Agreement between expression profiles calculated based on reference transcripts and trinity components at different *S. pombe* samples. (a) A heatmap showing the expression profiles for the 430 transcripts (rows) that were differentially expressed between the samples (columns) based on reference transcripts (Ref) and reciprocally-best-matched Trinty components (Trin). Genes and samples are clustered. (b) Gene expression correlation matrix between the expression profiles of each pair of samples calculated based on the reference transcripts (REF) or the matching Trinity components (TRIN).