



**Figure S1** No evidence for aneuploidy in any sequenced sample. The median expression value (log<sub>10</sub>-transformed RPKM) of each of the eight chromosomes in the *C. albicans* genome is plotted for each of the 96 sequenced transcriptome samples. Samples are grouped by time point (column) and replicate (row). None of the samples displayed observable deviations from uniformity between the chromosomes, which would have otherwise indicated a possible presence of aneuploidy.