

Supplementary Table S3. *in silico* cDNA library of assembled *M. anisopliae* JEF-197 and Japanese pine sawyer

	<i>M. anisopliae</i> JEF-197	Japanese pine sawyer
Total length of transcripts (bp)	38,664,210	71,371,911
Total number of transcripts	35,334	52,306
Total number of genes	9,286	19,046
N50 of genes (bp)	1,500	2,007
GC %	54.19	41.74