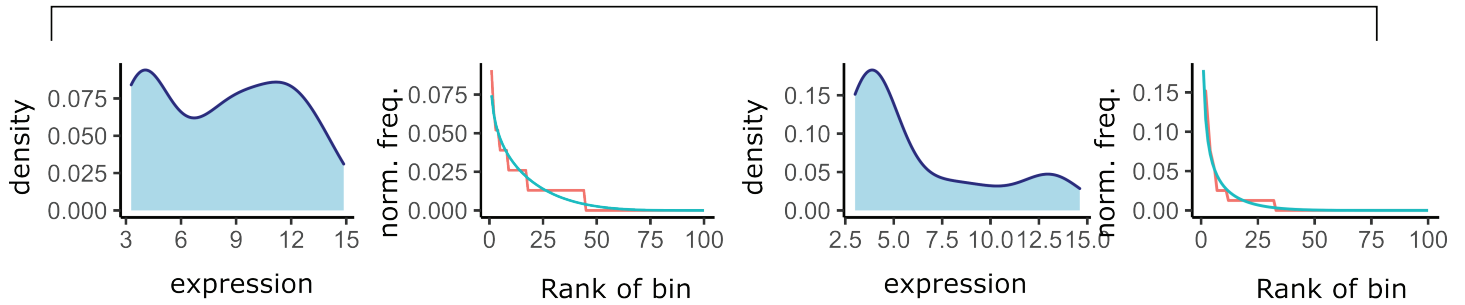
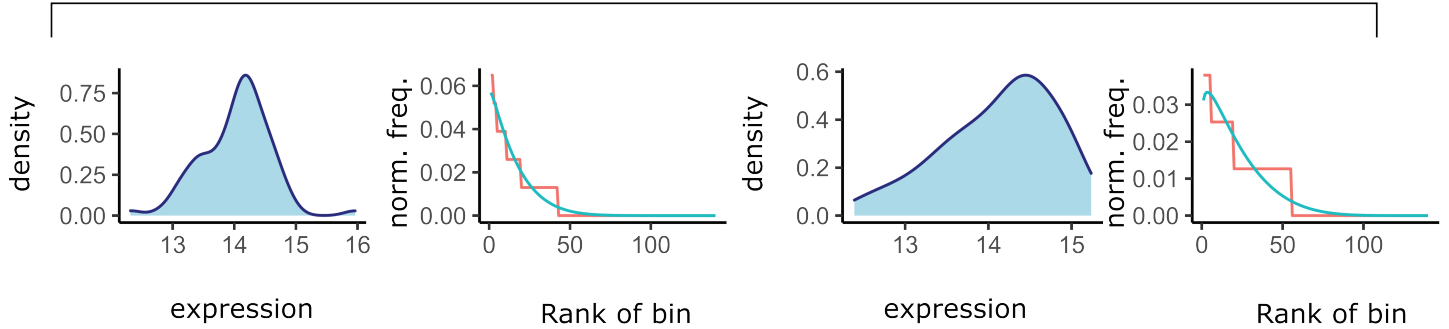


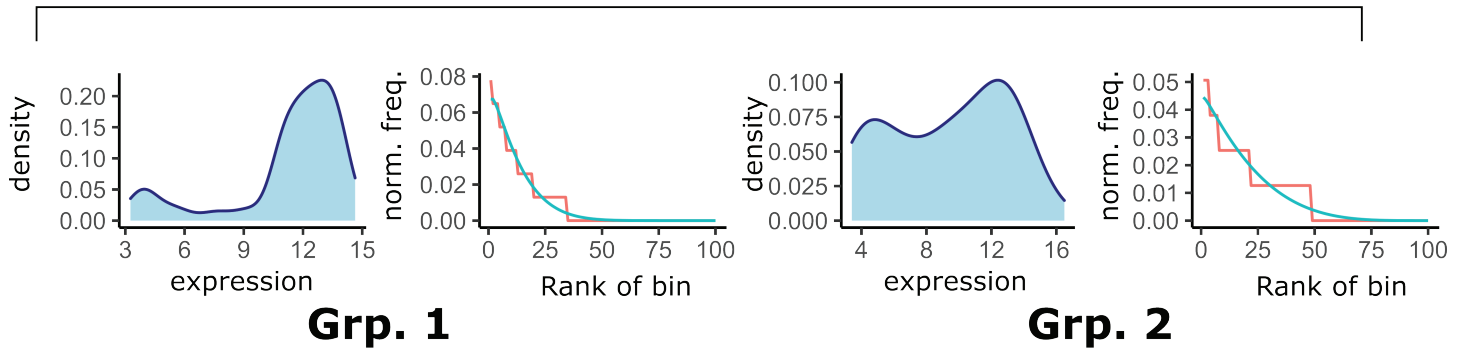
EPHA2



ATP5E



INIP



Supplemental Figure S2. Five randomly selected differential genes between myoblasts sampled before/24 hours after differentiation (Source: Trapnell data). Differential expression analysis was performed using ROSeq. Each row contains cell-group-wise expression density plot as well as a plot depicting DGBD based modeling of rank-ordered expression bins.