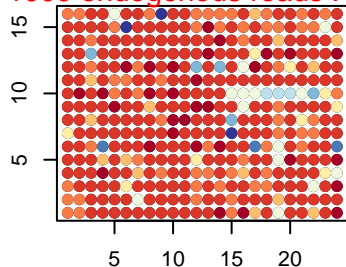
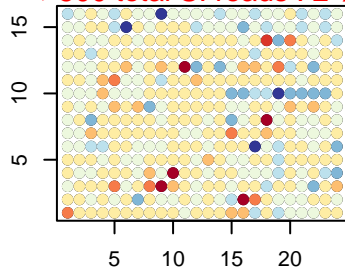


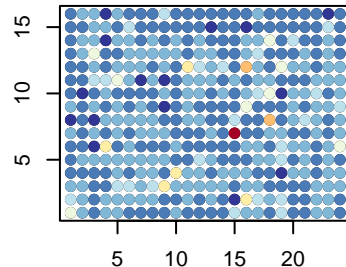
Total endogenous reads
 >1000 endogenous reads : 95 %



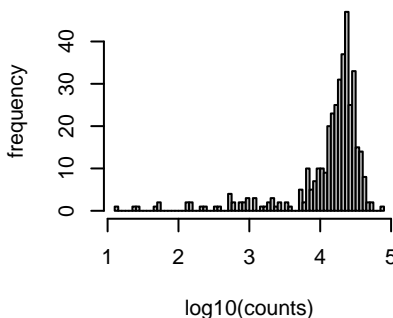
Total spike in reads
 >500 total SI reads : 2 %



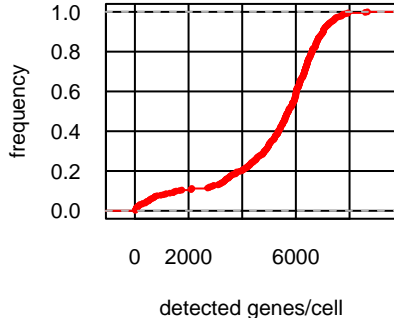
ratio SI/Endogenous reads



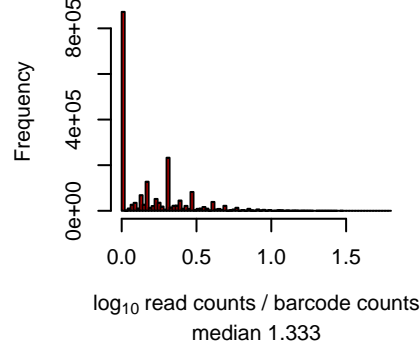
total unique reads w/o SI
 mean: 19157 median: 19565



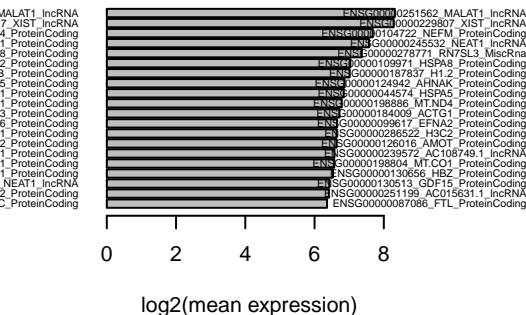
cumulative dist genes
 mean: 5166 median: 5728



oversequencing_molecules



top expressed genes



top noisy genes

