



## Anopheles albimanus mosquito rRNA gene internal transcribed spacer 2 (ITS2) unanimity alignment & molecular taxonomy primer map

Laboratorio de Genómica Viral y Humana, Facultad de Medicina, Universidad Autónoma de San Luis Potosí, México (08/Apr/2022, v4.0)





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Consensus L78065	CCAAACGTACCCTGAAGCAACGGCTCGAACGGTGTGCGACTCATCGAGCGCACGATCACCCGAACCCGATGATCAAGTGGCCTCAAATA		ATGTGTGACTACCCCTAAATTTAAGCAT
GU477268	-----		
GU477267	-----		
GU477269	-----		
GU477274	-----		
GU477270	-----		
JX212820	-----		*****
GU477271	-----		*****

Alignments prepared using ElimDups ([www.hiv.lanl.gov/content/sequence/elimdupserv2/elimdups.html](http://www.hiv.lanl.gov/content/sequence/elimdupserv2/elimdups.html)), CLUSTAL O ([www.ebi.ac.uk/Tools/msa/clustalo/](http://www.ebi.ac.uk/Tools/msa/clustalo/)), EMBOSS Seqret ([www.ebi.ac.uk/Tools/sfc/emboss\\_seqret/](http://www.ebi.ac.uk/Tools/sfc/emboss_seqret/)) and SURE v1.0 Sequence Unanimity Reformatting tool (<http://midasmap.uaslp.mx/suretool/>).