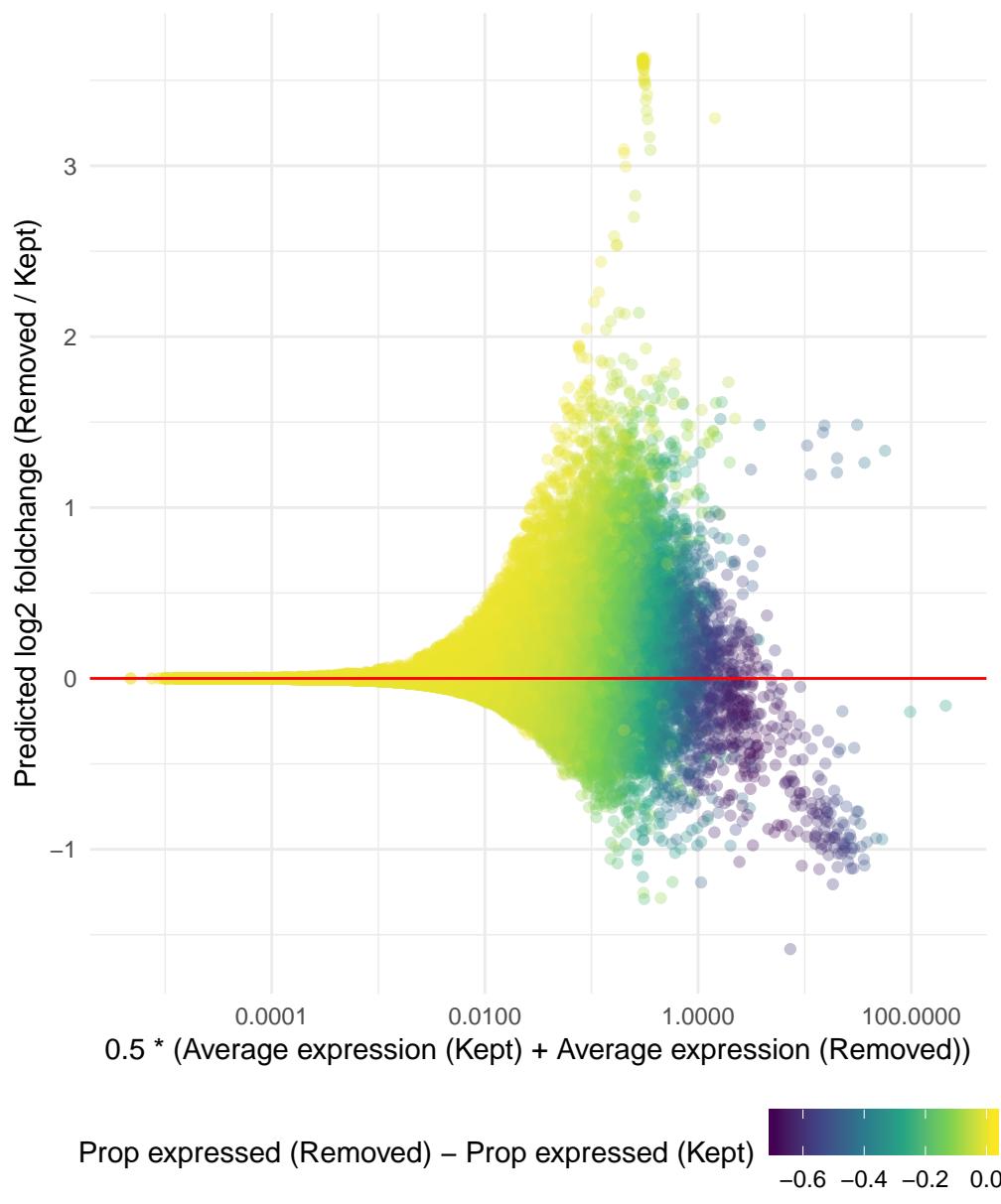
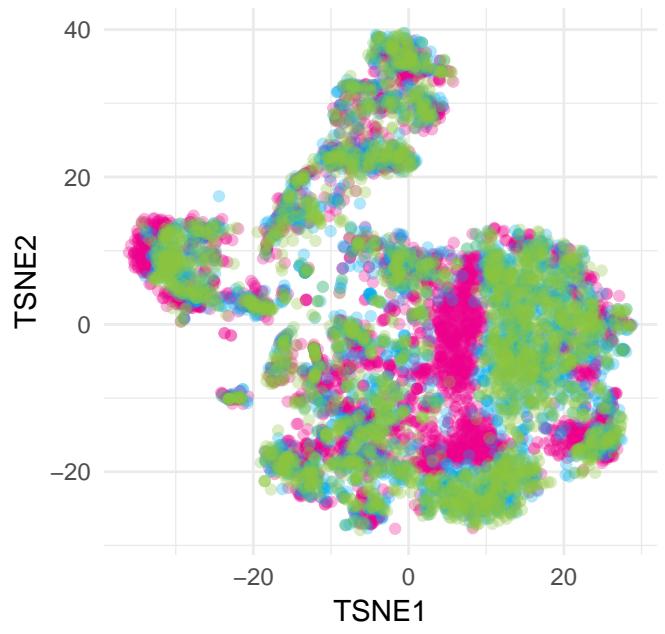
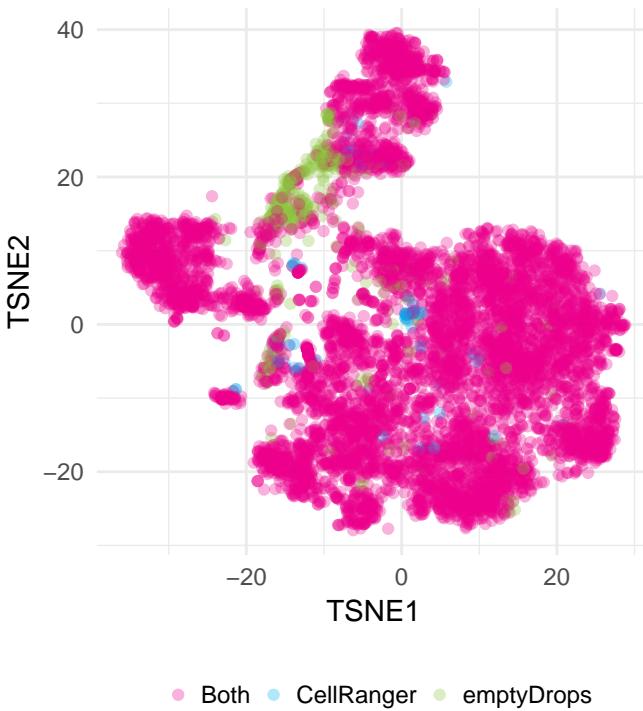


**A** Foldchange between removed and kept cells**B**

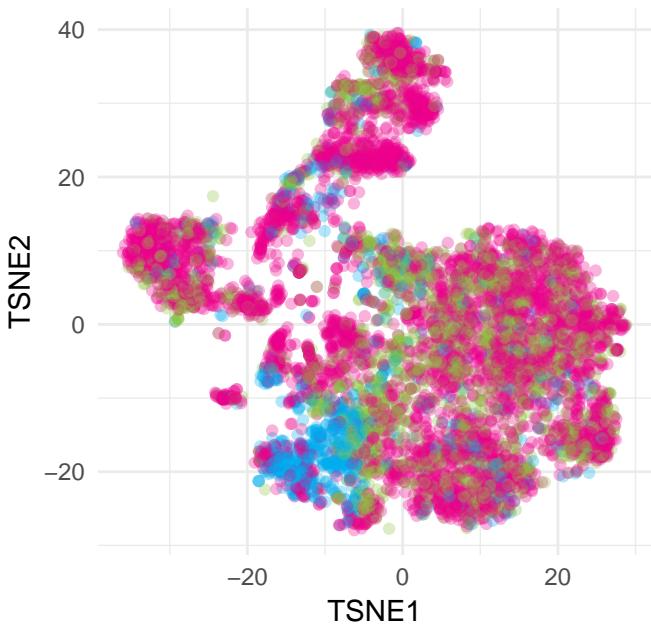
Gene	Predicted foldchange (log2)
AC005523.2	3.633061
OCLM	3.632253
LINC00843	3.625275
AL627309.1	3.624039
LINC02067	3.623951
AC025569.1	3.622870
ITPKA	3.622630
GCNT4	3.615836
AC010761.1	3.614835
LINC00921	3.614623
AC008738.2	3.609746
HSF4	3.608297
NR5A2	3.600864
AC091057.3	3.600186
AP000350.7	3.597255

**C** Sample

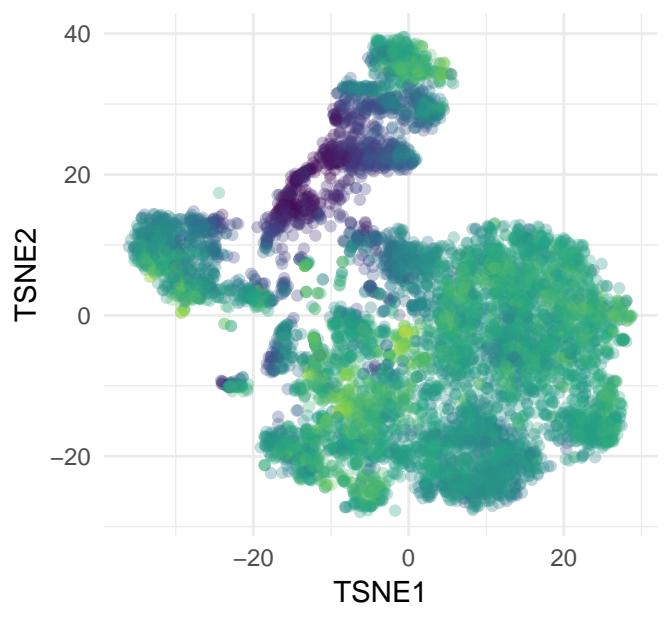
Selection method



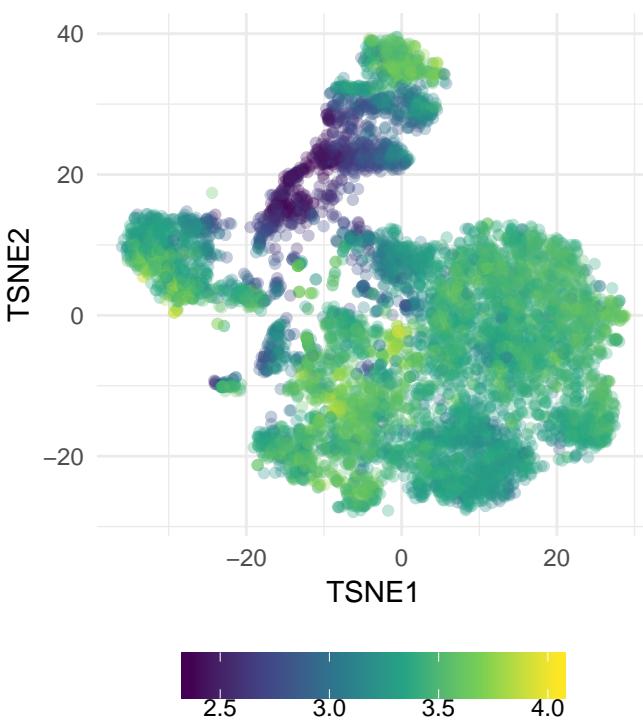
Cell cycle phase



Total counts (log10)



Total features (log10)



% Mitochondrial

