

SI Table 1. Additional data describing ML and BI analyses conducted.

		Mean support		
		value	Std. Dev. bs(ML) stdev. splfr (BI)	-lnL
ML				
<i>Sequence</i>				
	G1	99.07	0.84	-223665.52
	G2	98.36	0.16	-269023.15
	G1+G2	99.10	1.72	-290961.06
<i>Combined</i>				
	G1+RGC	99.07	0.82	-223882.83
	G2+RGC	98.40	0.27	-269358.13
	G1+G2+RGC	91.88	1.79	-291612.57
BI				
<i>Sequence</i>				
	G1	1.0	0.010476	-227764.13
	G2	1.0	0.012298	-275775.33
	G1+G2	0.99	0.002266	-302944.15
<i>Combined</i>				
	G1+RGC	0.97	0.007857	-230513.31
	G2+RGC	1.0	0.00739	-276119.53
	G1+G2+RGC	1.0	0.011224	-303590.55

SI.T.1 Data describing ML and BI analyses and resulting phylogenomic trees.

G1 = Group 1; G2 = Group 2; RGC = Rare Genomic Changes.