



Figure 1: Reconstructed TFAs of 11 yeast cell cycle related TFs from four sub-networks, by gNCA. The first three columns are for the wild type with synchronization methods of elutriation, α -factor and *cdc15* respectively, and the last column is for Fkh1 Fkh2 mutant. In each plot, TFAs reconstructed from all the 4 sub-networks are plotted, with the solid, dotted, dashdot and dashed lines representing for sub-network 1, 2, 3, and 4 respectively.