

logo

NTDB id 70274 V060002 RS02675 WP 000648511.1	MAKNGGFSLSFAKKEKRFIFEGRHSSASDKLVNGEVSFAFTEEEARKKLA	72
NTDB id 1169 A1552VC RS11075 WP 000648511.1	MAKNGGFSLSFAKKEKRFIFEGRHSSASDKLVNGEVSFAFTEEEARKKLA	72
NTDB id 1402 DSB67 RS12670 WP 010643257.1	MAKNGGFSLSFAKKEKRFIFEGRHSSASDKLVNGEVSFAFTEEEARKKLA	71
NTDB id 1112 NGFG RS09215 WP 003689811.1	MAKNGGFSLSFAKKEKRFIFEGRHSSASDKLVNGEVSFAFTEEEARKKLA	74
NTDB id 1113 AAA85695.1 219..1451()	MAKNGGFSLSFAKKEKRFIFEGRHSSASDKLVNGEVSFAFTEEEARKKLA	74
NTDB id 1252 GC085 RS07730 WP 011213805.1	MAKNGGFSLSFAKKEKRFIFEGRHSSASDKLVNGEVSFAFTEEEARKKLA	71
NTDB id 1198 PSJM300 03950 AFN76868.1	MAKNGGFSLSFAKKEKRFIFEGRHSSASDKLVNGEVSFAFTEEEARKKLA	70
NTDB id 1016 ACIAD RS01680 WP 004920476.1	MAKNGGFSLSFAKKEKRFIFEGRHSSASDKLVNGEVSFAFTEEEARKKLA	73
NTDB id 1059 ABD1 RS01610 WP 000279215.1	MAKNGGFSLSFAKKEKRFIFEGRHSSASDKLVNGEVSFAFTEEEARKKLA	73
consensus	MAKNGGFSLSFAKKEKRFIFEGRHSSASDKLVNGEVSFAFTEEEARKKLA	

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NTDB id 70274 V060002 RS02675 WP 000648511.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAH	152
NTDB id 1169 A1552VC RS11075 WP 000648511.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAH	152
NTDB id 1402 DSB67 RS12670 WP 010643257.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAH	151
NTDB id 1112 NGFG RS09215 WP 003689811.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAH	154
NTDB id 1113 AAA85695.1 219..1451()	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAH	154
NTDB id 1252 GC085 RS07730 WP 011213805.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAH	151
NTDB id 1198 PSJM300 03950 AFN76868.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAH	150
NTDB id 1016 ACIAD RS01680 WP 004920476.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAH	153
NTDB id 1059 ABD1 RS01610 WP 000279215.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAH	153
consensus	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAH	

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NTDB id 70274 V060002 RS02675 WP 000648511.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMF	232
NTDB id 1169 A1552VC RS11075 WP 000648511.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMF	232
NTDB id 1402 DSB67 RS12670 WP 010643257.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMF	231
NTDB id 1112 NGFG RS09215 WP 003689811.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMF	234
NTDB id 1113 AAA85695.1 219..1451()	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMF	234
NTDB id 1252 GC085 RS07730 WP 011213805.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMF	231
NTDB id 1198 PSJM300 03950 AFN76868.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMF	230
NTDB id 1016 ACIAD RS01680 WP 004920476.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMF	233
NTDB id 1059 ABD1 RS01610 WP 000279215.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMF	233
consensus	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMF	

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NTDB id 70274 V060002 RS02675 WP 000648511.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATS	312
NTDB id 1169 A1552VC RS11075 WP 000648511.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATS	312
NTDB id 1402 DSB67 RS12670 WP 010643257.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATS	311
NTDB id 1112 NGFG RS09215 WP 003689811.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATS	314
NTDB id 1113 AAA85695.1 219..1451()	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATS	314
NTDB id 1252 GC085 RS07730 WP 011213805.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATS	311
NTDB id 1198 PSJM300 03950 AFN76868.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATS	310
NTDB id 1016 ACIAD RS01680 WP 004920476.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATS	313
NTDB id 1059 ABD1 RS01610 WP 000279215.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATS	313
consensus	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATS	

