



Fig. S4 Differentiation regulatory network. **(a)** Network reconstruction pipeline. **(b)** Number of interactions in the gene regulatory network by DEG major groups colored by the TF expression profile. **(c)** Percentage of interactions in the gene regulatory network by DEG major groups colored by the TF expression profile. **(d)** Distribution of the number of targets each TF is associated to. **(e)** Distribution of the number of TF regulators each DEG is associated to.