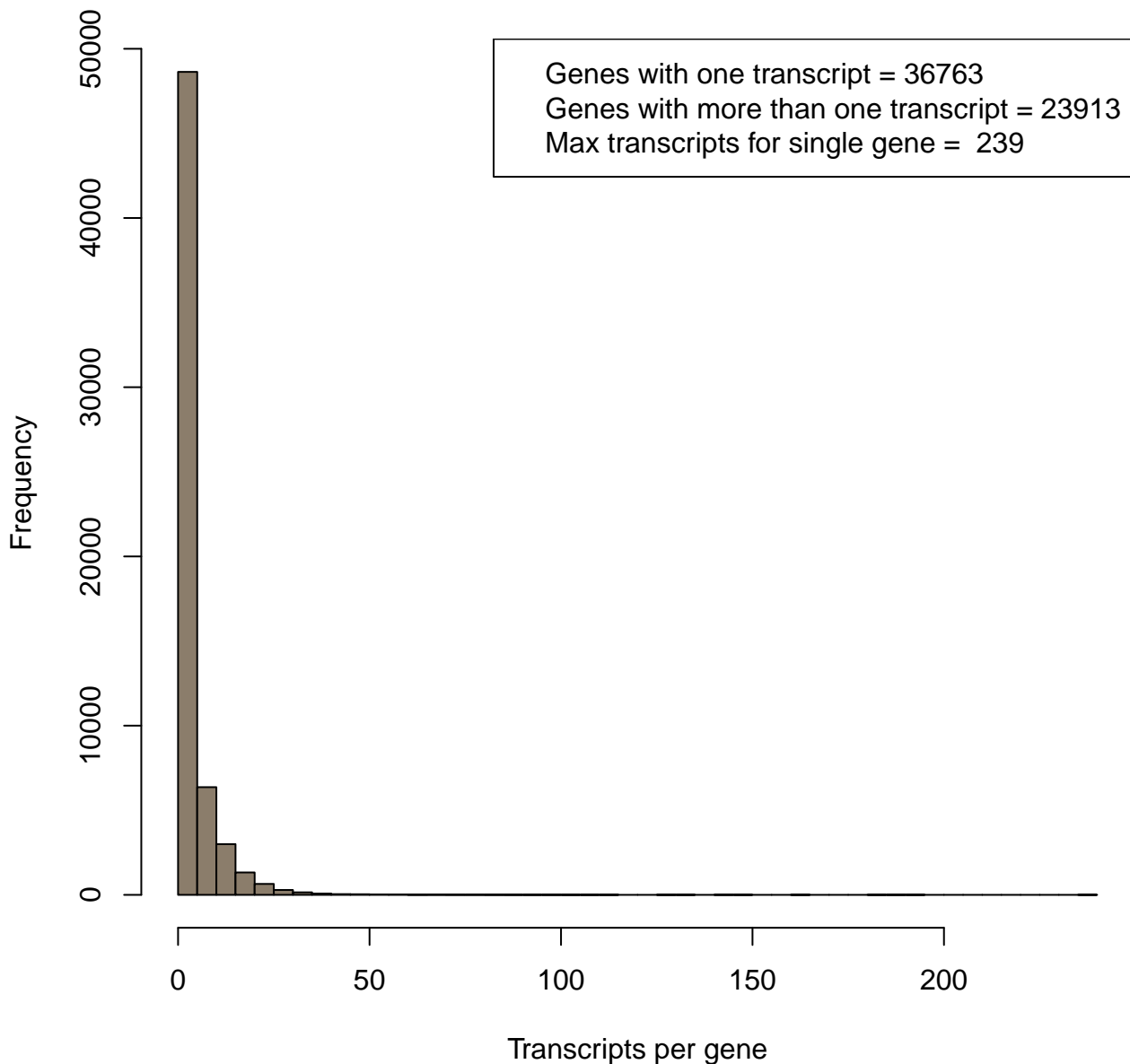
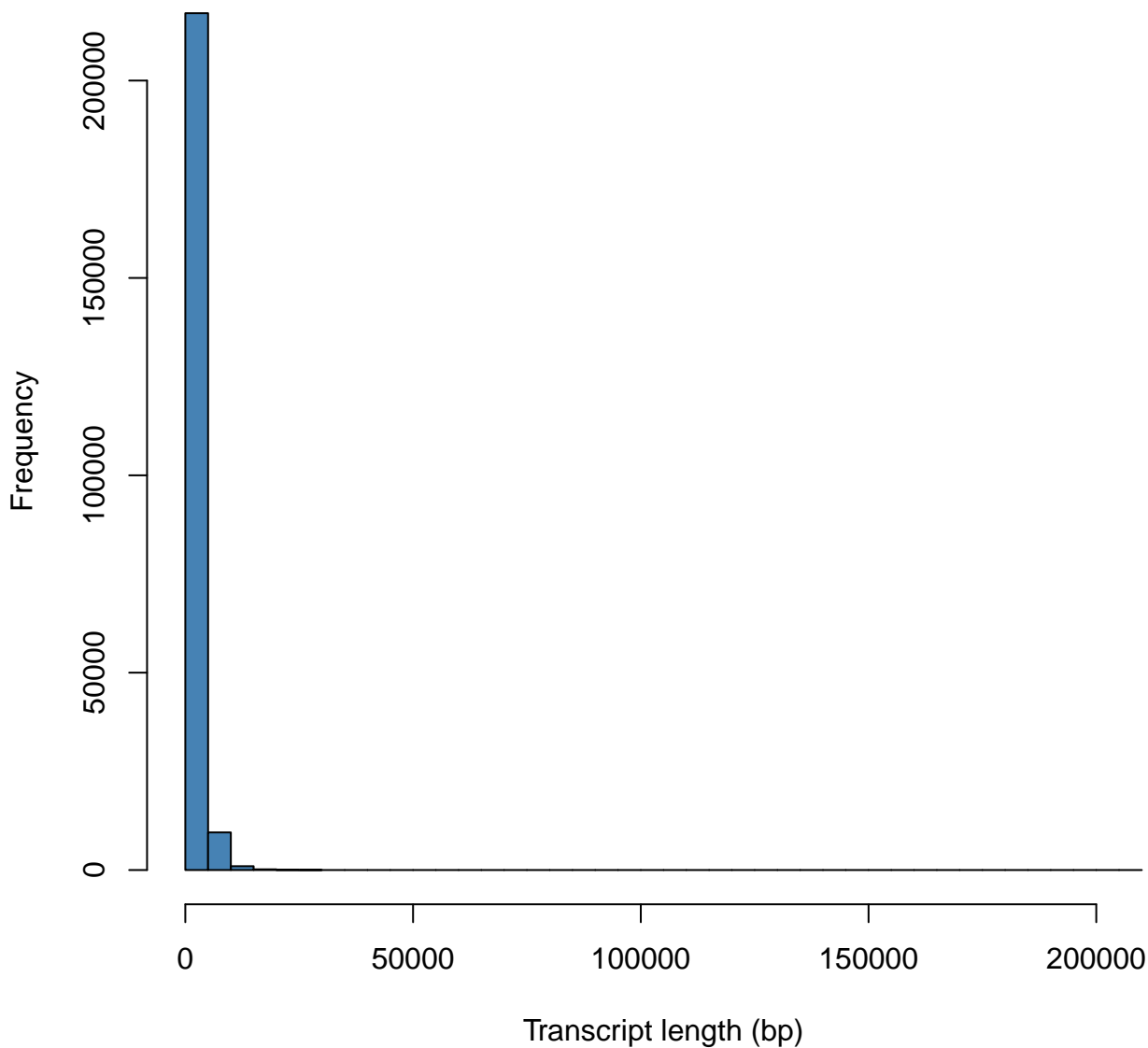


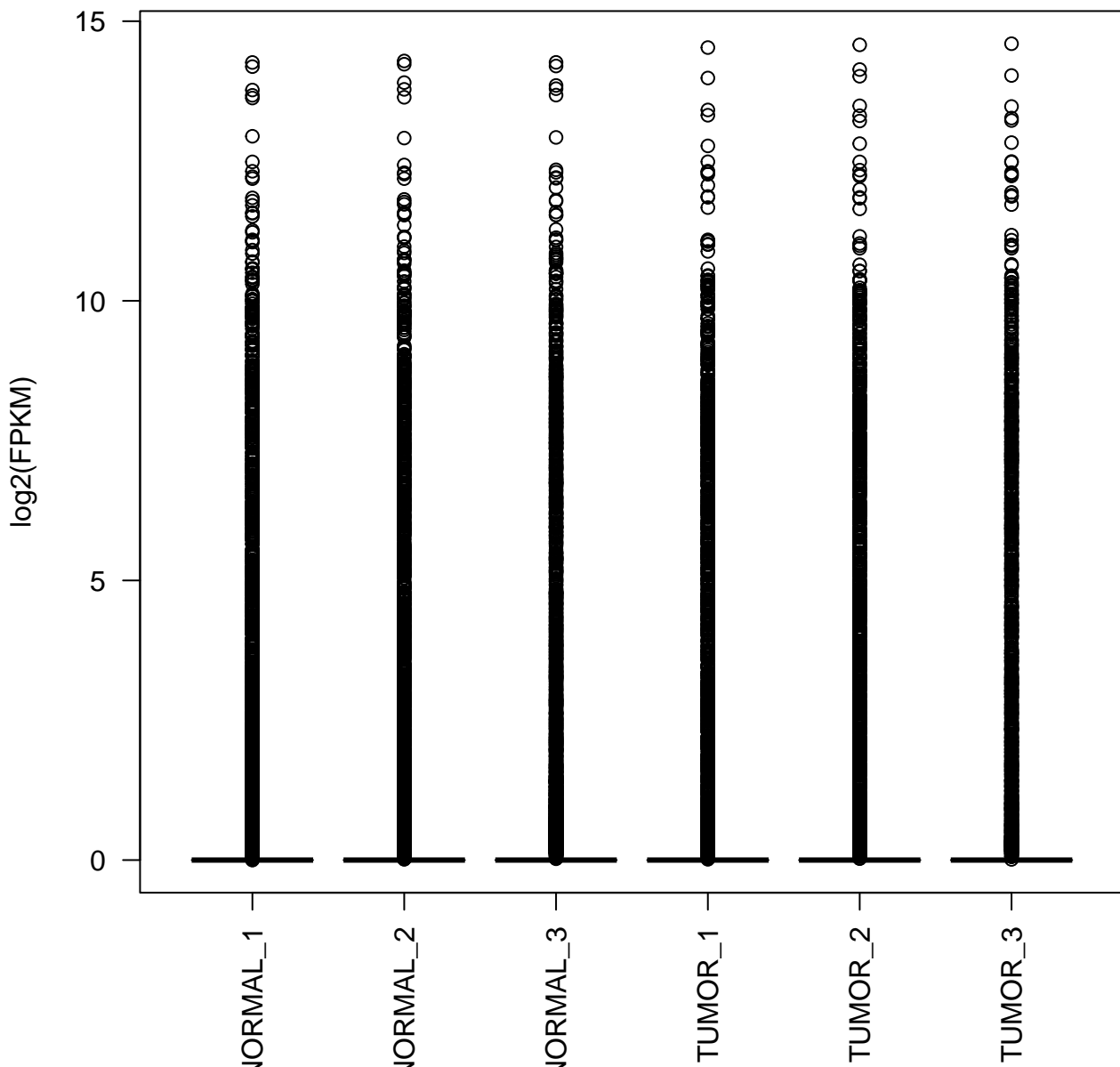
Distribution of transcript count per gene



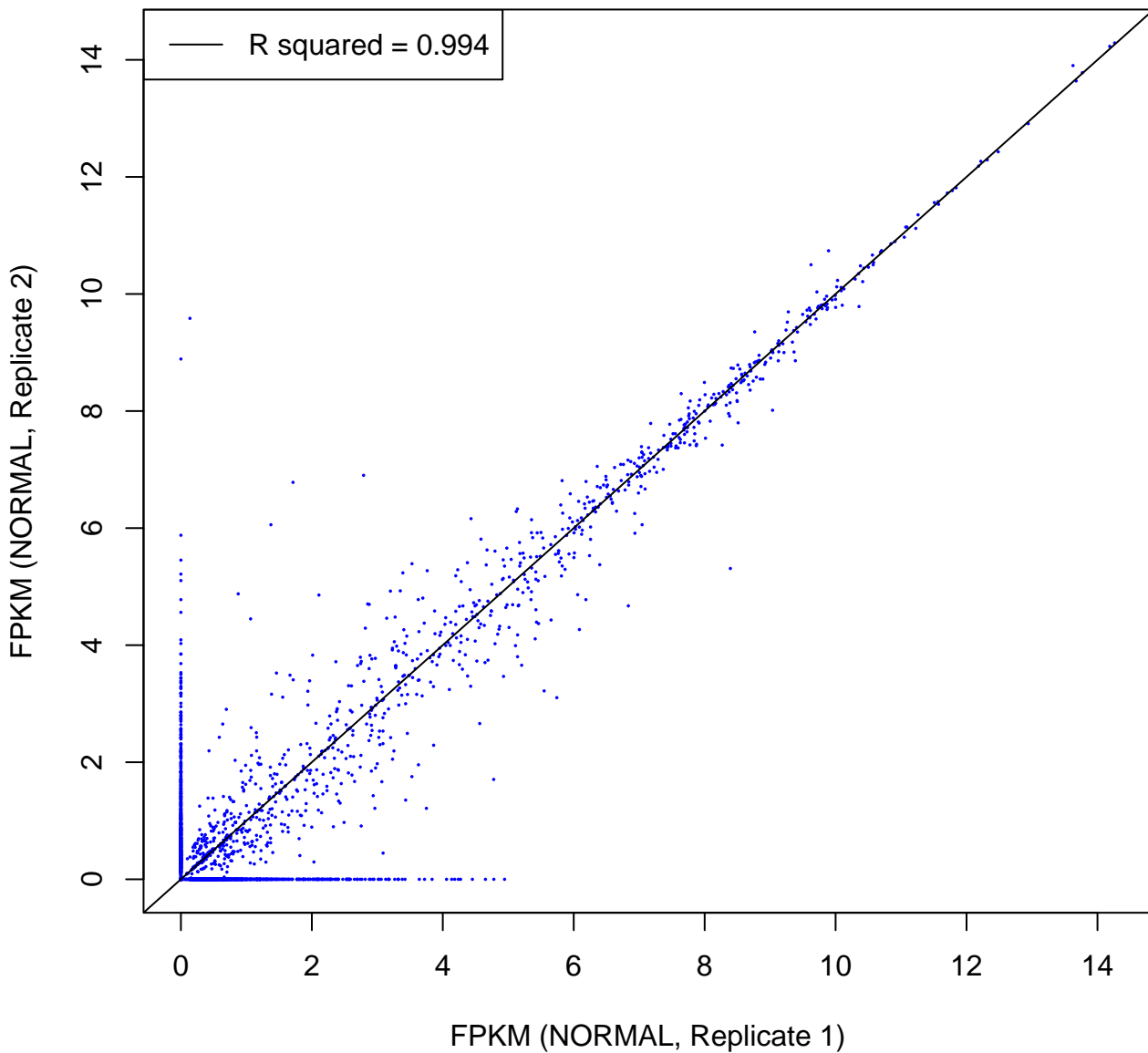
Distribution of transcript lengths



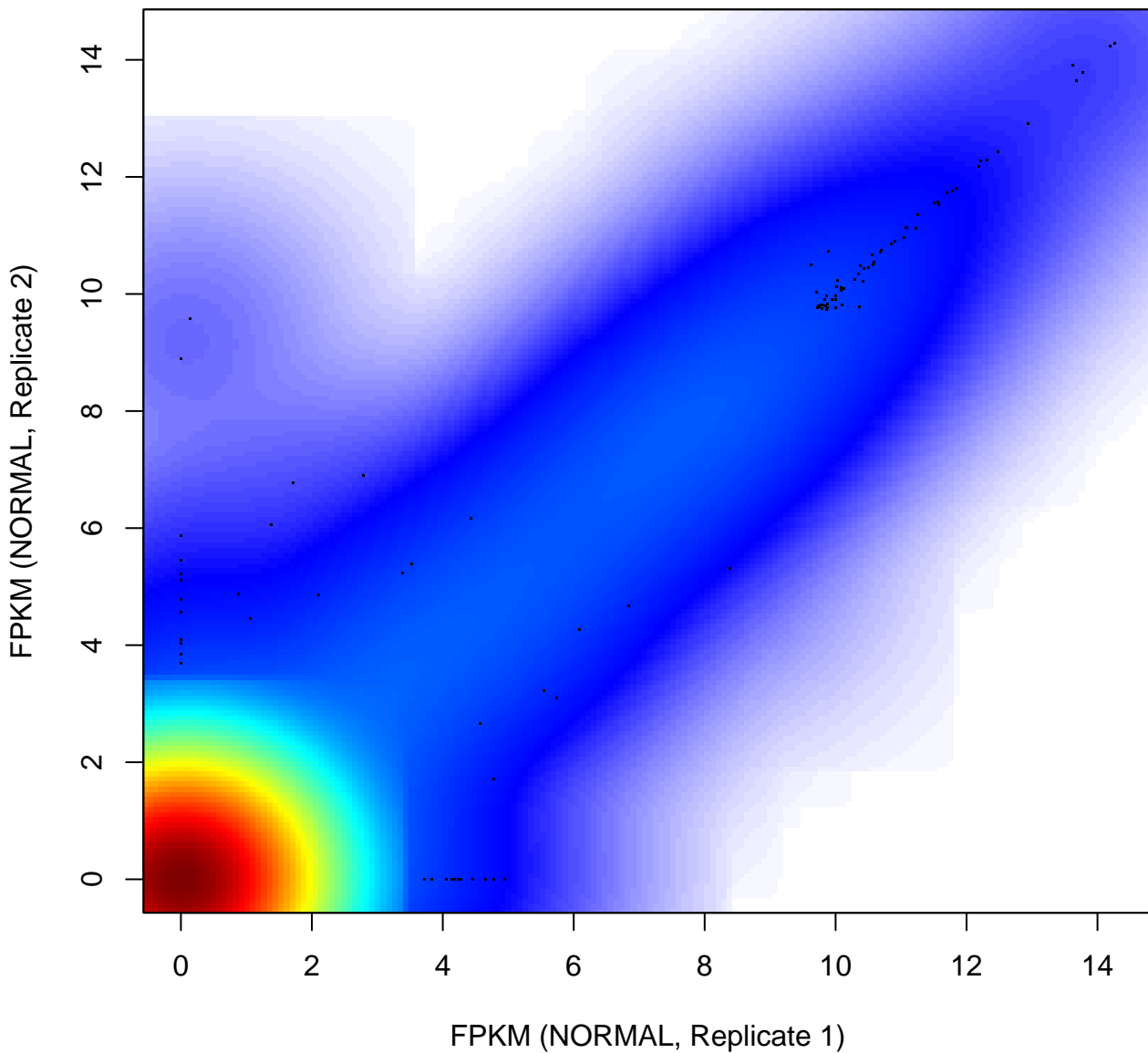
Distribution of FPKMs for all 6 libraries



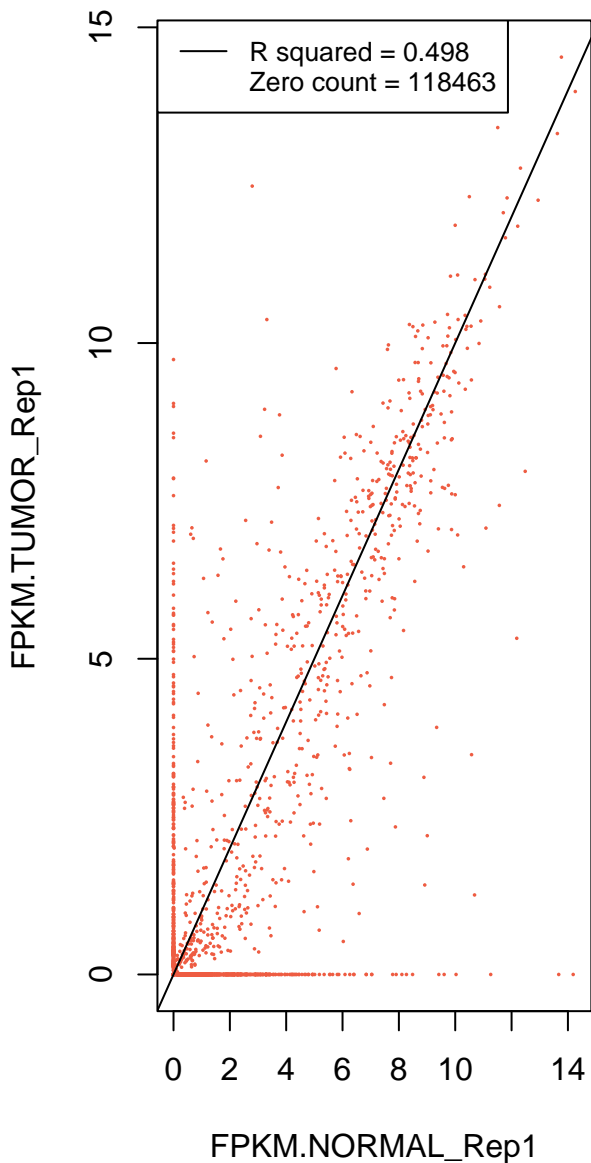
Comparison of expression values for a pair of replicates



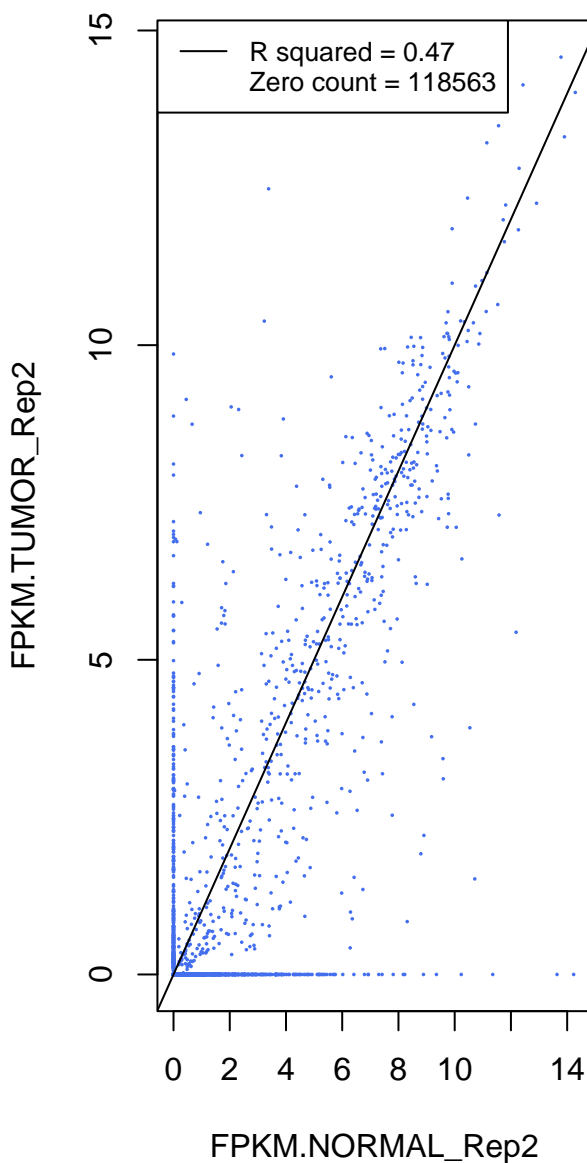
Comparison of expression values for a pair of replicates



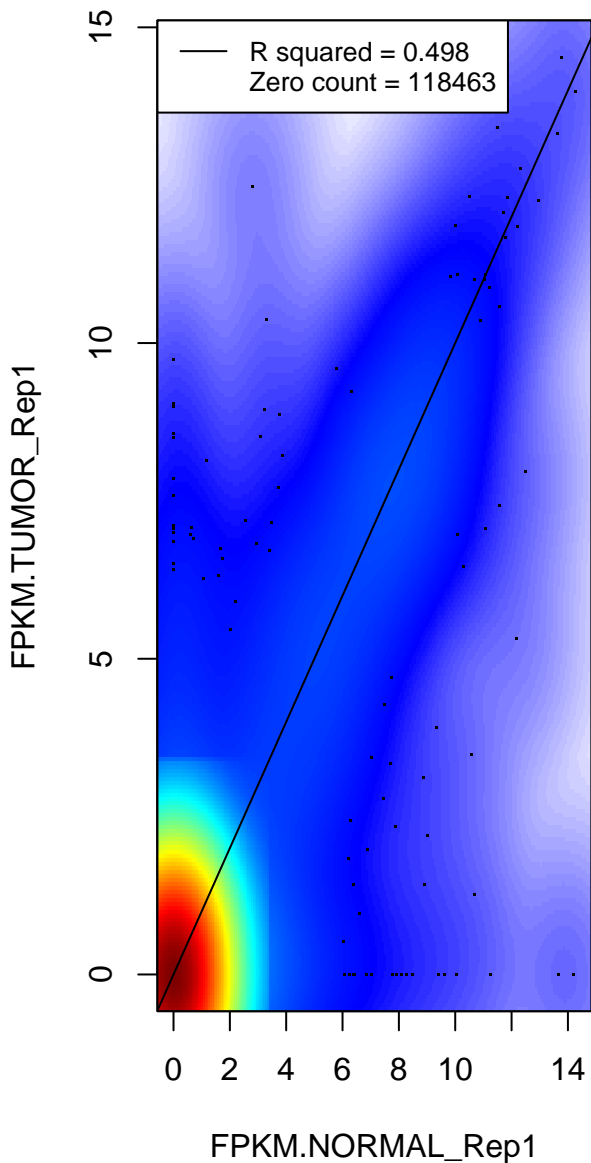
NORMAL_1 vs TUMOR_1



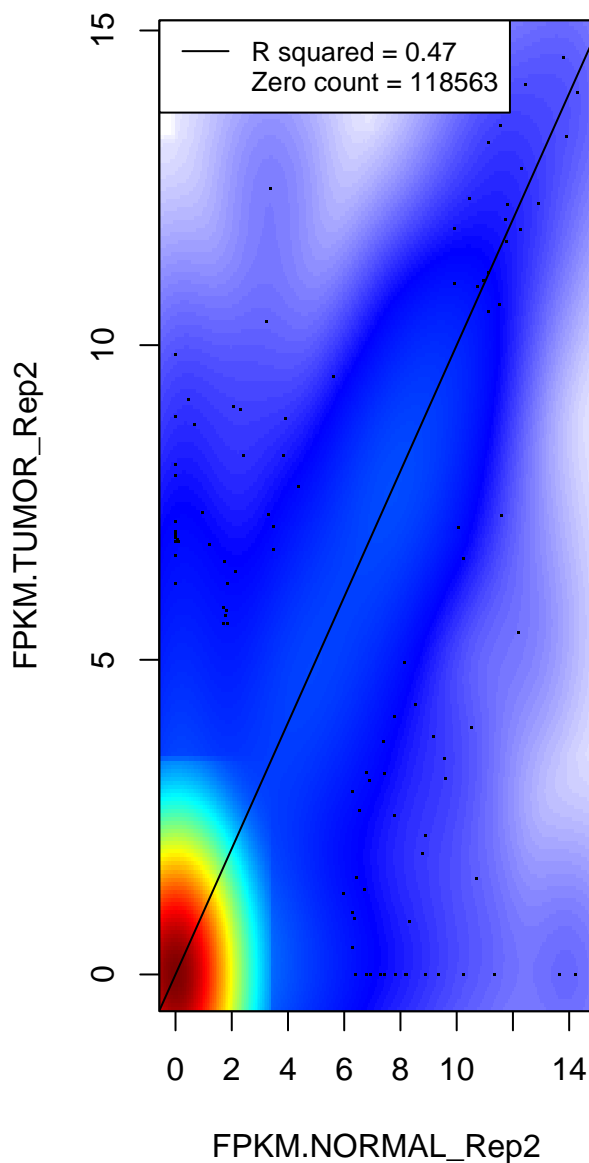
NORMAL_2 vs TUMOR_2



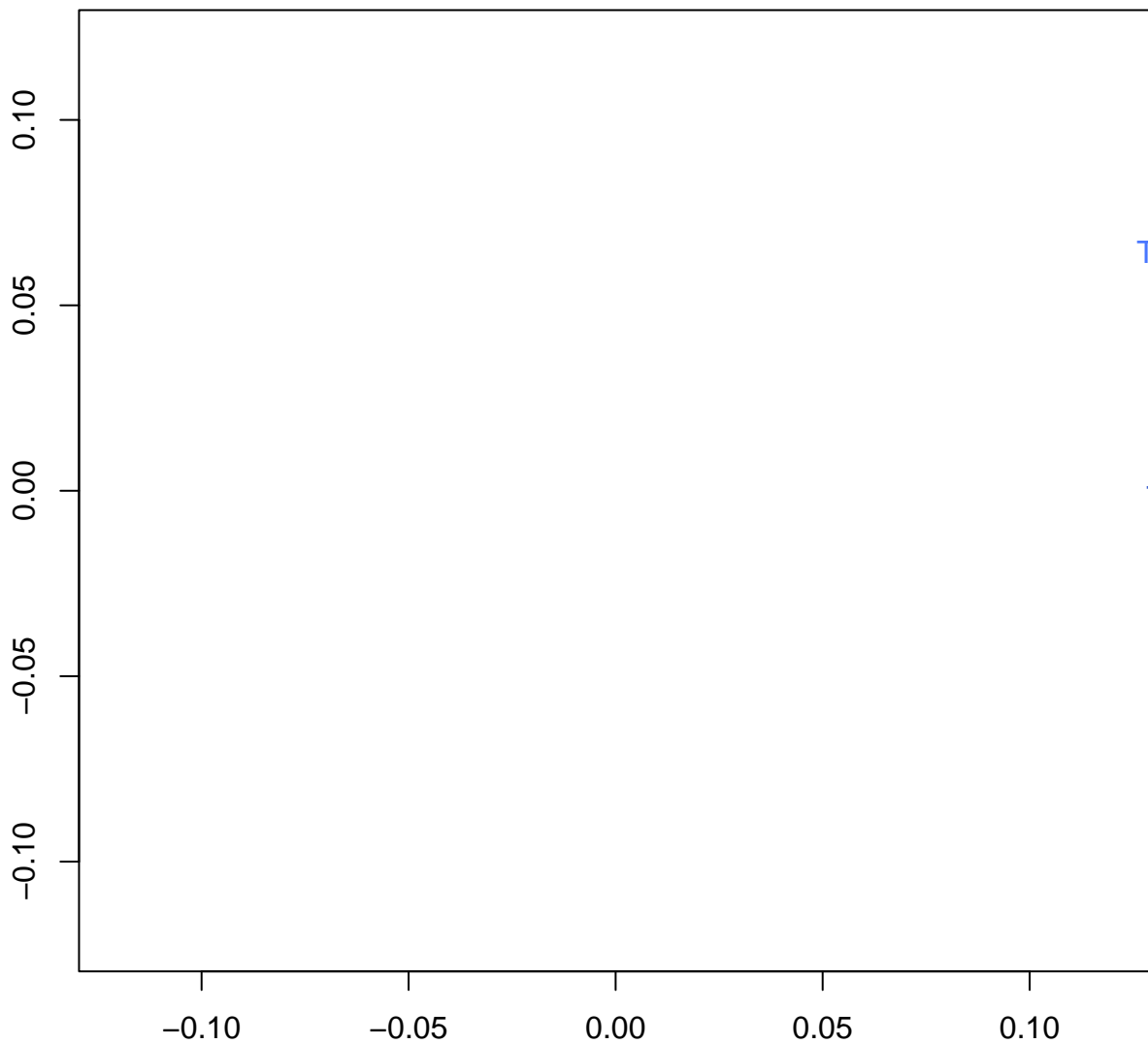
NORMAL_1 vs TUMOR_1



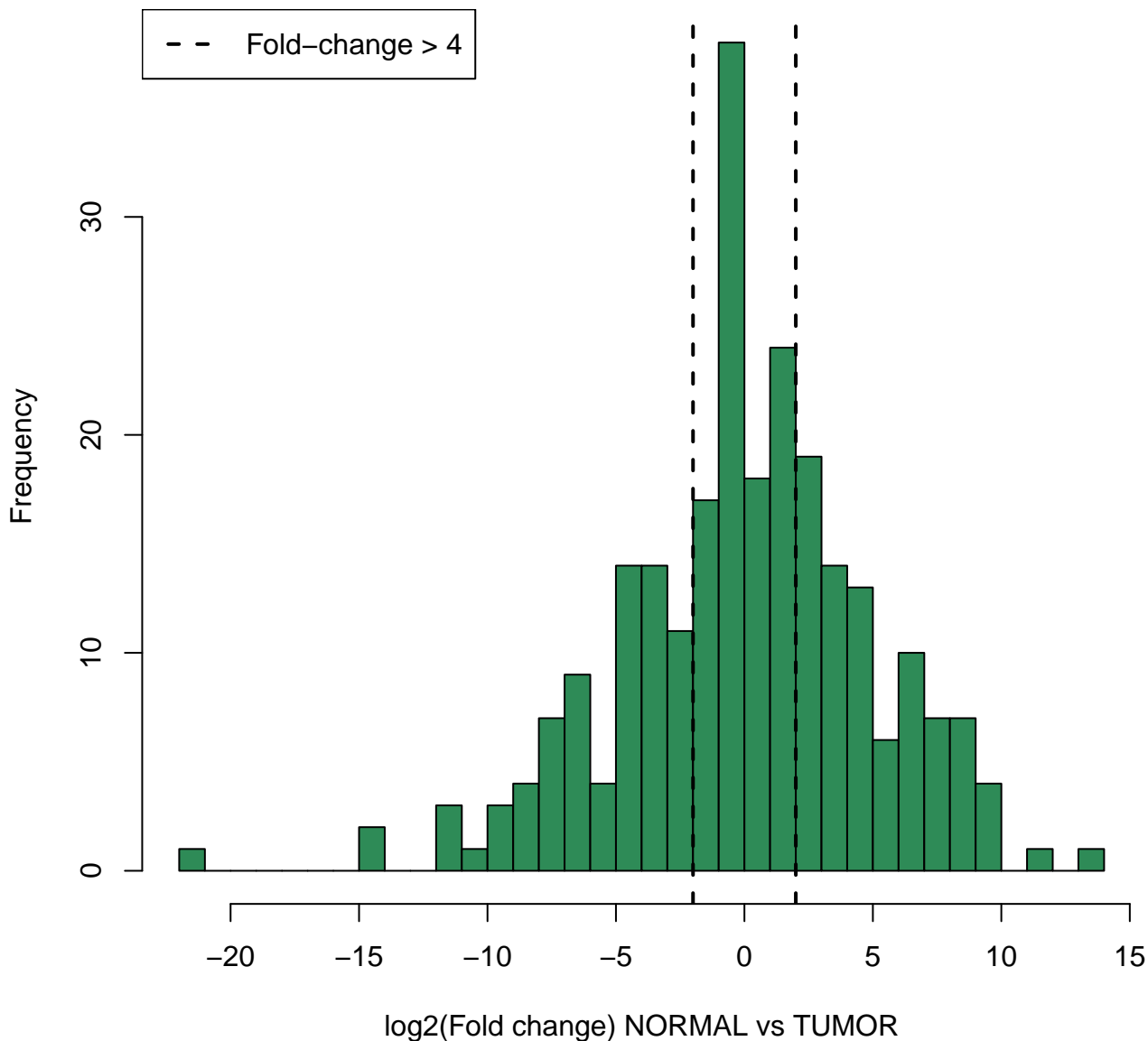
NORMAL_2 vs TUMOR_2



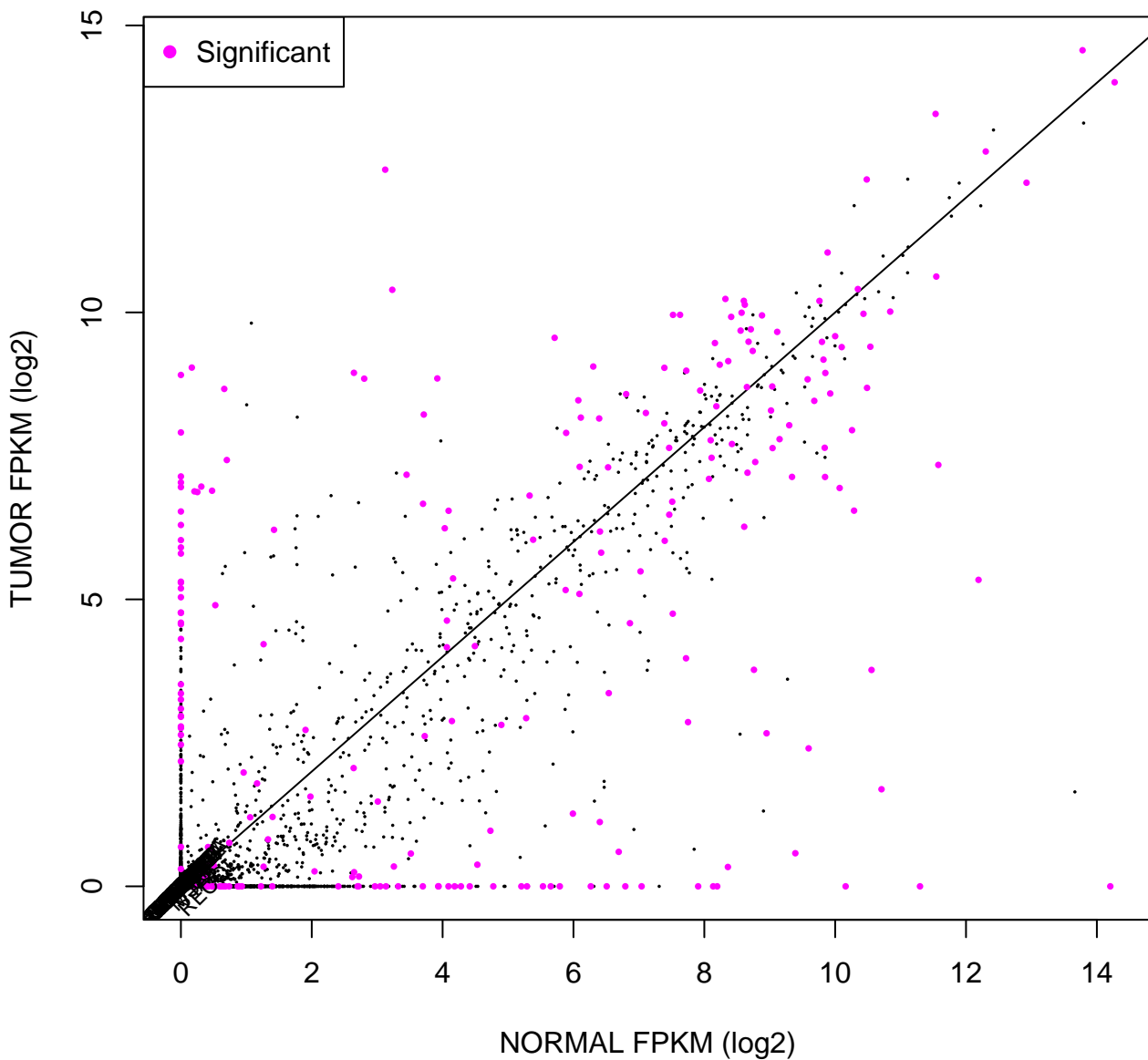
MDS distance plot (all non-zero genes)



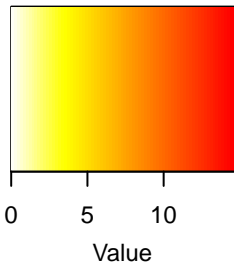
Distribution of differential expression values



NORMAL vs TUMOR FPKMs



Color Key



sig DE Transcripts

