

logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1	MAKN	72
NTDB id 1402 DSB67 RS12670 WP 010643257.1	MAKNGGFS	71
NTDB id 1113 AAA85695.1 219..1451()	MAKNGGFS	74
NTDB id 94596 AN159 RS07150 WP 069972465.1	MAKNGGFS	74
NTDB id 1112 NGFG RS09215 WP 003689811.1	MAKNGGFS	74
NTDB id 1252 GC085 RS07730 WP 011213805.1	MAKNGGFS	71
NTDB id 1198 PSJM300 03950 AFN76868.1	MAKNGGFS	70
NTDB id 1016 ACIAD RS01680 WP 004920476.1	MAKNGGFS	73
NTDB id 1059 ABD1 RS01610 WP 000279215.1	MAKNGGFS	73
consensus	MAKNGGFS	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	FTRQLATM	152
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NTDB id 1252 GC085 RS07730 WP 011213805.1	FTRQLATM	151
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NTDB id 1059 ABD1 RS01610 WP 000279215.1	FTRQLATM	153
consensus	FTRQLATM	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	ERLATYREK	232
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NTDB id 1059 ABD1 RS01610 WP 000279215.1	ERLATYREK	233
consensus	ERLATYREK	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1	ALGAIYGL	312
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NTDB id 1059 ABD1 RS01610 WP 000279215.1	ALGAIYGL	313
consensus	ALGAIYGL	

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NTDB id 1169 A1552VC RS11075 WP 000648511.1
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NTDB id 1016 ACIAD RS01680 WP 004920476.1
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consensus

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NTDB id 1169 A1552VC RS11075 WP 000648511.1
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NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus

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[illegible]

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
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
I T 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