

SUPPLEMENTARY TABLE 1. Primers used for *Lb2169*-LRV sequencing. Forward (f) and reverse (r) primer sequences were designed from the LRV infecting the following strains (as indicated in the “strain” column): *L. guyanensis* M4147 and CUMC-1, as well as *Lb2169* itself after partial sequencing. Primer position is indicated relative to the complete *LgM4147*-LRV sequence (5283bp).

Primer	Position	Strain	Sequence
1f	236-255	<i>LgM4147</i>	TTG TAT CCA GGT TAC TGC CG
2f	1955-1974	<i>Lb2169</i>	CAG GTT ACG CCC GTT CAA TG
3f	2091-2113	<i>LgM4147</i>	TGG GGC AGT ACG TCA TGT GAC CT
3r	2091-2113	<i>LgM4147</i>	AGG TCA CAT GAC GTA CTG CCC CA
4f	2704-2726	<i>Lb2169</i>	ATT AGT GGT ATT TTG GAT CTA TC
5r	2796-2816	<i>LgM4147</i>	GCT GGC ATC TGG GAA AAG AAG
6f	3614-3641	<i>Lb2169</i>	GTG ATT TCT GGT CGT CAC GTT GGT TGT G
7r	4250-4376	<i>Lb2169</i>	CTA AAC ACC TGC TTA CTA GCA TTT AGC C
8f	4828-4854	<i>Lb2169</i>	GGA TCG GAT ATT GCT GAT ATC ATG GCC
9r	5255-5277	<i>LgCUMC-1</i>	TTA TGA CAC TTT CTG CCA TAT TCC