



S9 Fig: Differentially expressed genes (DEGs) between bulk and single cell data for deletion strains with few DEGs (< 63 , 1% of yeast genome). Top panel is delta AUROC between bulk DEGs and single cell DEGs ordered by adjusted p-value. Bottom panel is delta Jaccard index for single-cell DEGs at FDR = 0.01 for each separate deletion strain. *preprocessed* is count normalized and log-transformed with no denoising method. Delta is taken between denoised and preprocessed.