S10 Fig. Dendogram representation for Deng et al (2014) developmental phase single cell data.

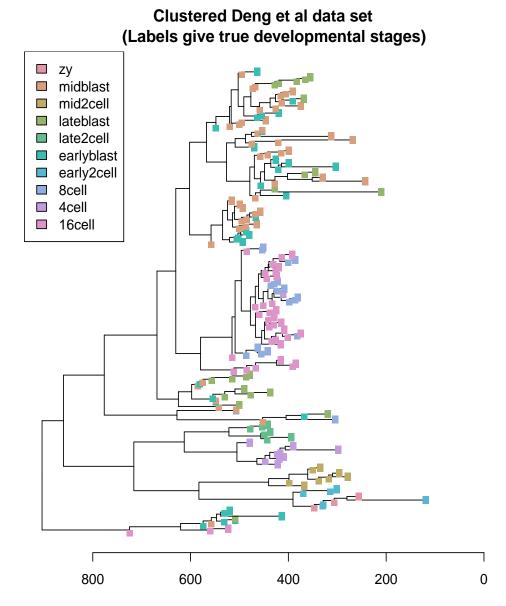


Fig 1. Dendogram representation of the 259 cells collected across different developmental phases by Deng et al (2014). The hierarchical clustering in this case has been performed on the log counts per million (cpm) data. Note that the labels in this case are difficult to interpret and also the hierarchical clustering fails to represent the continuum among the different developmental phases as in PCA or the GoM model.