S4 Table. Multiple comparisons for the generalized linear model of the topology weighting of the Cbp_{Cg} subgenome of *C. bursa-pastoris* and *C. rubella*.

Comparison	Estimate	Std. Error	z value	p-value
ASI_CR - species	-3.711	0.022	-166.33	0.0000
ASI_CR - EUR_CR	-1.425	0.016	-90.97	0.0000
ASI_CR - ME_CR	-0.797	0.016	-50.48	0.0000
EUR_CR - species	-2.286	0.021	-107.07	0.0000
EUR_CR - ME_CR	0.628	0.014	43.65	0.0000
ME_CR - species	-2.914	0.021	-135.94	0.0000

With the binomial family, the null deviance was 274261 on 198584 degrees of freedom, and residual deviance of 225829 on 198580 degrees of freedom.