S3 Table Summary of Max $\chi^2,$ GARD and PHI based on the ITS and BT2 data

Group	Max chi ²		GARD		PHI	
	ITS	BT2	ITS	BT2	ITS	BT2
C.carrionii	13.71 (p = 0.03)	17.75 (p = 0.02)	-	+	P = 0.30	P = 0.02
Fonsecaea spp	25.62 (p = 0.00)	9.32 (p = 0.46)	-	-	P = 0.82	P = 0.21

in Cladophialophora carrionii_and Fonsecaea spp.

GARD: genetic analysis for recombination detection PHI: the pairwise homoplasy index SplitsTree 4