

1 Mosq-F 130

Consensus TGTGAAC TG CAGG ACAC AT
TGTGAAC TG CAGG ACAC AT GAAC ACCG GACA AG TT GAAC GCAT ATT GCAC AT CGTACA AAC AGTAC GAT GTAC A CATT TT GAG TGC CTAT ATT TAT CTATT CA ACT GT GT CGG CACC ACCT CGACT GGT GC

U22138 -----
U22137 ----- G -----
U22135 -----
U22136 ----- C -----
U22134 -----
U22133 -----

131 Mosq-RCX 260

Consensus CGGCACGCAGAATGGTGTGCTGCCTTCGGTG GCTGGCAAAACATTCAAGAC
GCTGGCAAAACATTCAAGA TACTCCCGTTGGCGGTTGC GGCG ACCAGCTAACACTGGTGC GCGACACACAGAC GCGTGAGAATA CAT CCCA

U22138 ----- • -----
U22137 ----- C -----
U22135 ----- • -----
U22136 ----- • -----
U22134 ----- • -----
U22133 ----- C ----- • -----

261 390

Consensus CACACCAACCTGGCTGGCGACGATGTAACCTCTCACACTCCTGCGTTGACGTGTCGCGTACCGGGTCTCGCAACCAACCGTGCACACACCGCATAGATAAACCCCCATGTAGGCCTCAAATAA

U22138 -----
U22137 -----
U22135 ----- C · · · · · - A - ATG -
U22136 ----- C · · · · · - A - ATG -
U22134 ----- A ----- G ----- G -----
U22133 ----- G ----- G -----



Culex restuans 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)

391 Mosq-R2 418

Consensus	TGTGTGACTACCCCTAAATTTAAGCAT
U22138	
U22137	
U22135	
U22136	
U22134	
U22133	