WP 010643257.1	SFVSLISGKIMS	311
NTDB id 1112 NGFG RS09215 WP 003689811.1	ALGFAIYGFLLK	314
NTDB id 1113 AAA85695.1 219..1451()	ALGFAIYGFLLK	314
NTDB id 1252 GC085 RS07730 WP 011213805.1	ALGGVVSFFYAKN	311
NTDB id 1198 PSJM300 03950 AFN76868.1	FVFGAAYAFKQAH	310
NTDB id 1016 ACIAD RS01680 WP 004920476.1	VIGAATAGFLEAK	313
NTDB id 1059 ABD1 RS01610 WP 000279215.1	AIGAVIAAFLEAK	313
consensus	* * * * * ! * * * * * ! * * * * * ! * * * * * ! * * * * * ! * * *	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 407355 GPY19 RS03185 WP 158131160.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1252 GCO85 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus

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logo

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NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 407355 GPY19 RS03185 WP 158131160.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
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NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus

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V V A M Y L P I F N L M S V L G . 408
V T A M Y L P I F N L M S V L G . 407
L V A M Y L P I F N L G N V V A . 410
L V A M Y L P I F N L G N V V A * 410
V V A M Y L P I F K L G S A V . 406
T I A M Y L P I F Q M G S V V . 405
V I A M Y L P I F Q M G S V V . 408
V I A M Y L P I F Q M G S V V . 408

- ☐ non conserved
- ☒ similar
- ☒ $\geq 50\%$ conserved