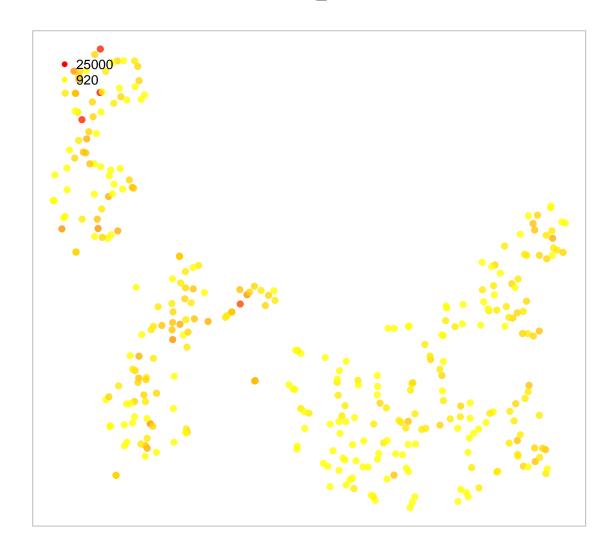
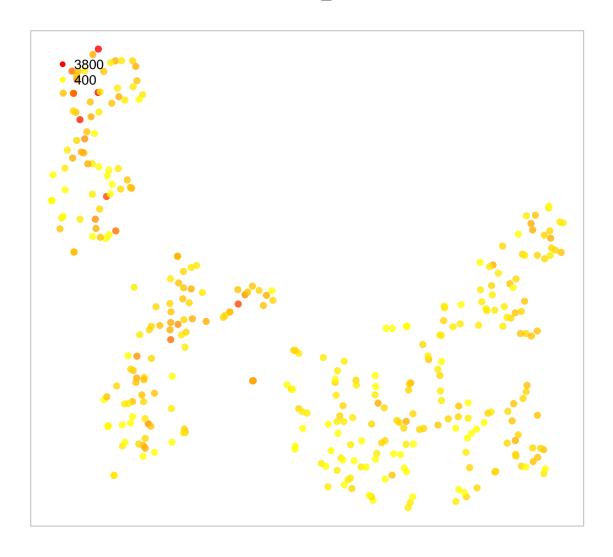
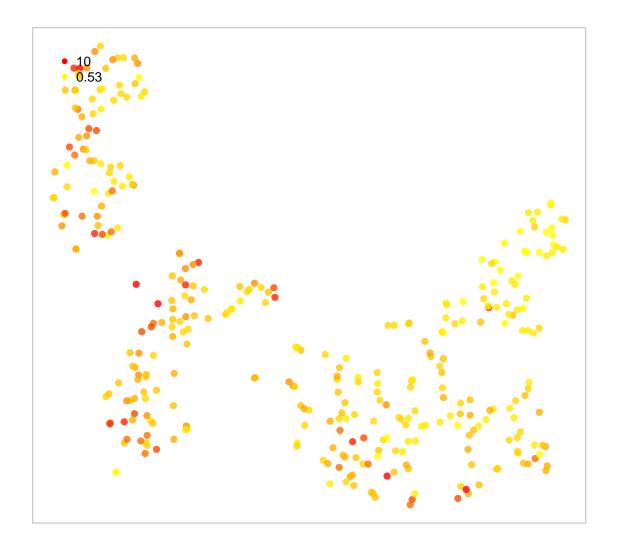
$n Count_RNA$



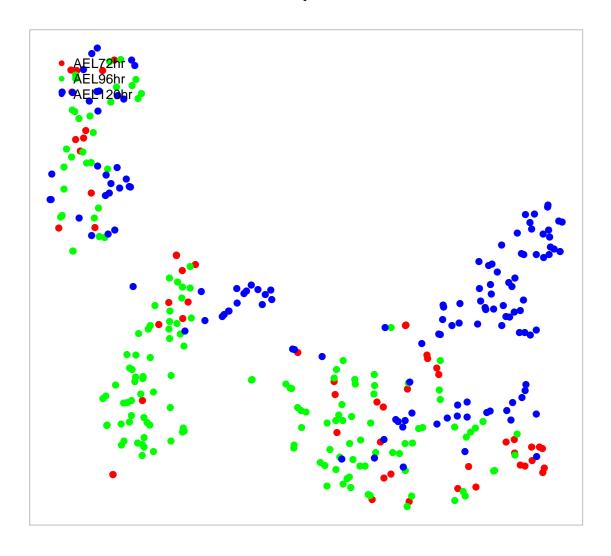
nFeature_RNA



percent.mt



timepoint



new_subclustering

