

**Table S1:** Summary statistics for transcriptome profiling of *Chrysoporthe austroafricana*

Samples	RIN	Total PE Reads	% Mapped reads to <i>C. austroafricana</i>			% Mapped reads to <i>E.grandis</i> genome	Q20 Percentage (%)	GC Percentage (%)	Expressed genes
			Paired	Singletons	Total				
1. Complete Media_BR1	7.9	32.141161	94.0	1.9	95.9	-	98.1	56.84	12477

2. Complete Media_BR2	7.6	30.008628	94.5	1.9	96.4	-	98.14	56.38	12434
3. Complete Media_BR3	7.8	34.342786	94.3	1.8	96.1	-	98.07	57.19	12477
4. Minimal media_BR1	9.1	33.951504	93.3	2.0	95.4	-	98.03	57.19	12498
5. Minimal media_BR2	8.7	34.477061	93.8	1.9	95.7	-	98.06	57.16	12572
6. Minimal media_BR3	6.9	34.256842	94.3	1.9	96.2	-	98.04	57.07	12520
7. TAG5 Infected_BR1	9.6	37.617103	1.6	0.5	2.1	75.5	97.75	48.67	9166
8. TAG5 Infected_BR2	9.5	37.390551	2.7	0.5	3.2	76.9	97.58	49.31	9609
9. TAG5 Infected_BR3	9.5	38.684116	1.5	0.4	1.9	78.6	97.66	49.41	9067
10. ZG14 Infected_BR1	9.7	38.062937	2.4	0.5	2.9	75	98.13	49.33	9412
11. ZG14 Infected_BR2	9.6	34.373634	2.7	0.6	3.2	76.2	98.18	48.36	9289
12. ZG14 Infected_BR3	9.7	36.626103	2.7	0.5	3.2	75.8	98.13	48.77	9500

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