



Aedes taeniorhynchus mosquito rRNA gene internal transcribed spacer 2 (ITS2) unanimity alignment & molecular taxonomy primer map

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1 | 130 |
MG232636 | ACAAGTTGAACGCATATTGCACATCGTACAACAGTACGATGTACACATTTTTGAGTGCCTATATTTATCCATTCAACTATACGTGTGCGTGCGCGTCCCACCTTCGGGTGGACAGGCGCACGGGCTCCCC |

131 | Mosq-RAE2 | 260 |
MG232636 | GCGTGCTCTCTCTCCGGAGAGCGCGGGGCA CGTATGCRKVGTTGAYGTTTTTC CGTATGCGTAGTGACGTTTTTC CCGCTCCGCCGCGAGCGATAAAACGTTGAAGATAGTCAGGCGGCCTCGTGCGGGCTGGTGAATACATCCCCTAGG |

261 | 316 |
MG232636 | CCTGCTCTCGCGAAGGCTATGTTGTATTCCACCGCGAGGGACGTTATCTCAGTAG |
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Alignments prepared using ElimDupes (www.hiv.lanl.gov/content/sequence/elimdupesv2/elimdupes.html), CLUSTAL O (www.ebi.ac.uk/Tools/msa/clustalo/), EMBOSS Seqret (www.ebi.ac.uk/Tools/sfc/emboss_seqret/) and SURE v1.0 Sequence Unanimity Reformatting tool (<http://midasmap.uaslp.mx/suretool/>).