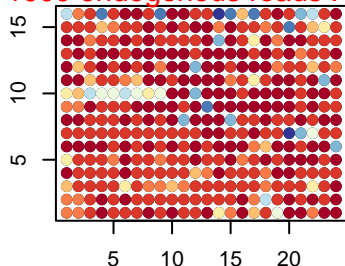
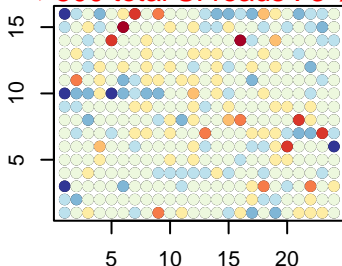


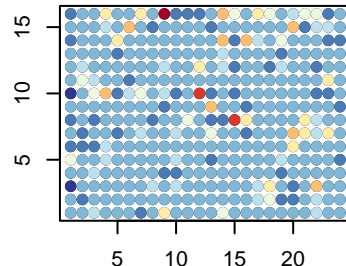
Total endogenous reads
 >1000 endogenous reads : 93 %



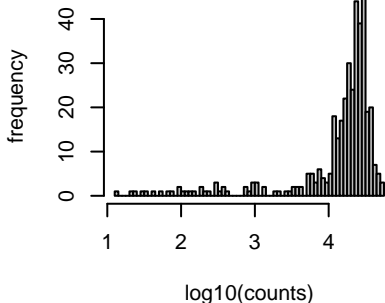
Total spike in reads
 >500 total SI reads : 3 %



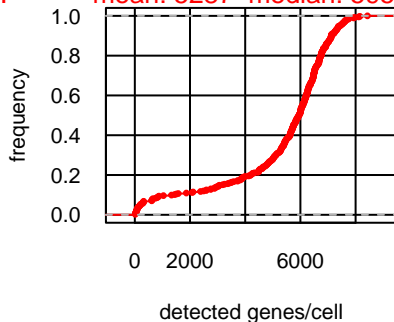
ratio SI/Endogenous reads



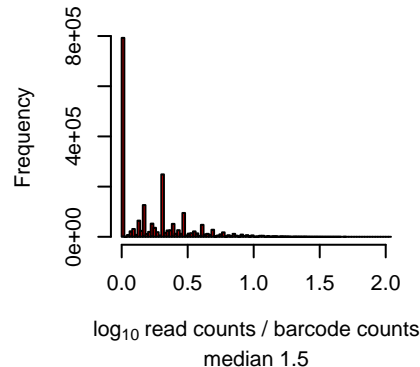
total unique reads w/o SI
 mean: 20490 median: 21091



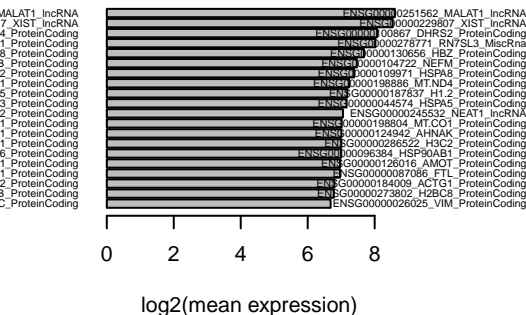
cumulative dist genes
 mean: 5287 median: 5952



oversequencing_molecules



top expressed genes



top noisy genes

