



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

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1
|
MG232611 CGTACGAGGTACACATTTTTGAGTGCCTATATTTAACGATTCAACTATACGTGCATGCGCGTTTCACTCCGGGTGGACAGGCGCAGGGCCCATGAGCA CGTATGCRKVGTGAYGTTTTTC CGTATGCGTAGTGACGTTTTTC CCGCCTTCGGT
|
Mosq-RAE2 130

131
|
MG232611 GCGCGGTAAAACGTTCAAGATAGTCAGGCGCGTTCCTTCGCGGACGCGGTTGATGAATACATCCCATACACCCCATGCGCTTCGGCCGGGTATGTTGTATTCATCACCAACAACA
|
249
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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/>).