Figure S11. Pairwise comparison of the expression of all expressed genes and Fip1-APA target genes during ESC EB differentiation time course. The expression data for the EB differentiation time course was downloaded from GEO (accession GSE3749) and processed as described in the methods. The mean log2 Fold-Change was calculated by contrasting each time point against 0 hours (the ESC state), for all expressed genes or Fip1 APA target genes. The p-values were calculated by Wilcoxon rank sum test.