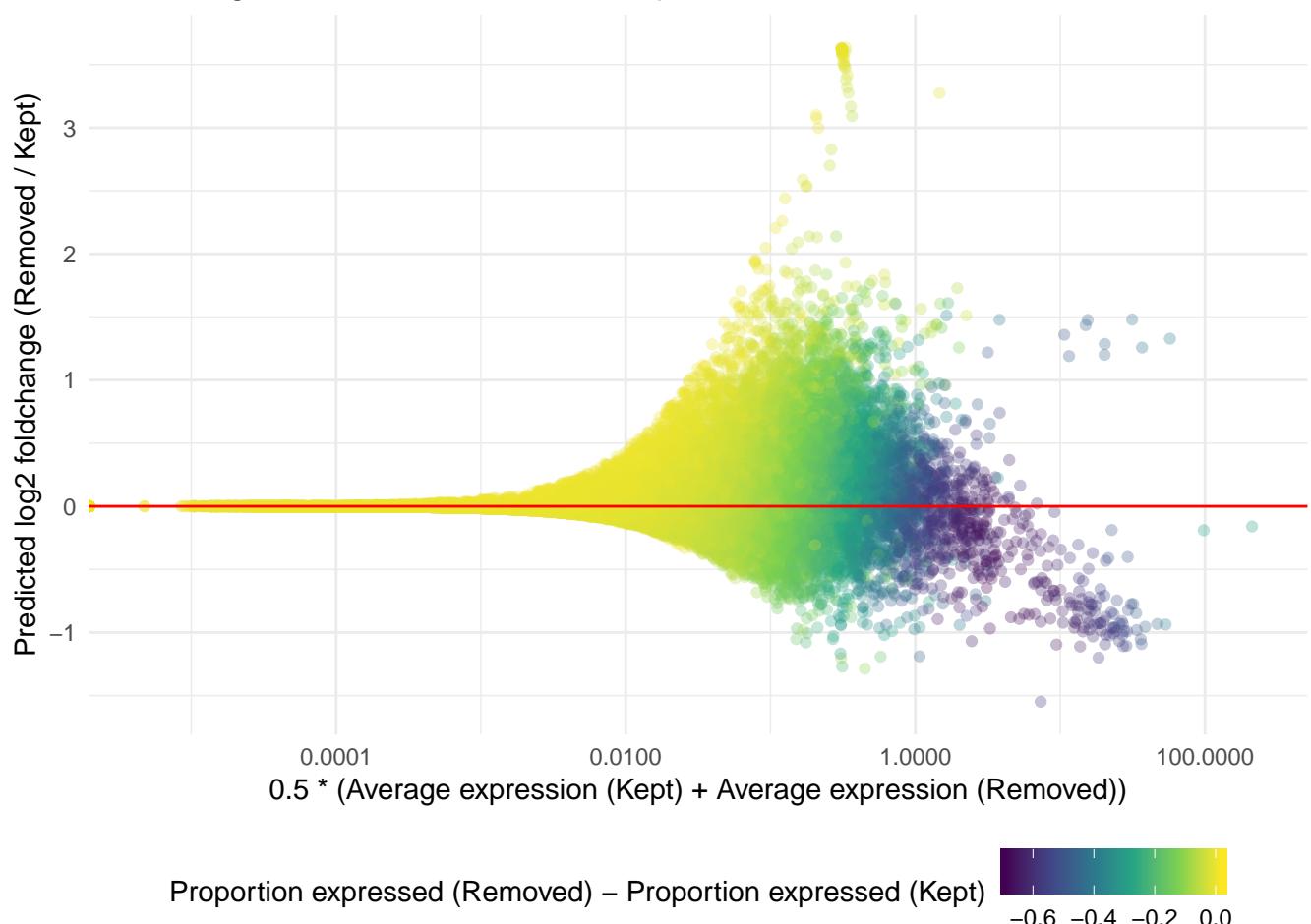
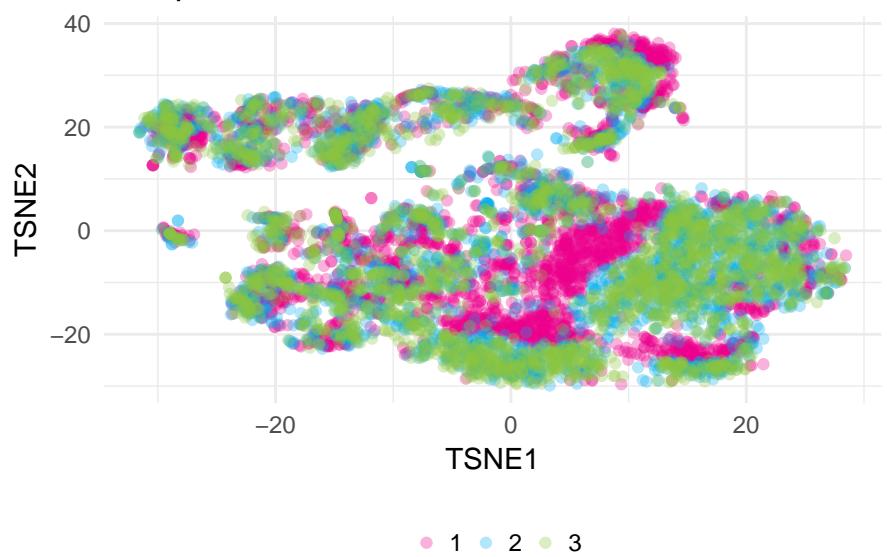
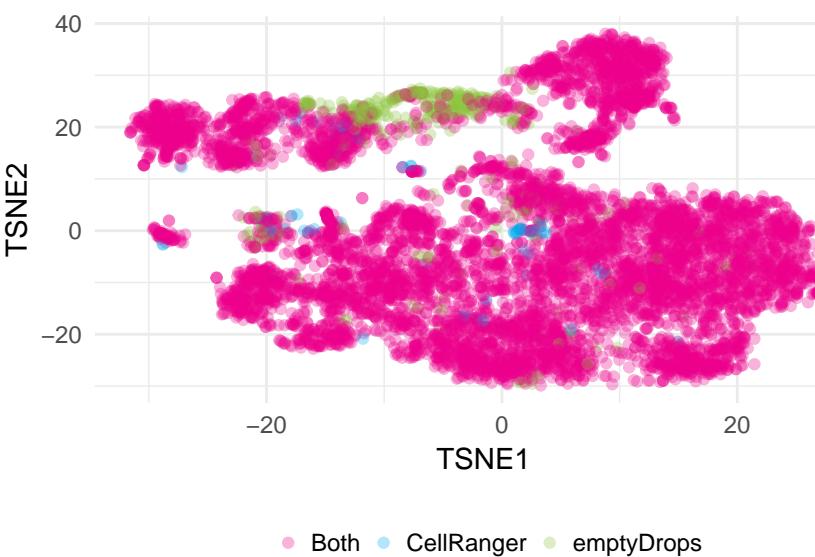


A Foldchange between removed and kept cells**B**

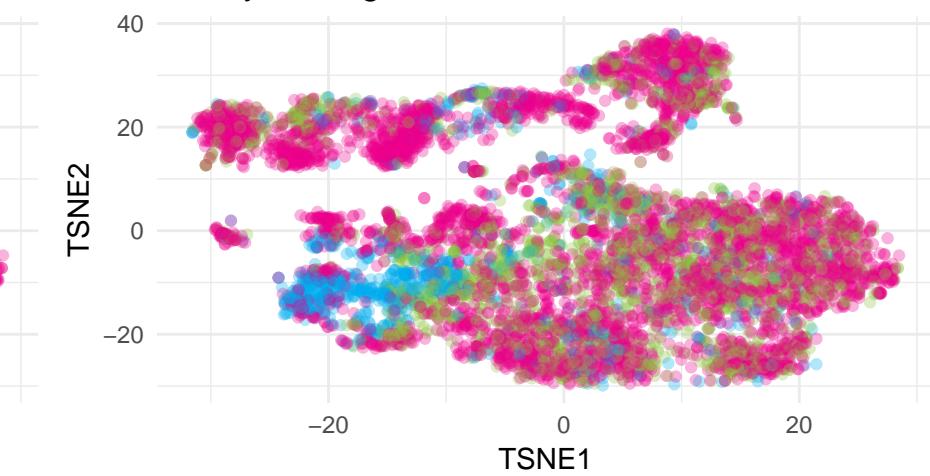
Gene	Predicted foldchange (log2)
AC005523.2	3.635019
OCLM	3.634619
LINC00843	3.627595
AL627309.1	3.626357
LINC02067	3.626263
AC025569.1	3.625177
ITPKA	3.624939
GCNT4	3.618081
AC010761.1	3.617089
LINC00921	3.616814
AC008738.2	3.611973
HSF4	3.610516
NR5A2	3.602700
AC091057.3	3.602354
AP000350.7	3.599396

C Sample

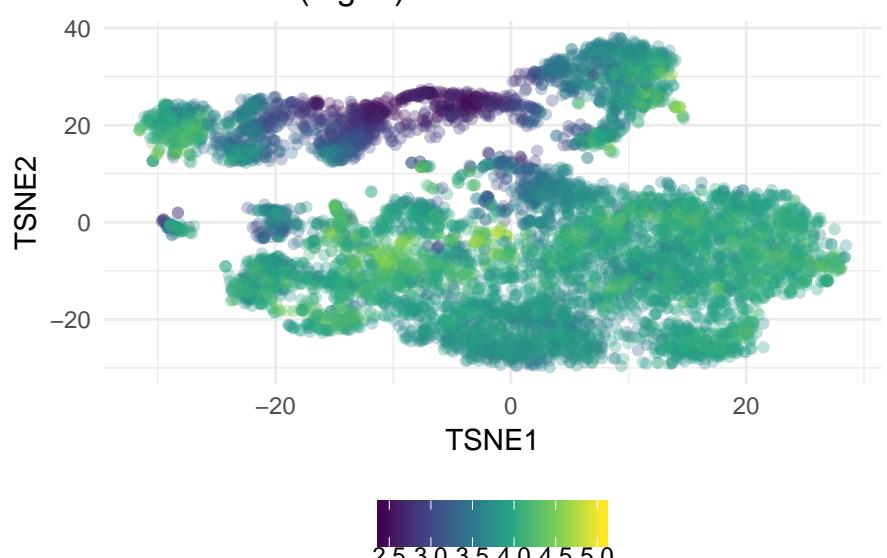
Selection method



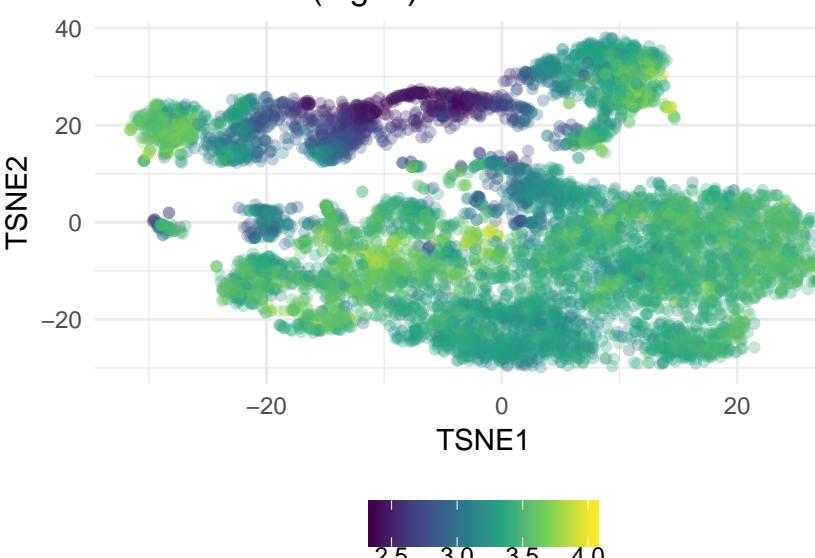
Cell cycle stage



Total counts (log10)



Total features (log10)



% Mitochondrial

