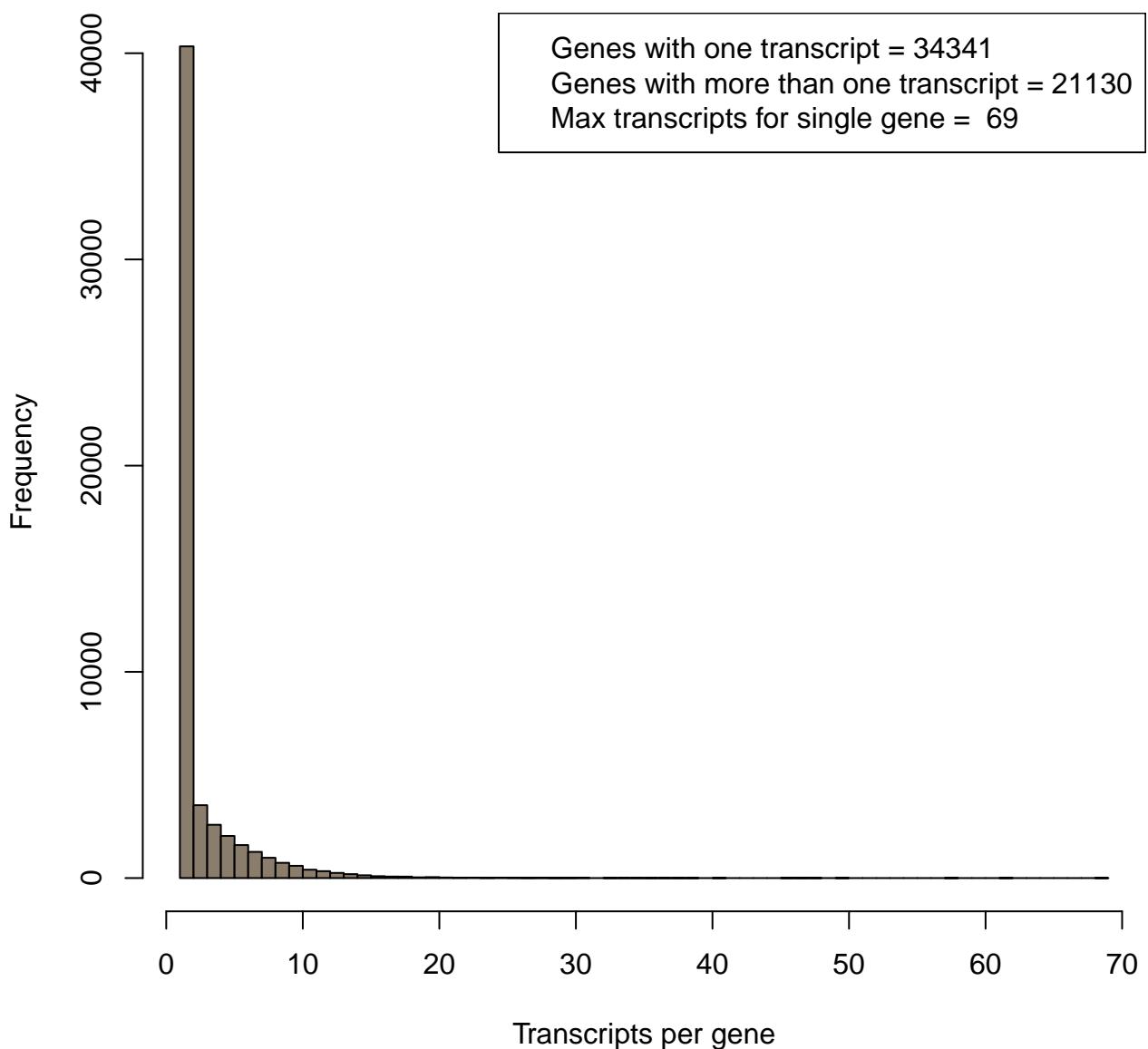
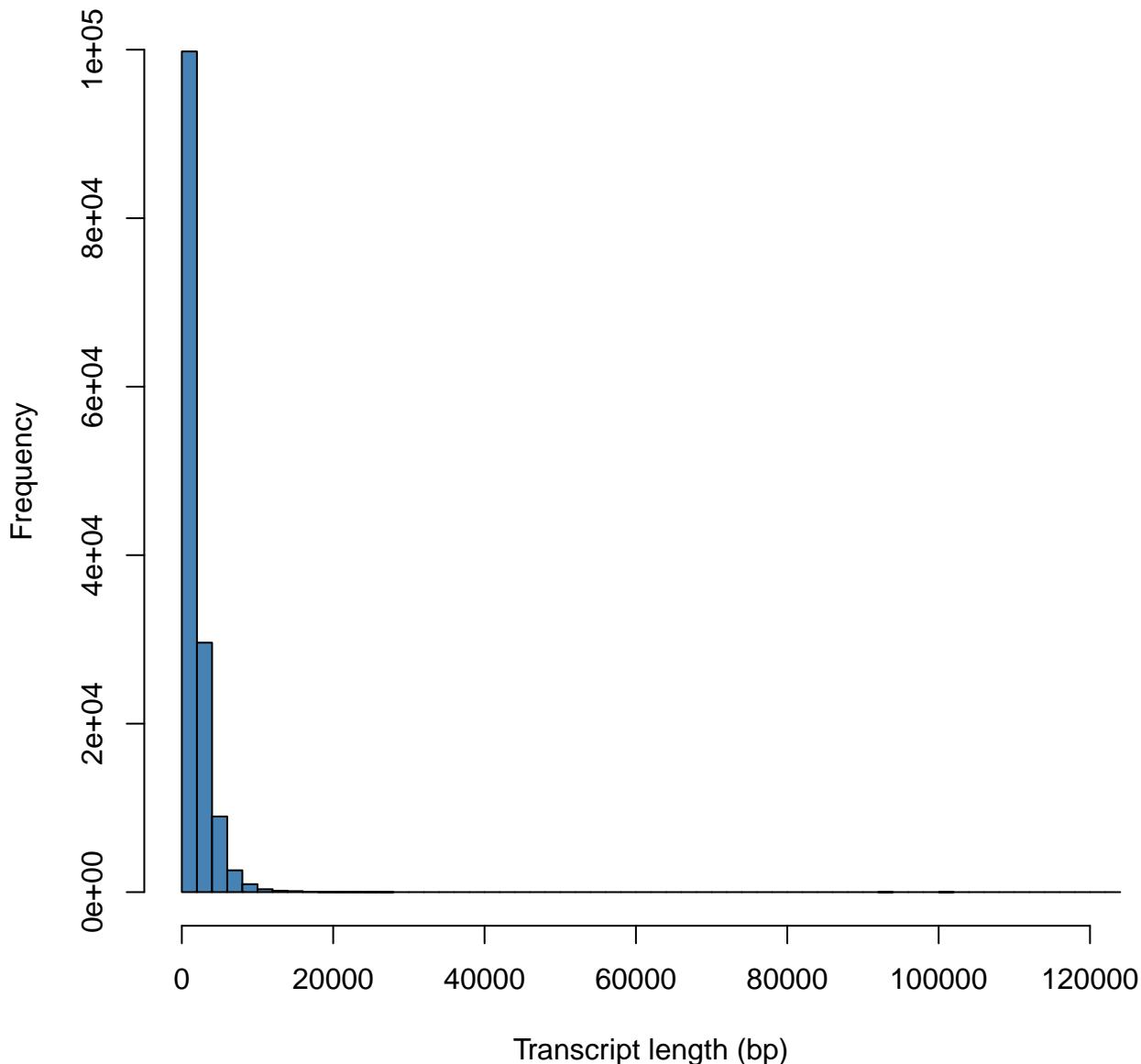


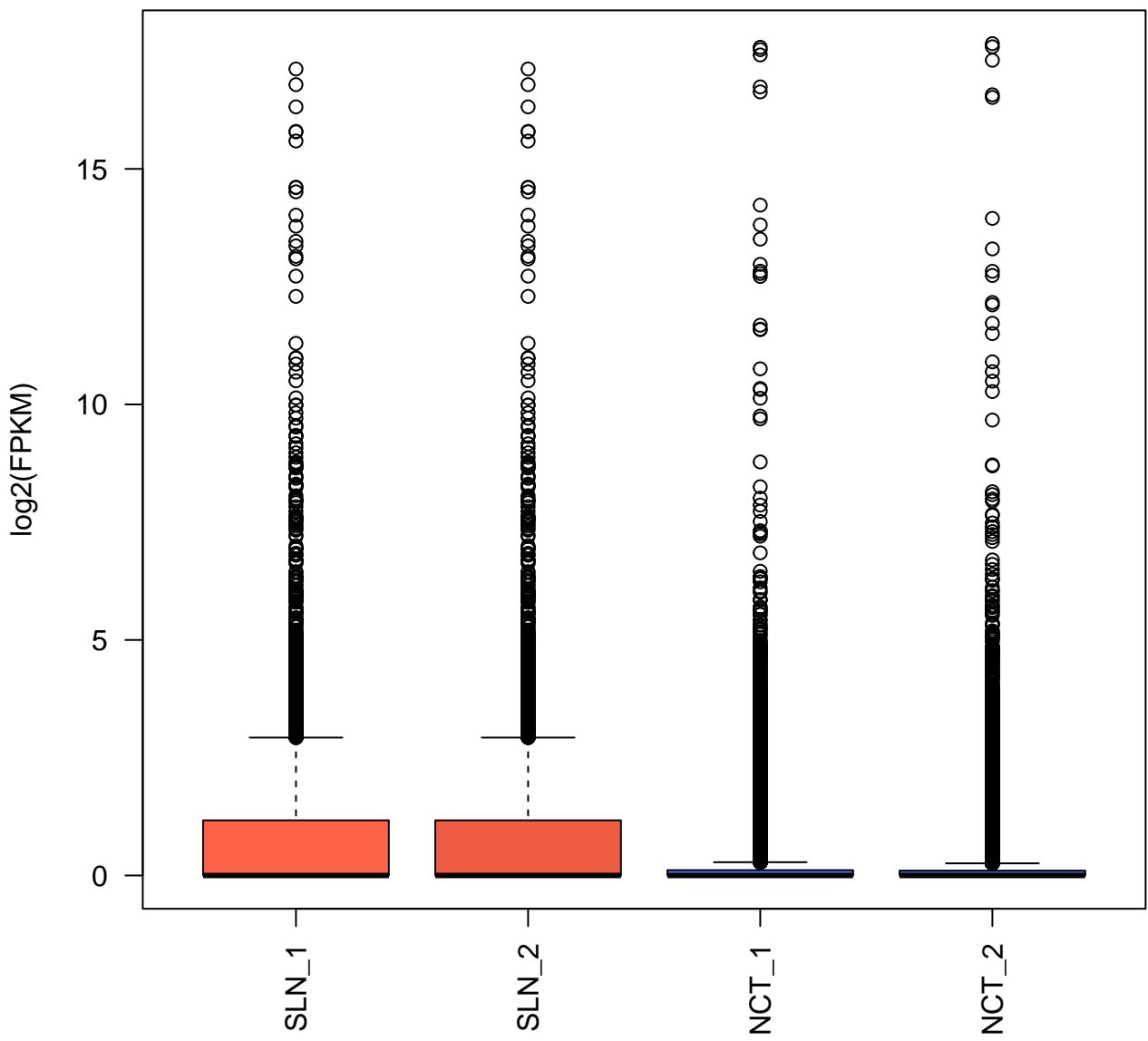
## Distribution of transcript count per gene



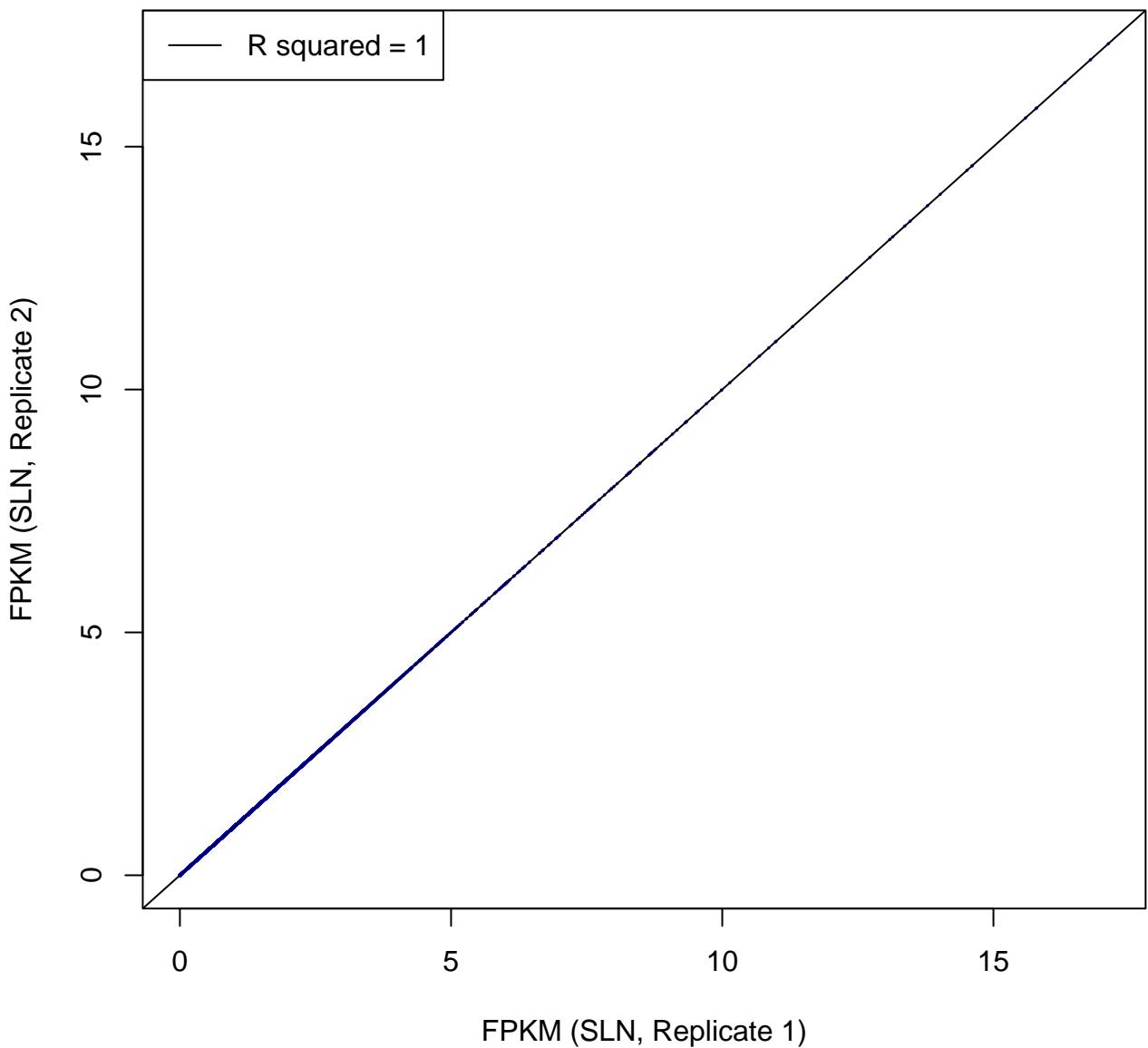
## Distribution of transcript lengths



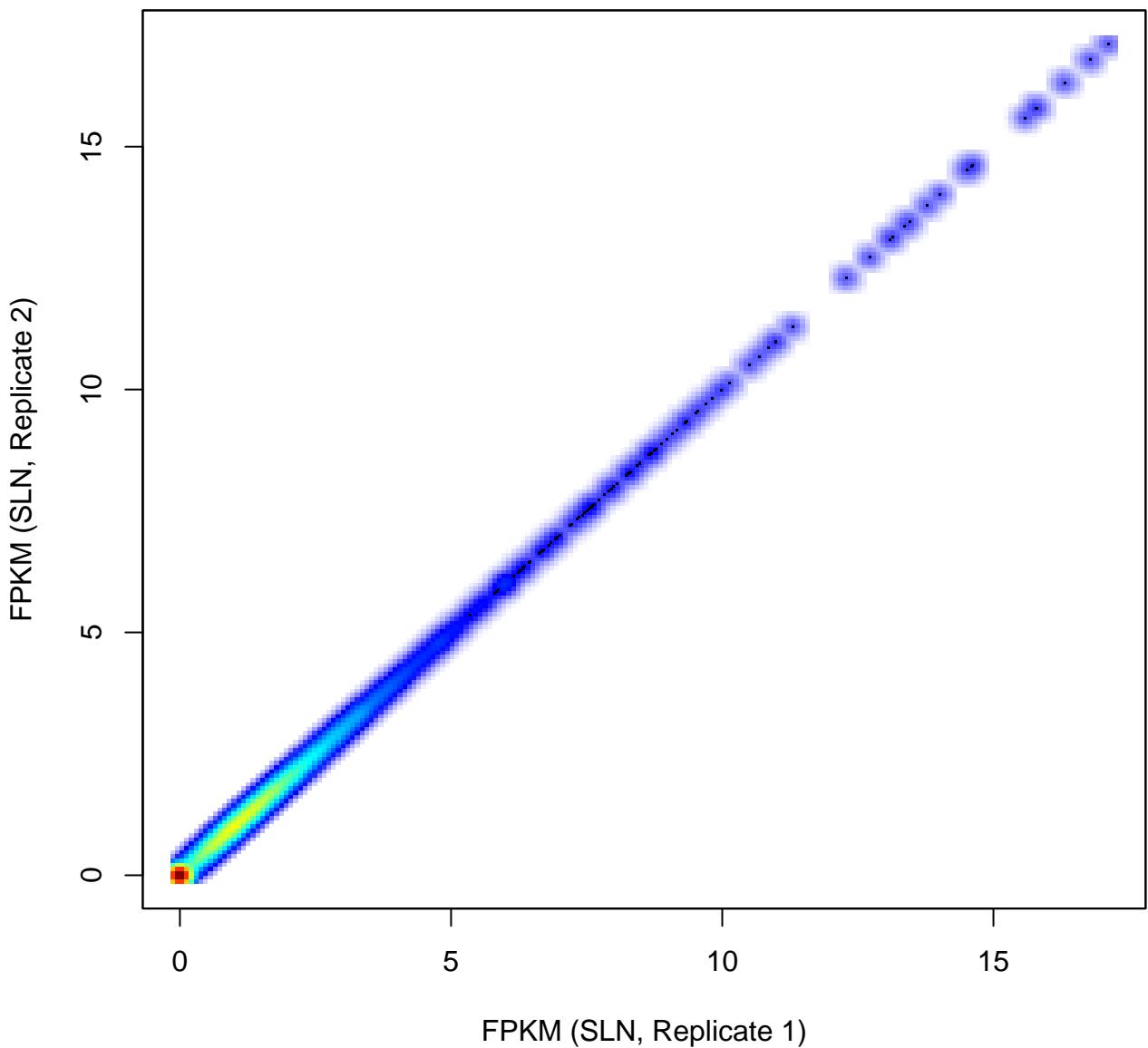
## Distribution of FPKMs for all 4 libraries



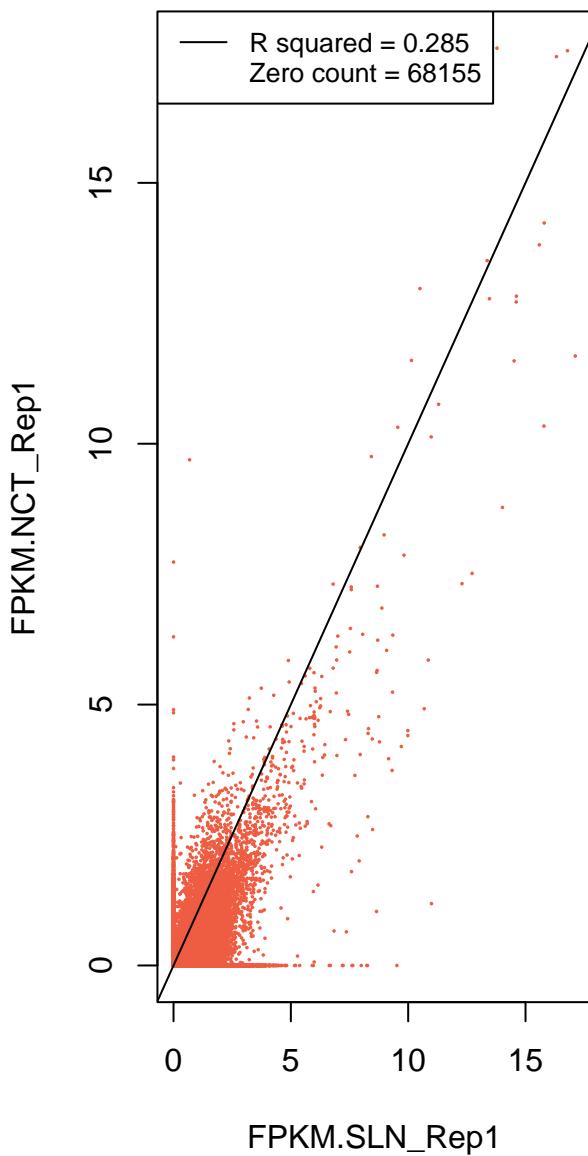
## Comparison of expression values for a pair of replicates



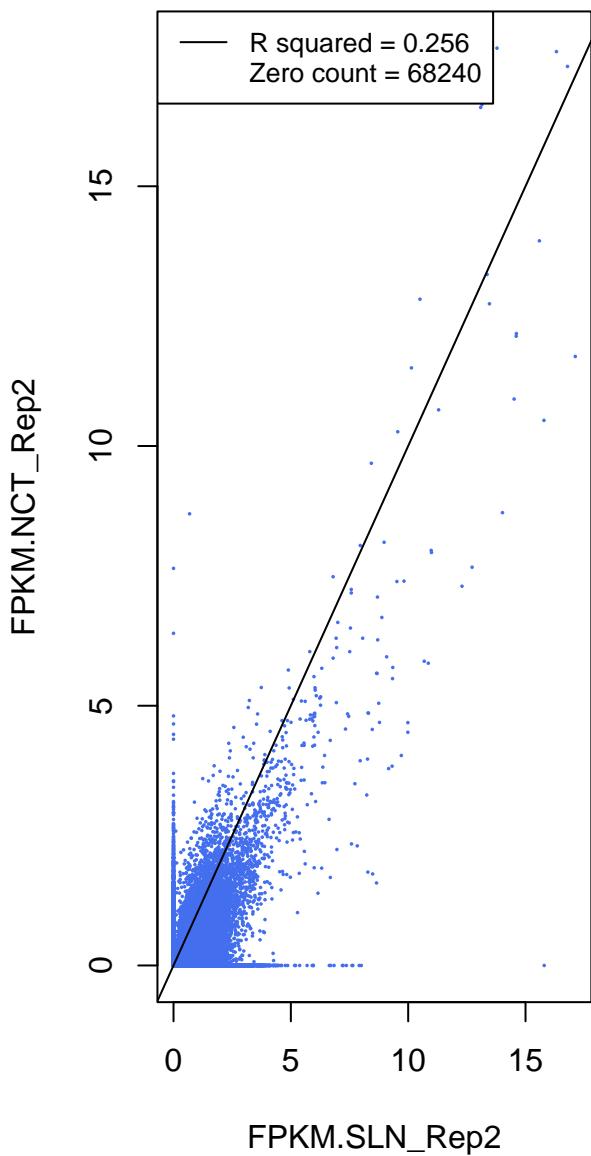
## Comparison of expression values for a pair of replicates



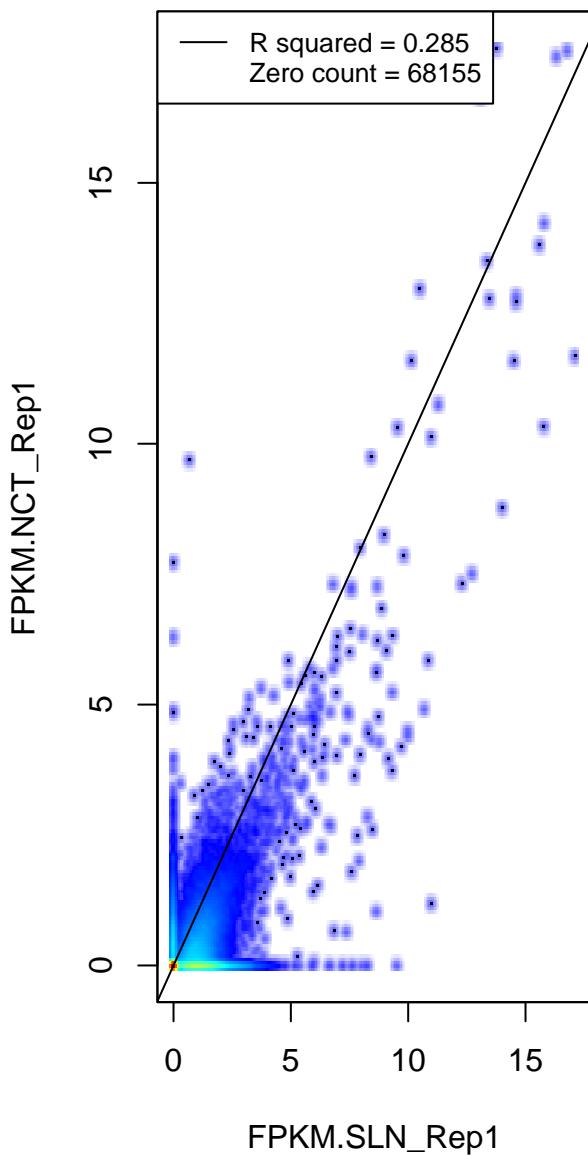
**SLN\_1 vs NCT\_1**



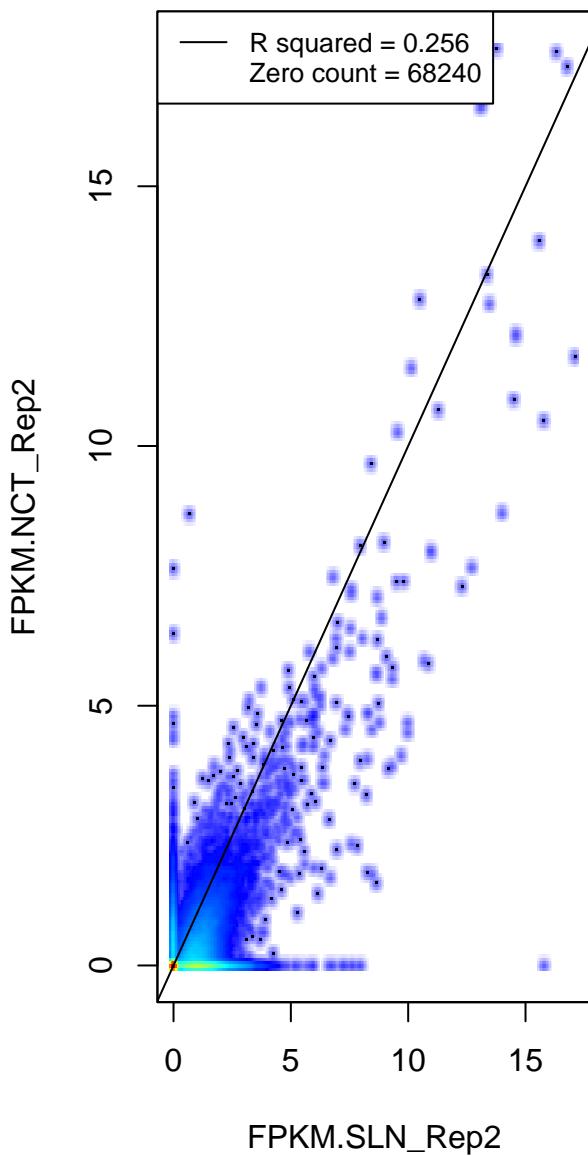
**SLN\_2 vs NCT\_2**



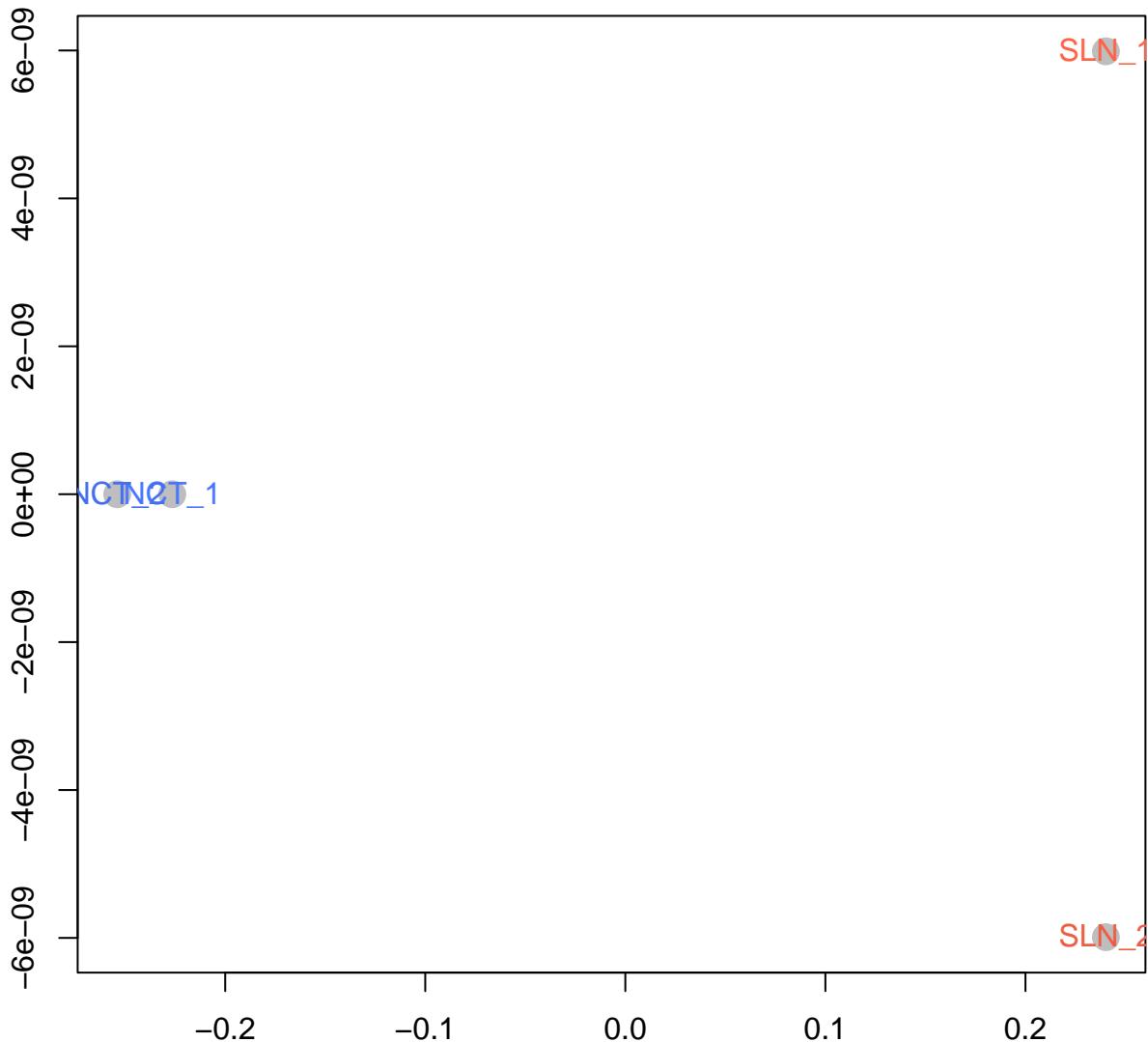
**SLN\_1 vs NCT\_1**



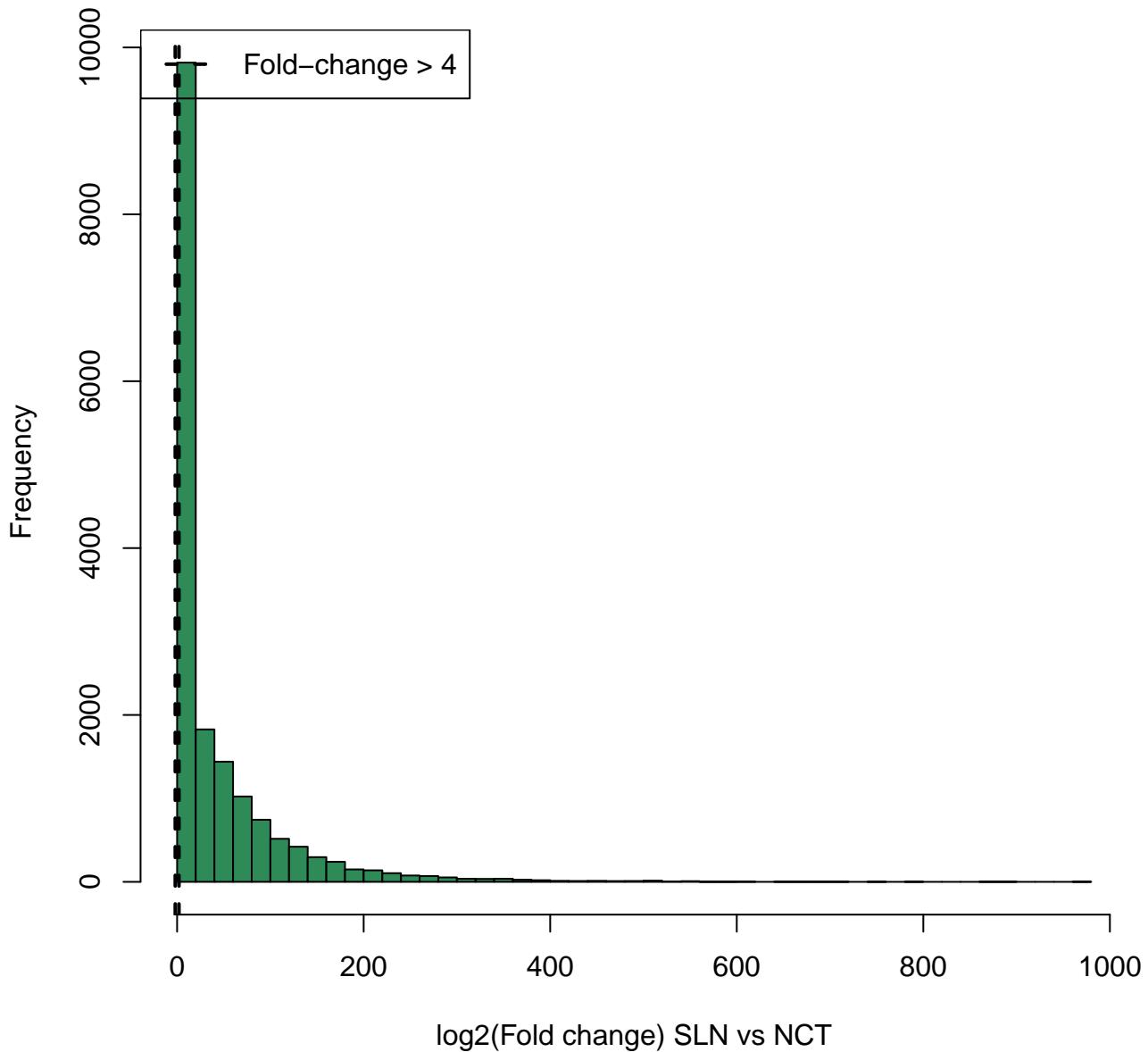
**SLN\_2 vs NCT\_2**



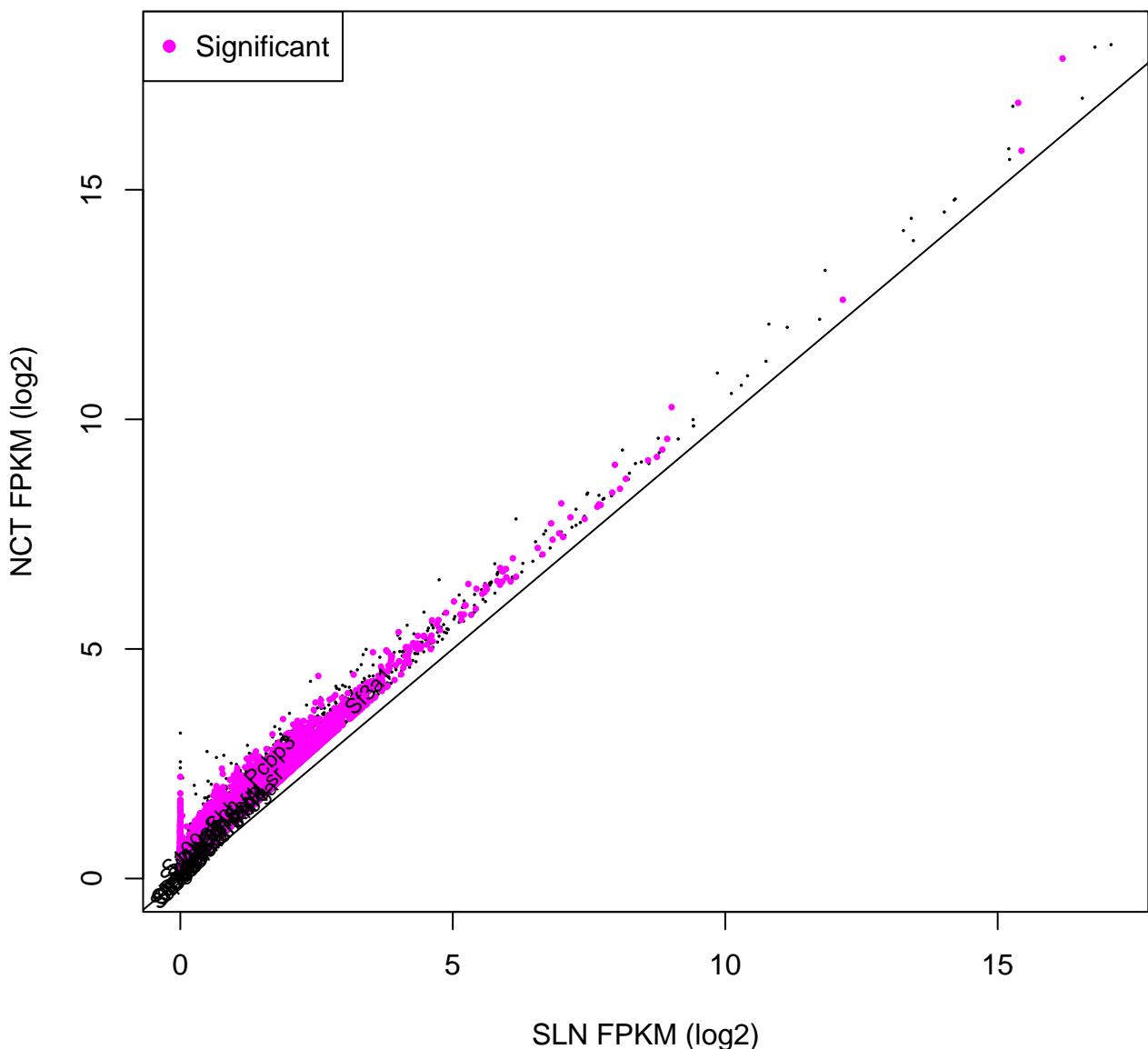
## MDS distance plot (all non-zero genes)



## Distribution of differential expression values



## SLN vs NCT FPKMs



Color Key

## sig DE Transcripts

