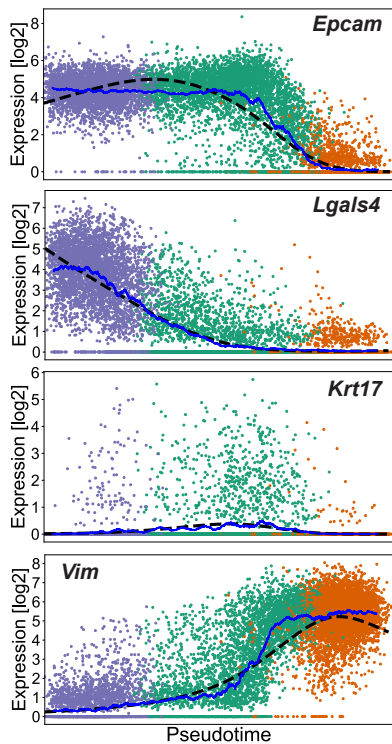
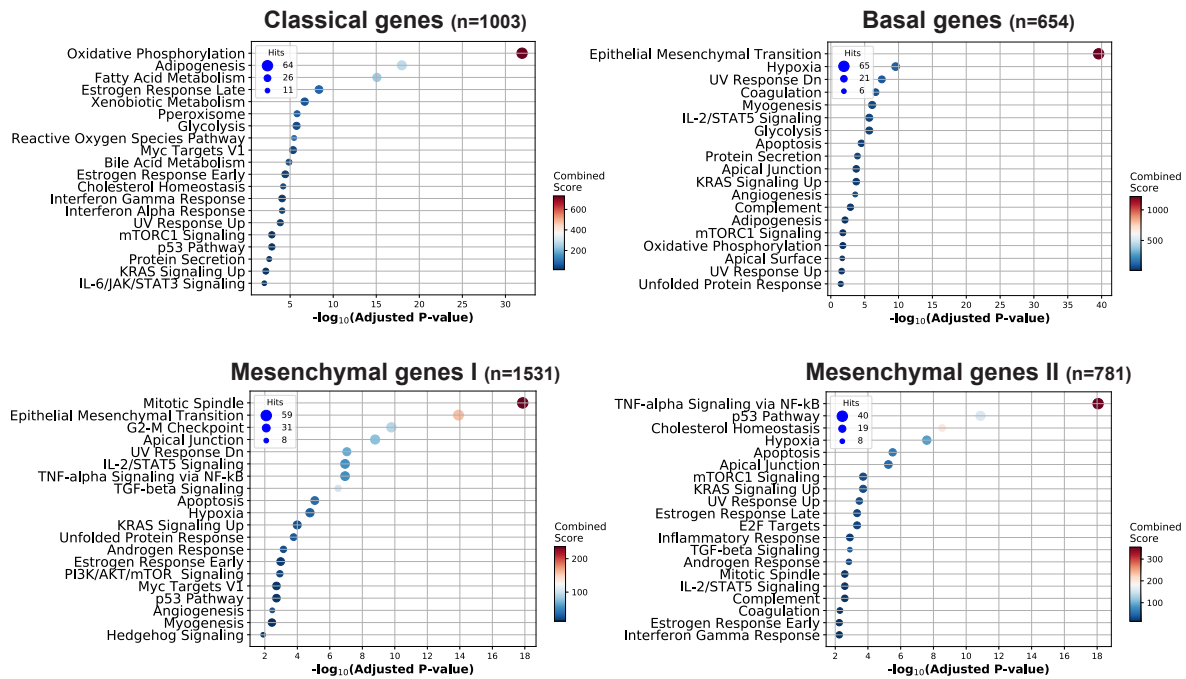


D Examples of gene expression modeled along pseudotime



E Gene set enrichment analysis of gene clusters introduced in panel (C)



F Additional marker genes for classical, basal and mesenchymal cell states

