

logo

NTDB id 251722 FORC73 RS02485 WP 000648511.1	MAKNGGFSLEAKKEKRFIFEGRHSSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	72
NTDB id 1169 A1552VC RS11075 WP 000648511.1	MAKNGGFSLEAKKEKRFIFEGRHSSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	72
NTDB id 1402 DSB67 RS12670 WP 010643257.1	MAKNGGFSLEAKKEKRFIFEGRHSSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	71
NTDB id 1112 NGFG RS09215 WP 003689811.1	MAKNGGFSLEAKKEKRFIFEGRHSSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	74
NTDB id 1113 AAA85695.1 219..1451()	MAKNGGFSLEAKKEKRFIFEGRHSSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	74
NTDB id 1252 GC085 RS07730 WP 011213805.1	MAKNGGFSLEAKKEKRFIFEGRHSSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	71
NTDB id 1198 PSJM300 03950 AFN76868.1	MAKNGGFSLEAKKEKRFIFEGRHSSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	70
NTDB id 1016 ACIAD RS01680 WP 004920476.1	MAKNGGFSLEAKKEKRFIFEGRHSSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	73
NTDB id 1059 ABD1 RS01610 WP 000279215.1	MAKNGGFSLEAKKEKRFIFEGRHSSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	73
consensus	MAKNGGFSLEAKKEKRFIFEGRHSSASDKLVNGEVSATFEEEARKKLAKRGIRPLQITRV....KTSSKRKITQEDITV	

logo

NTDB id 251722 FORC73 RS02485 WP 000648511.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPFVF	152
NTDB id 1169 A1552VC RS11075 WP 000648511.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPFVF	152
NTDB id 1402 DSB67 RS12670 WP 010643257.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPFVF	151
NTDB id 1112 NGFG RS09215 WP 003689811.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPFVF	154
NTDB id 1113 AAA85695.1 219..1451()	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPFVF	154
NTDB id 1252 GC085 RS07730 WP 011213805.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPFVF	151
NTDB id 1198 PSJM300 03950 AFN76868.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPFVF	150
NTDB id 1016 ACIAD RS01680 WP 004920476.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPFVF	153
NTDB id 1059 ABD1 RS01610 WP 000279215.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPFVF	153
consensus	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPFVF	

logo

NTDB id 251722 FORC73 RS02485 WP 000648511.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMFKGFGAELPWFTQQVVKLSHWVQAYSLWAFI	232
NTDB id 1169 A1552VC RS11075 WP 000648511.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMFKGFGAELPWFTQQVVKLSHWVQAYSLWAFI	232
NTDB id 1402 DSB67 RS12670 WP 010643257.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMFKGFGAELPWFTQQVVKLSHWVQAYSLWAFI	231
NTDB id 1112 NGFG RS09215 WP 003689811.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMFKGFGAELPWFTQQVVKLSHWVQAYSLWAFI	234
NTDB id 1113 AAA85695.1 219..1451()	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMFKGFGAELPWFTQQVVKLSHWVQAYSLWAFI	234
NTDB id 1252 GC085 RS07730 WP 011213805.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMFKGFGAELPWFTQQVVKLSHWVQAYSLWAFI	231
NTDB id 1198 PSJM300 03950 AFN76868.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMFKGFGAELPWFTQQVVKLSHWVQAYSLWAFI	230
NTDB id 1016 ACIAD RS01680 WP 004920476.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMFKGFGAELPWFTQQVVKLSHWVQAYSLWAFI	233
NTDB id 1059 ABD1 RS01610 WP 000279215.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMFKGFGAELPWFTQQVVKLSHWVQAYSLWAFI	233
consensus	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMFKGFGAELPWFTQQVVKLSHWVQAYSLWAFI	

logo

NTDB id 251722 FORC73 RS02485 WP 000648511.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHFFETAIN	312
NTDB id 1169 A1552VC RS11075 WP 000648511.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHFFETAIN	312
NTDB id 1402 DSB67 RS12670 WP 010643257.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHFFETAIN	311
NTDB id 1112 NGFG RS09215 WP 003689811.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHFFETAIN	314
NTDB id 1113 AAA85695.1 219..1451()	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHFFETAIN	314
NTDB id 1252 GC085 RS07730 WP 011213805.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHFFETAIN	311
NTDB id 1198 PSJM300 03950 AFN76868.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHFFETAIN	310
NTDB id 1016 ACIAD RS01680 WP 004920476.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHFFETAIN	313
NTDB id 1059 ABD1 RS01610 WP 000279215.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHFFETAIN	313
consensus	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPPIICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHFFETAIN	

