

seq depth vs gene count coloured by mt perc.

nFeature_RNA

10000

3000

1000

1e+03

1e+04

1e+05

nCount_RNA

- B_Cells
- CD4+_T_Cells
- CD8+_T_Cells
- DCIS 1
- DCIS 2
- Endothelial
- Invasive_Tumor
- IRF7+_DCs
- LAMP3+_DCs
- Macrophages_1
- Macrophages_2
- Mast_Cells
- Myoepi_ACTA2+
- Myoepi_KRT15+
- Perivascular-Like
- Prolif_Invasive_Tumor
- Stromal
- Stromal_&_T_Cell_Hybrid
- T_Cell_&_Tumor_Hybrid

