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NTDB id 139936 VAA049 RS02410 WP 000648510.1	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	

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NTDB id 139936 VAA049 RS02410 WP 000648510.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPPEVF	152
NTDB id 1169 A1552VC RS11075 WP 000648511.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPPEVF	152
NTDB id 1402 DSB67 RS12670 WP 010643257.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPPEVF	151
NTDB id 1112 NGFG RS09215 WP 003689811.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPPEVF	154
NTDB id 1113 AAA85695.1 219..1451()	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPPEVF	154
NTDB id 1252 GC085 RS07730 WP 011213805.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPPEVF	151
NTDB id 1198 PSJM300 03950 AFN76868.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPPEVF	150
NTDB id 1016 ACIAD RS01680 WP 004920476.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPPEVF	153
NTDB id 1059 ABD1 RS01610 WP 000279215.1	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPPEVF	153
consensus	LTRQLATMTTGVPIVQALKLVGDNHRKAEMKSLAQITKSVEACTPLSKAMRTASAHFDTLYVDLVETGEMSGNLPPEVF	

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NTDB id 1112 NGFG RS09215 WP 003689811.1	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMFKGFGAELPWFTQQVVKLSHWVQAYSLWAFI	234
NTDB id 1113 AAA85695.1 219..1451()	ERLATYREKSEQLRAKVIKALITYPSMVVLVALGVSYLMMLTVIPEFESMFKGFGAELPWFTQQVVKLSHWVQAYSLWAFI	234
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NTDB id 139936 VAA049 RS02410 WP 000648510.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHVFETAIN	312
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NTDB id 1198 PSJM300 03950 AFN76868.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHVFETAIN	310
NTDB id 1016 ACIAD RS01680 WP 004920476.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHVFETAIN	313
NTDB id 1059 ABD1 RS01610 WP 000279215.1	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHVFETAIN	313
consensus	AIAAAIFGLKALRNKNSFQIRLKTSLRLGLKFPICNVLAKASIAKFSRTLATSFAACGPIILASLKITAKTSGNVHVFETAIN	

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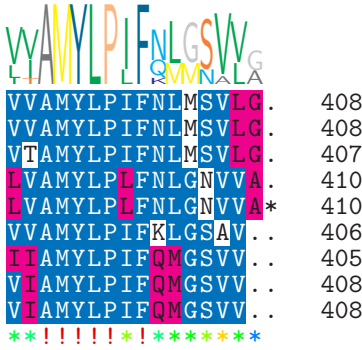
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NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus



☐ non conserved
☒ similar
☒ ≥ 50% conserved