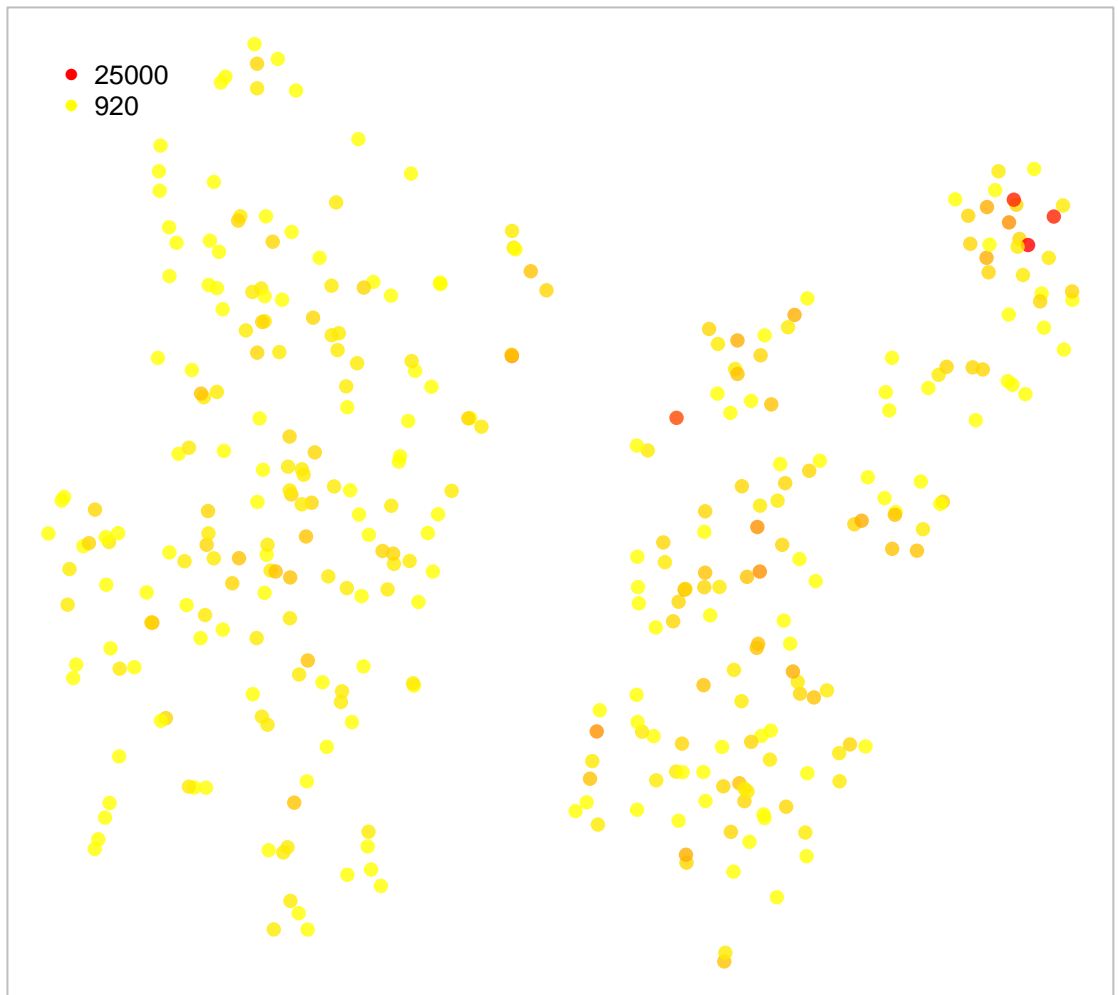
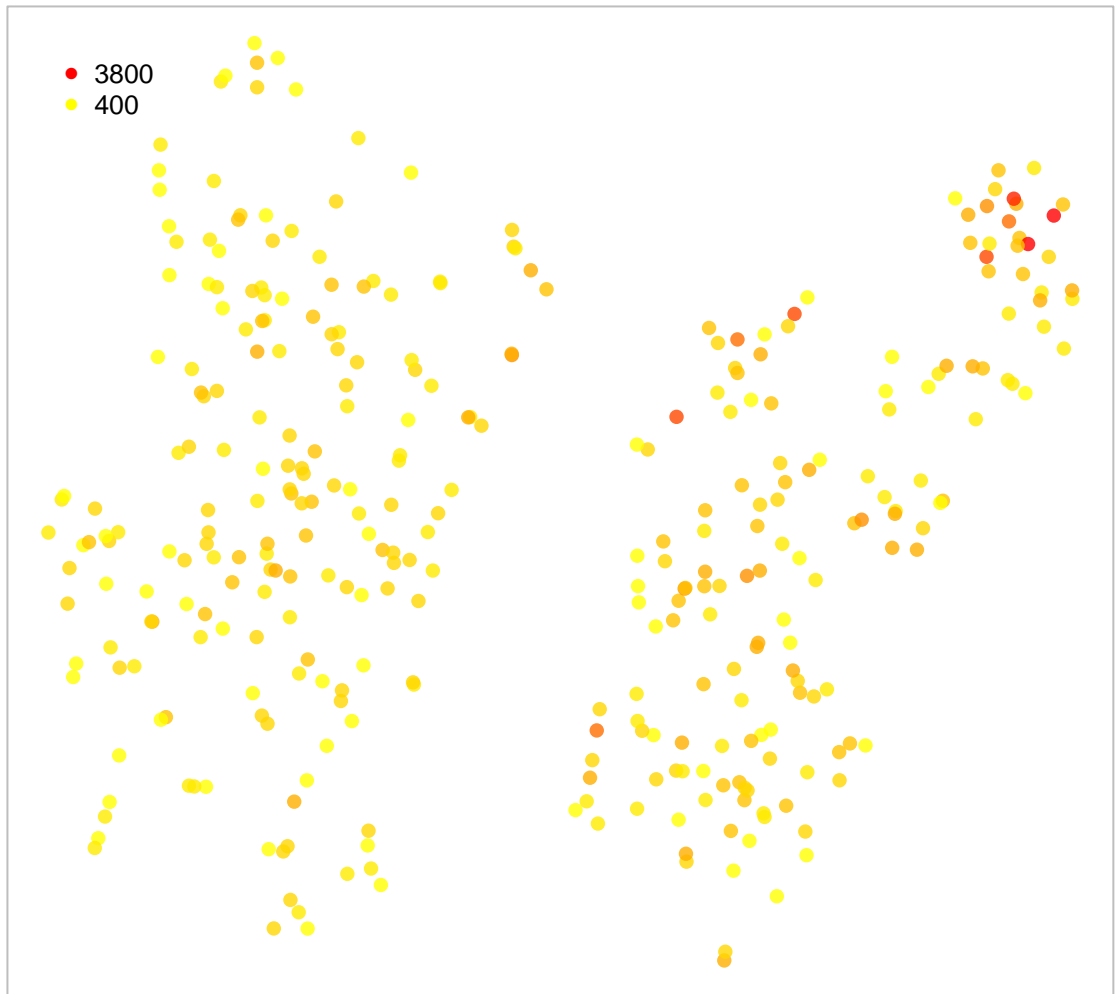


nCount_RNA



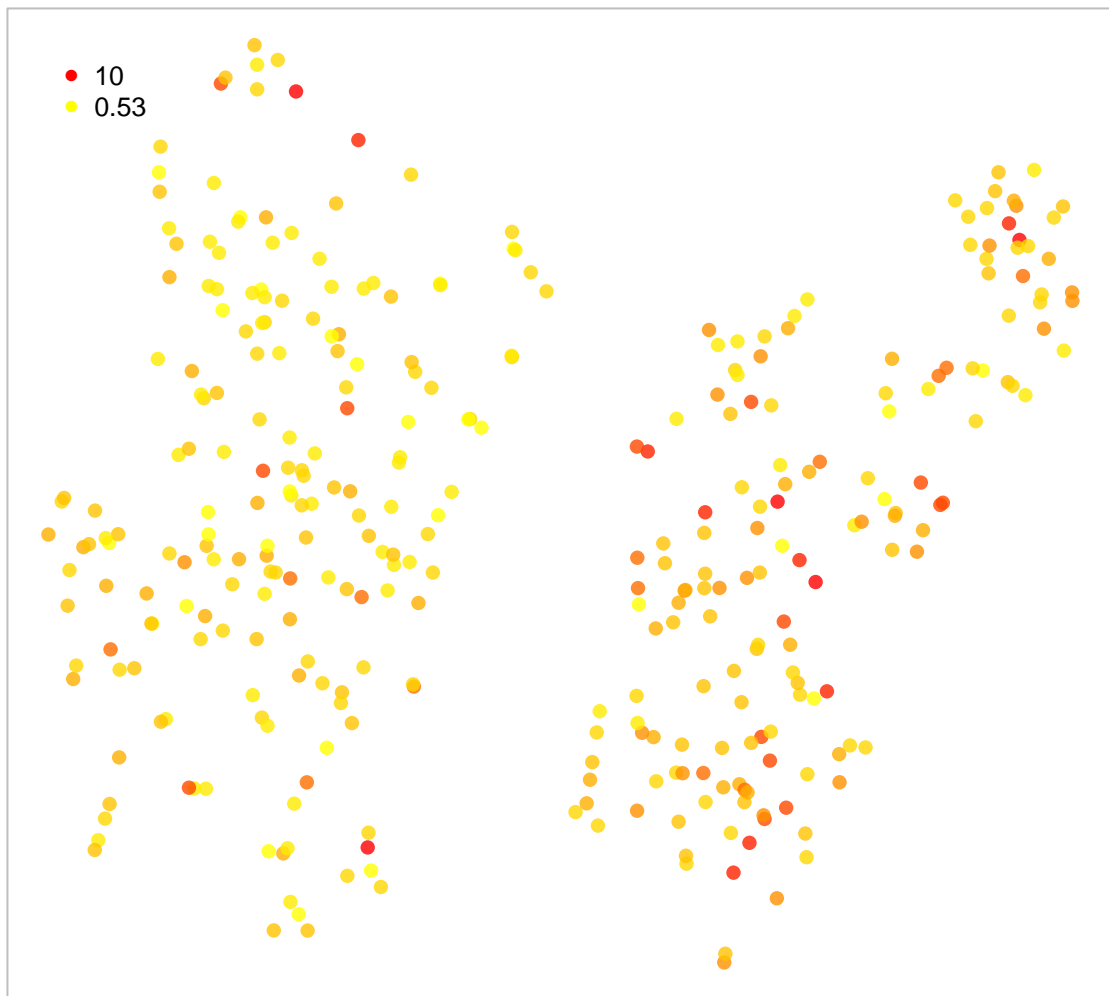
t-SNE on Binary 46 regulons (45PCs, 10 perplexity)

nFeature_RNA



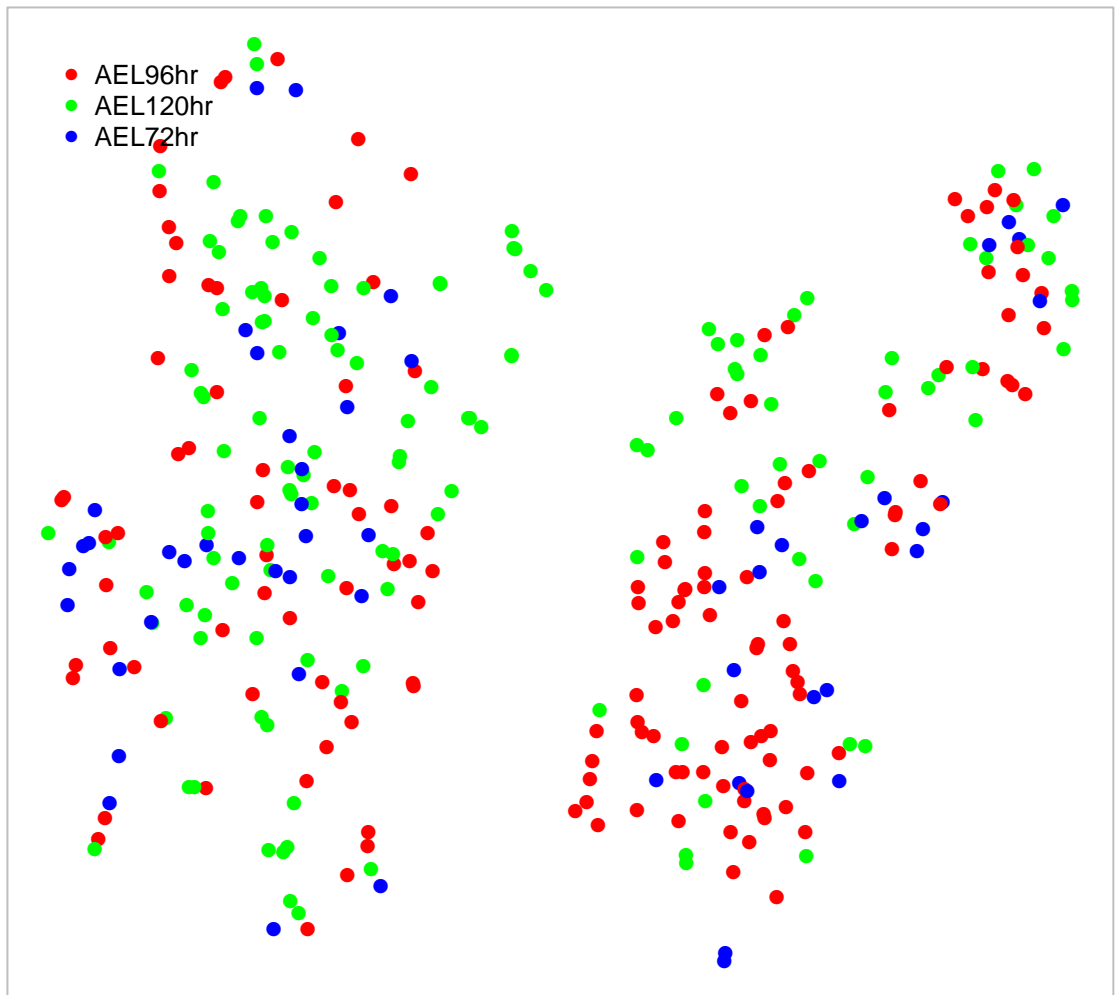
t-SNE on Binary 46 regulons (45PCs, 10 perplexity)

percent.mt



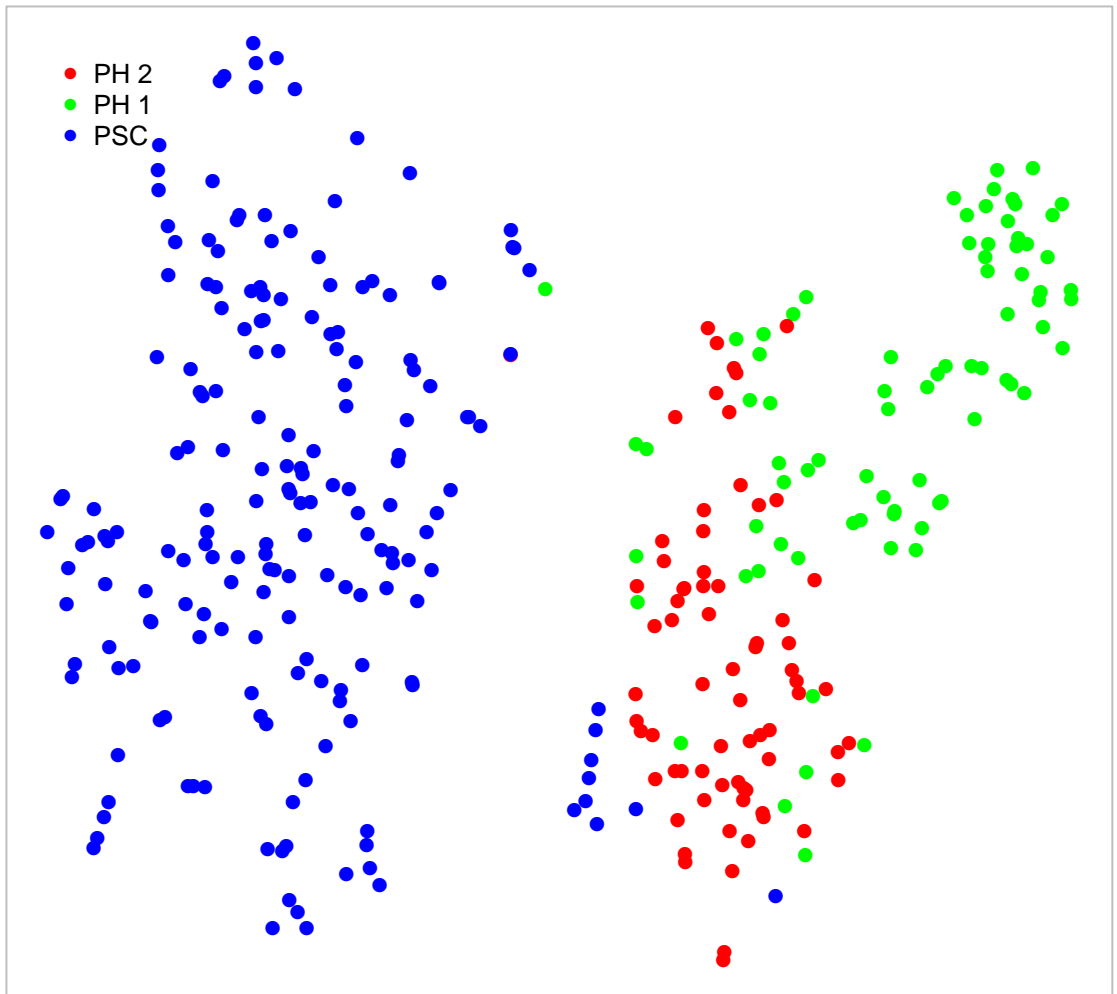
t-SNE on Binary 46 regulons (45PCs, 10 perplexity)

timepoint



t-SNE on Binary 46 regulons (45PCs, 10 perplexity)

new_subclustering



t-SNE on Binary 46 regulons (45PCs, 10 perplexity)