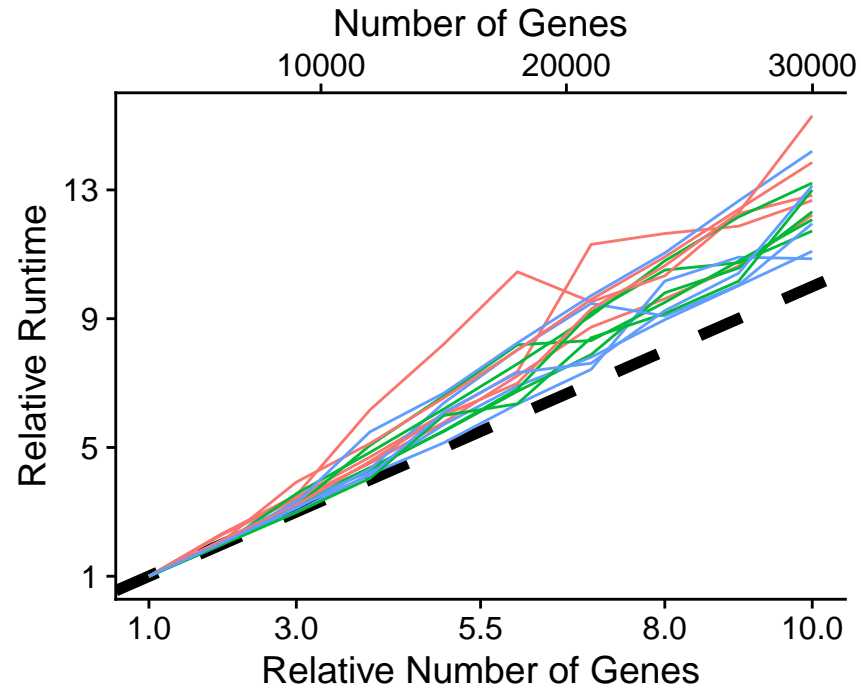
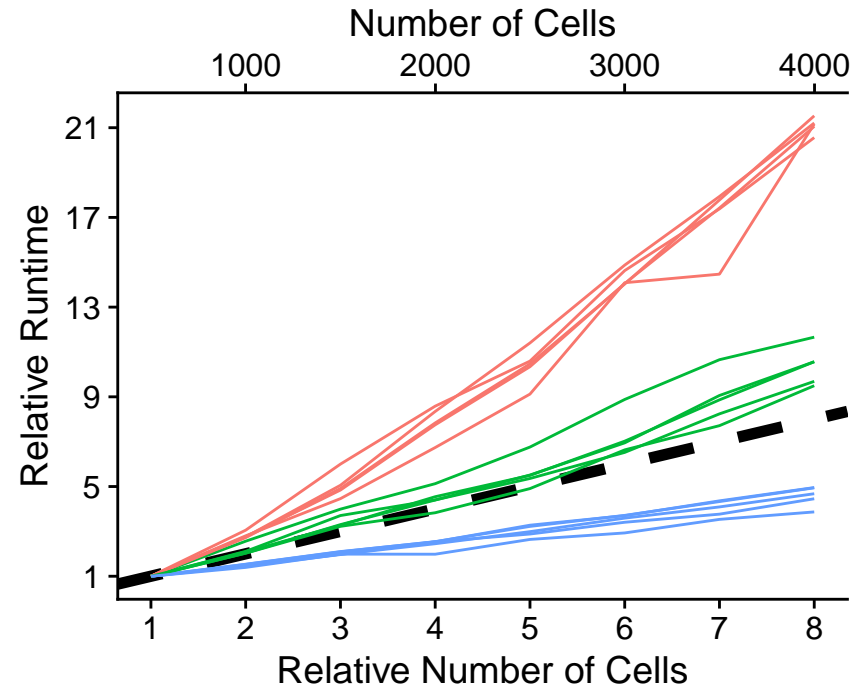
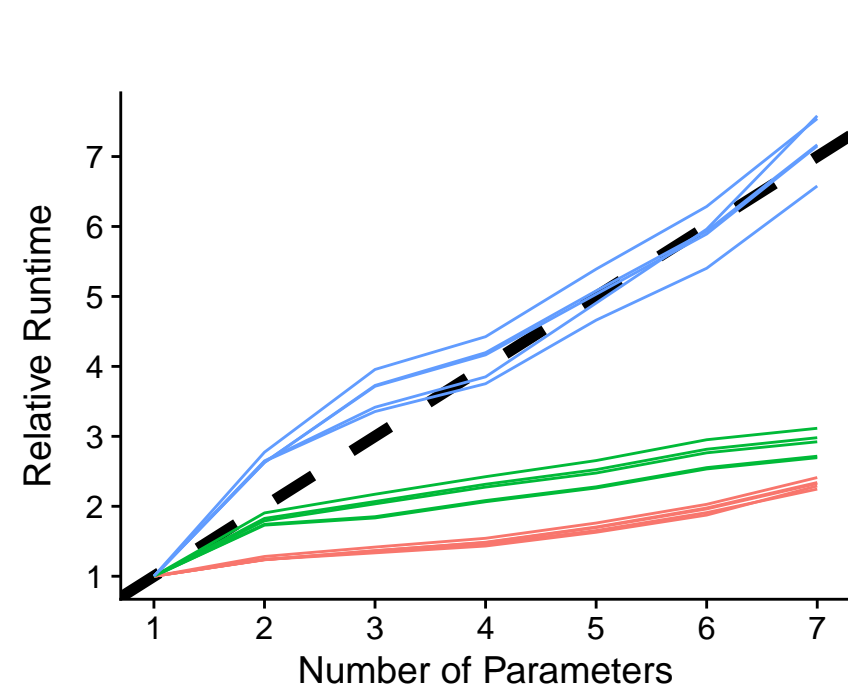


A**Runtime vs Number of Genes****B****Runtime vs Number of Cells****C****Runtime vs Number of Parameters**

DESeq2 edgeR glm_gp