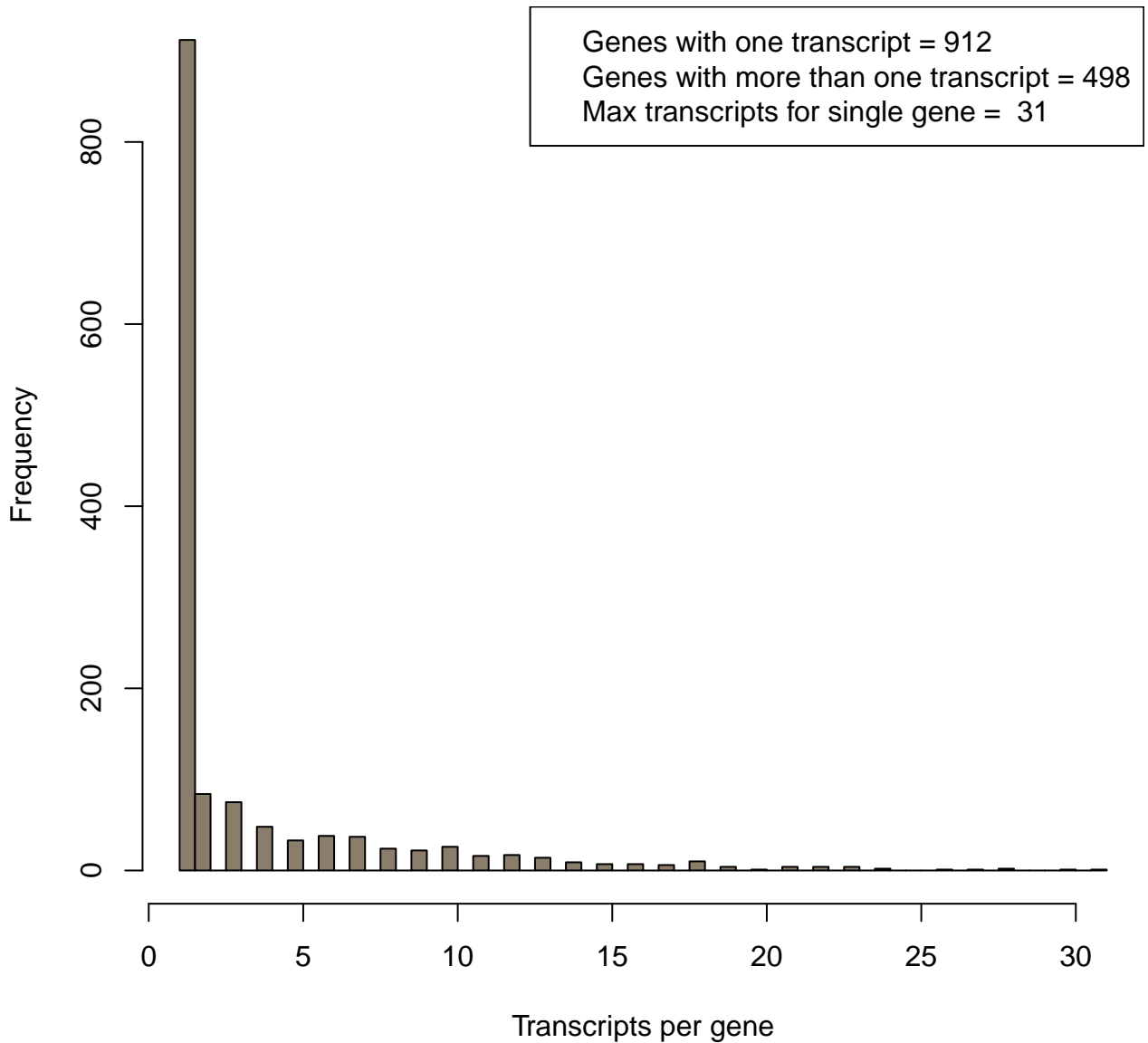
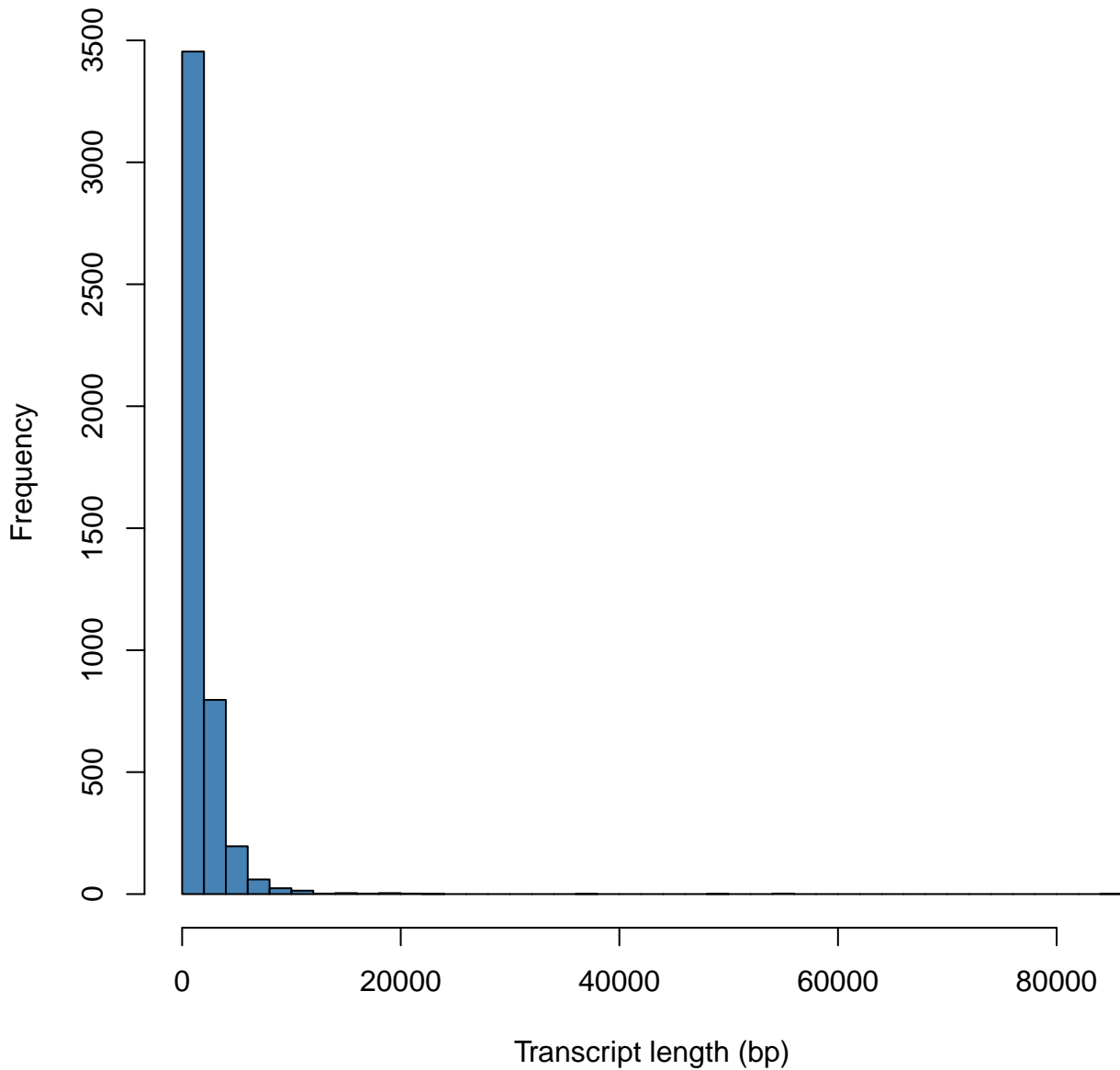


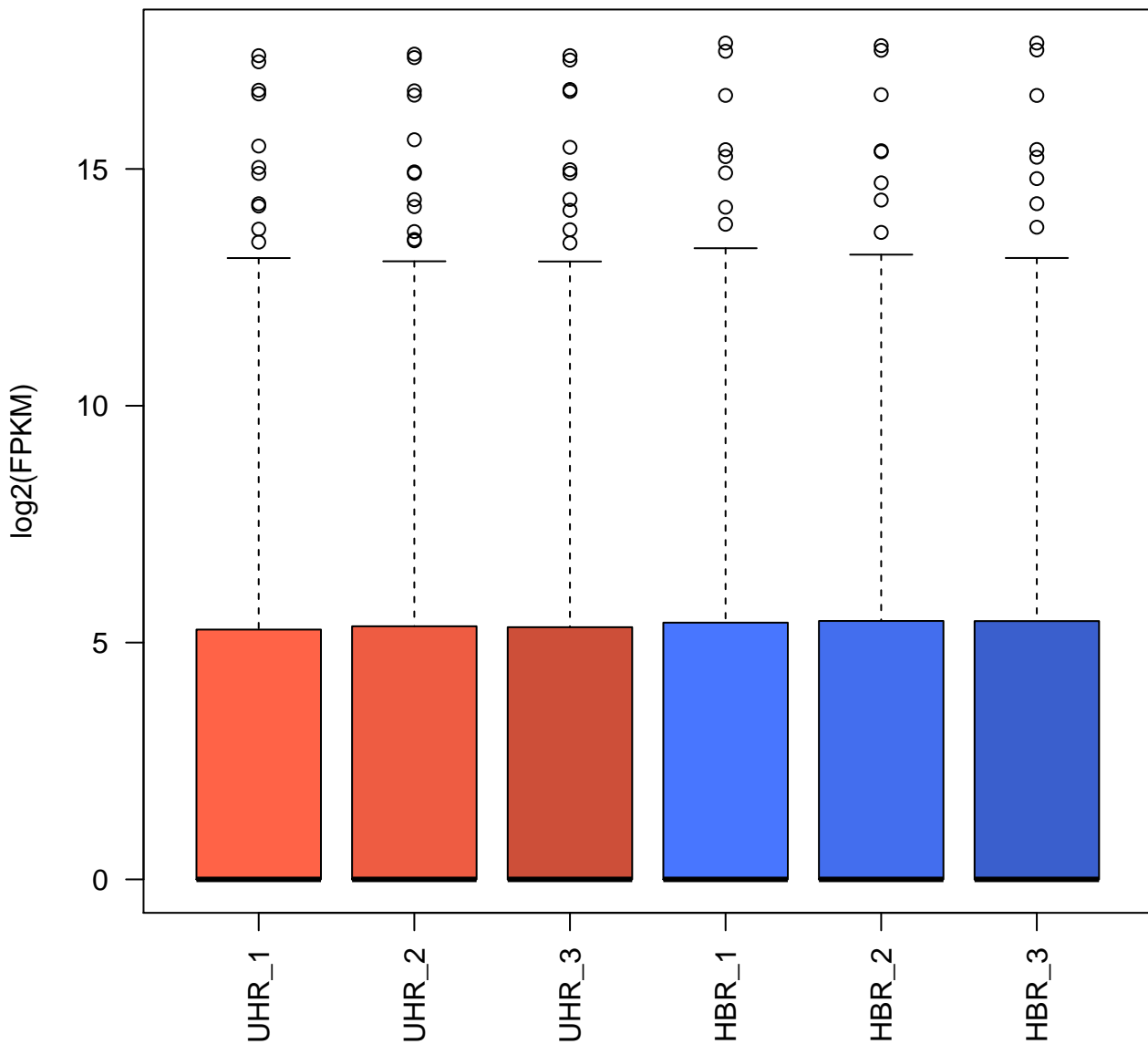
Distribution of transcript count per gene



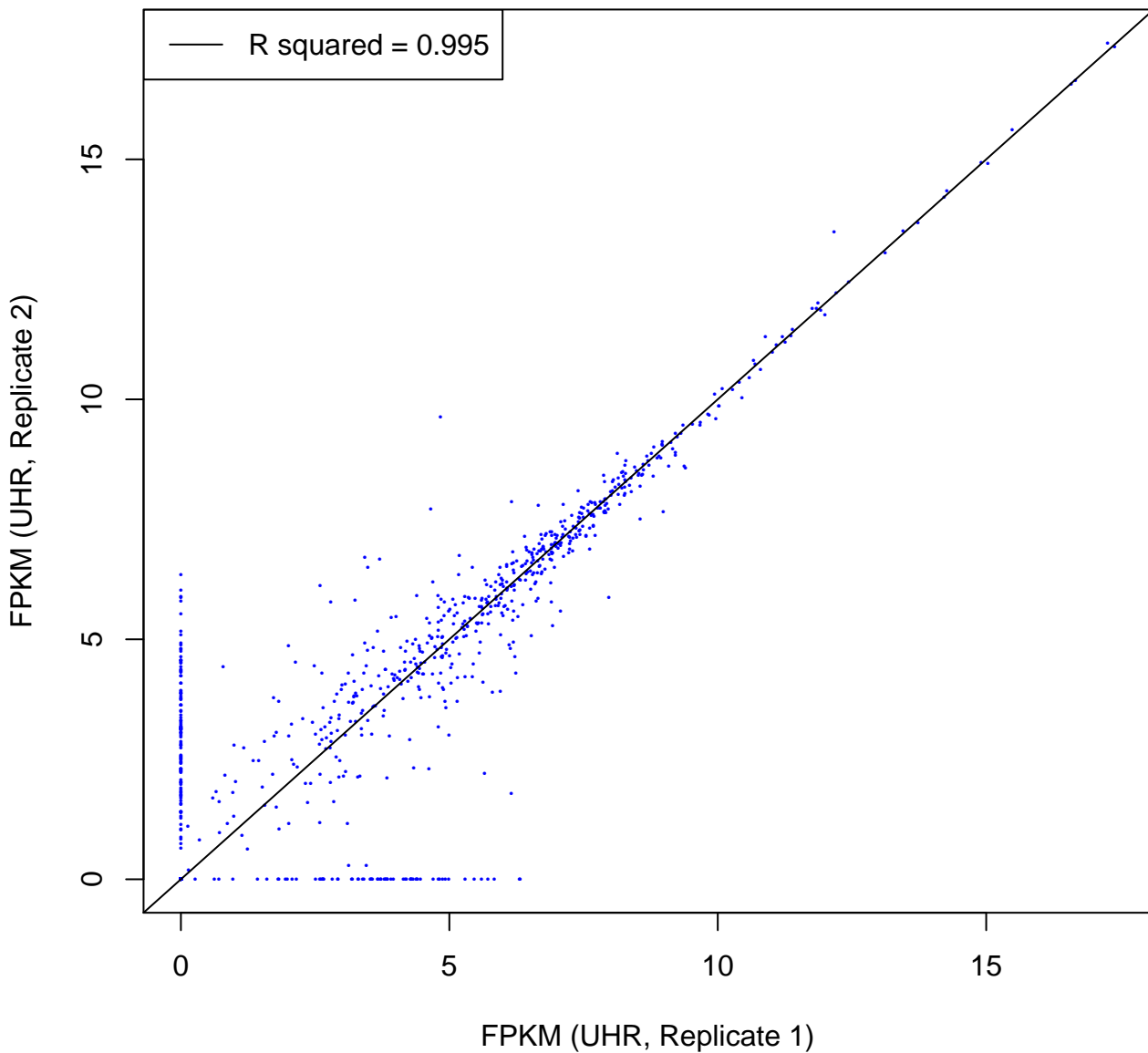
Distribution of transcript lengths



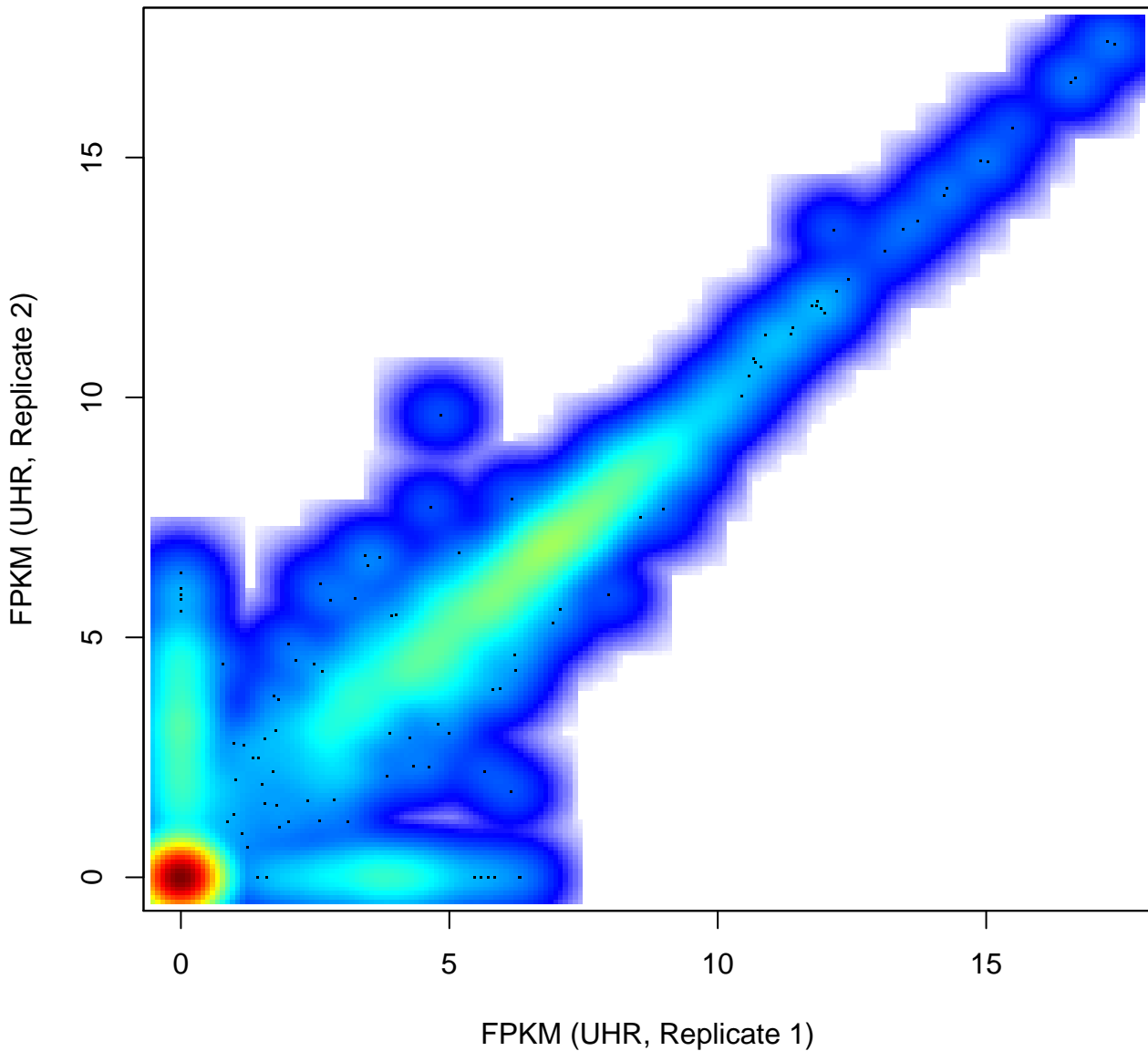
Distribution of FPKMs for all 6 libraries



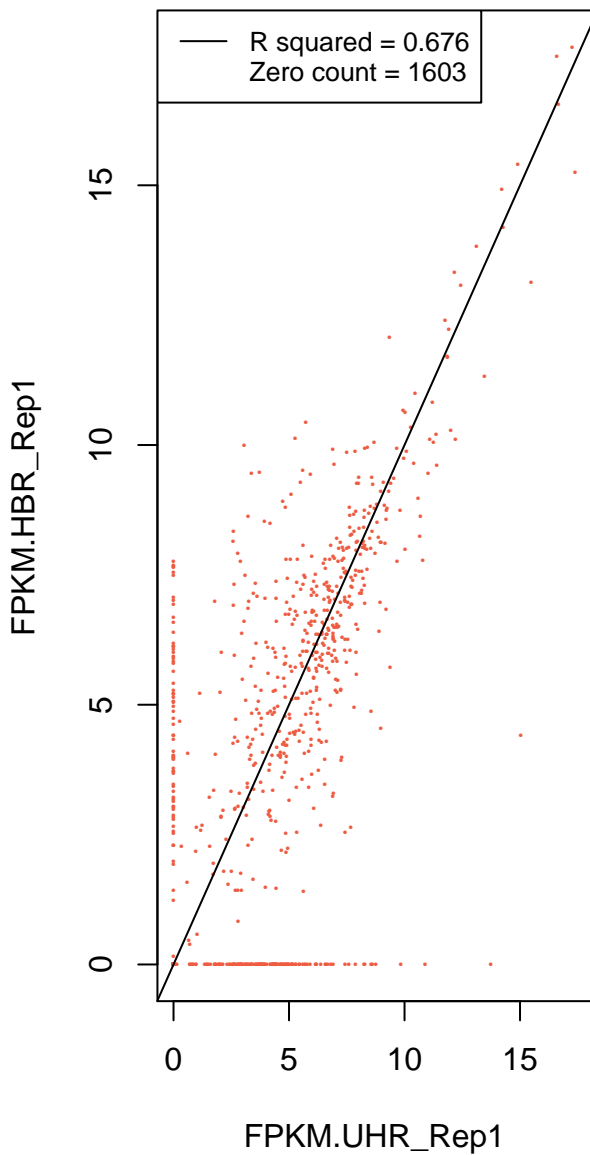
Comparison of expression values for a pair of replicates



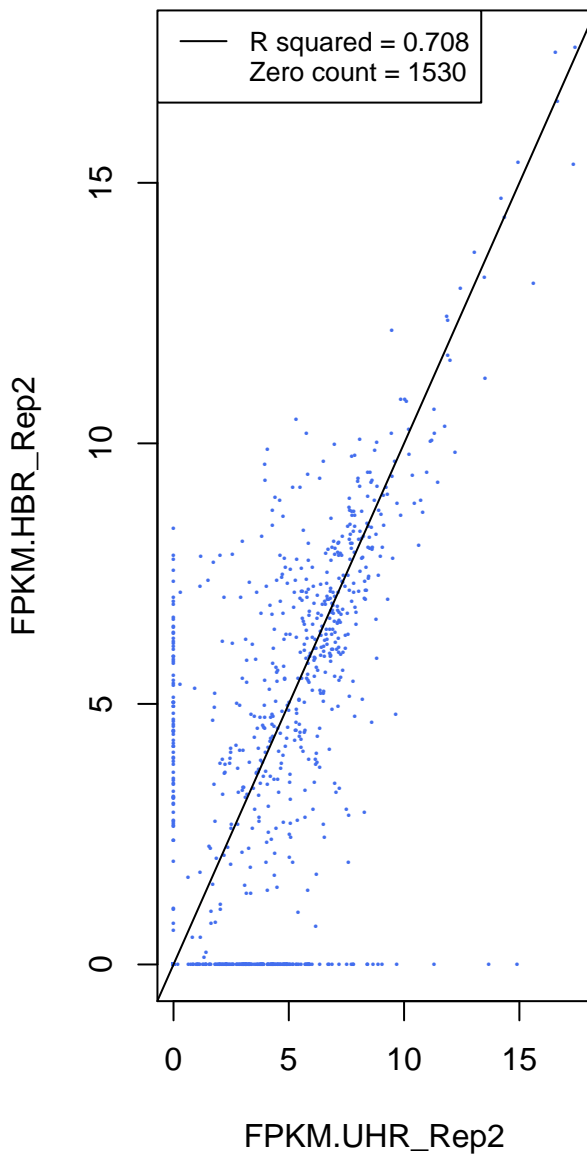
Comparison of expression values for a pair of replicates



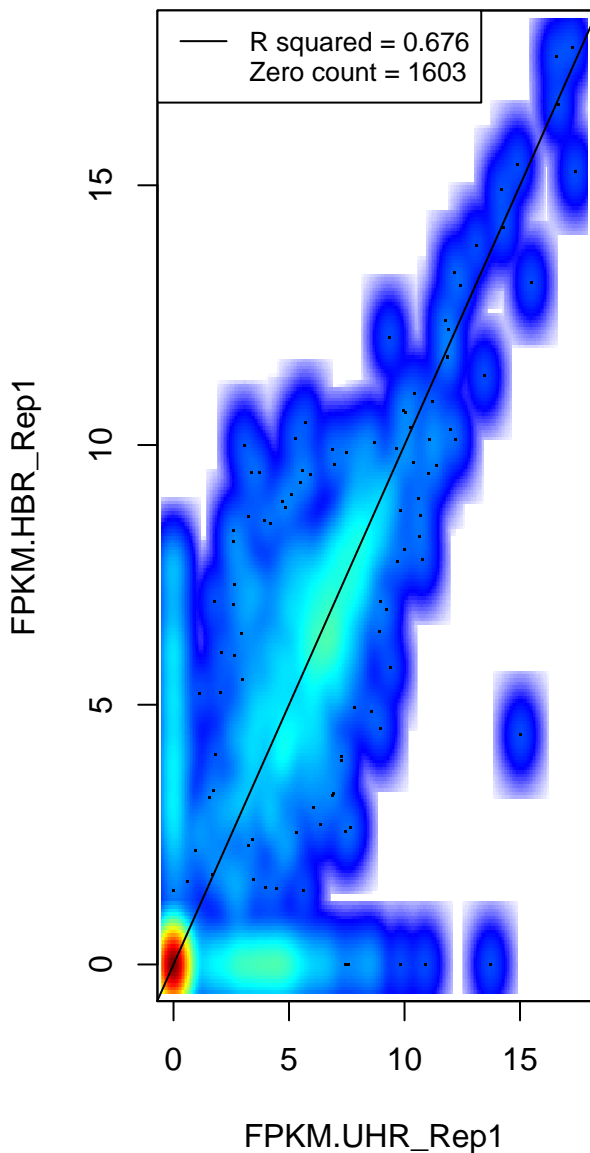
UHR_1 vs HBR_1



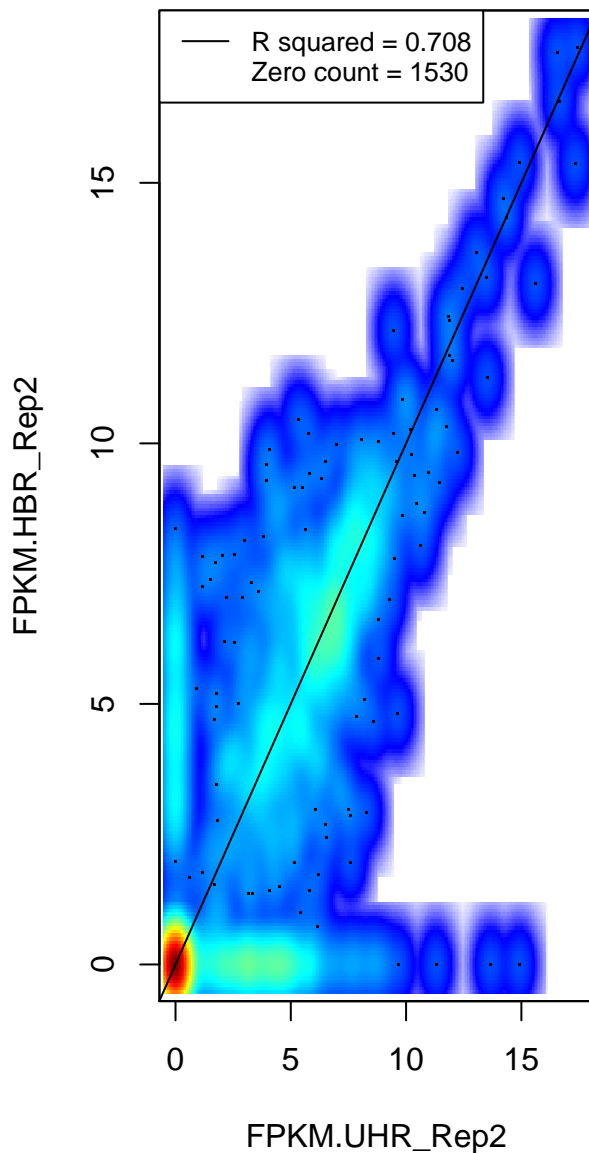
UHR_2 vs HBR_2



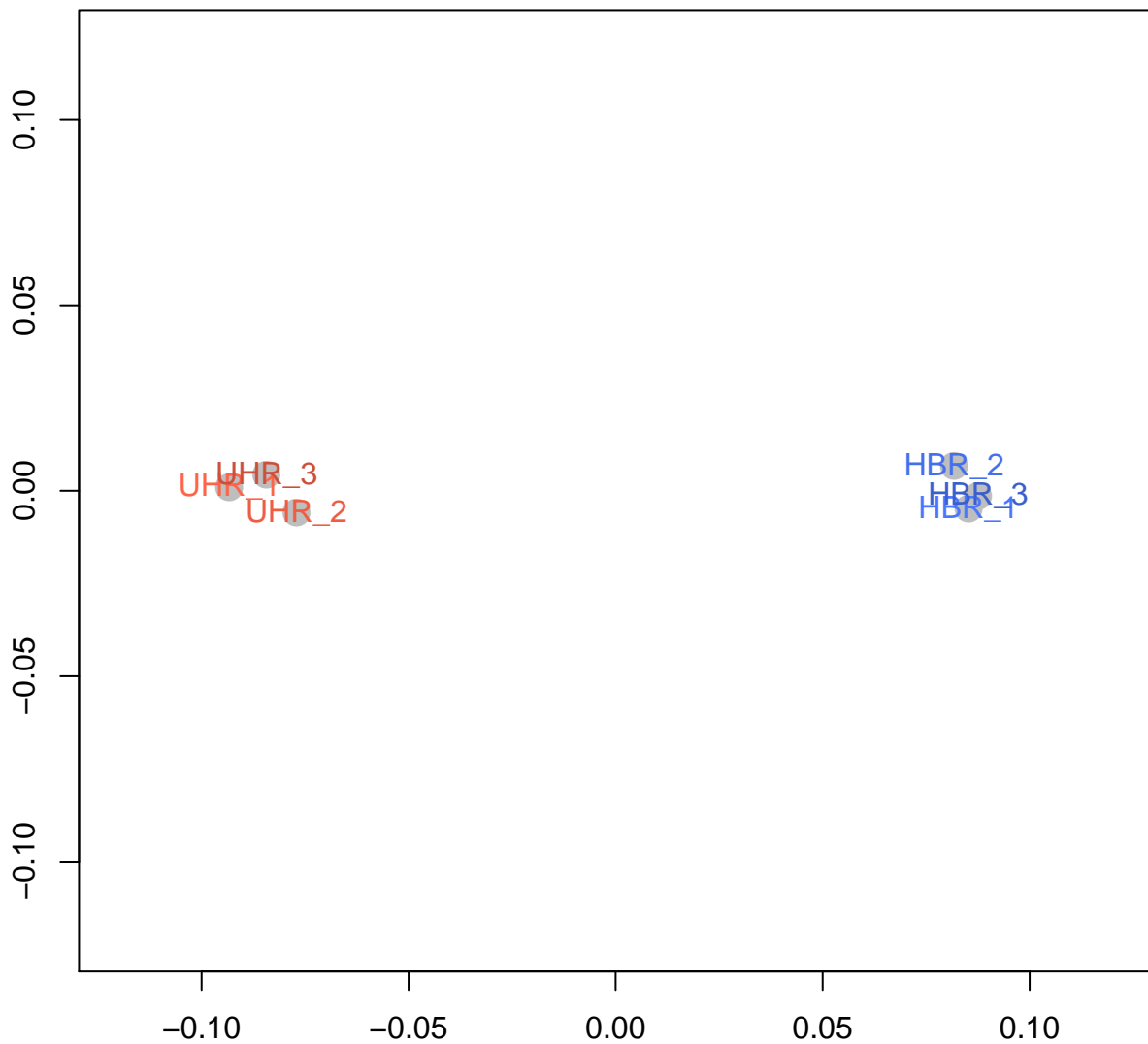
UHR_1 vs HBR_1



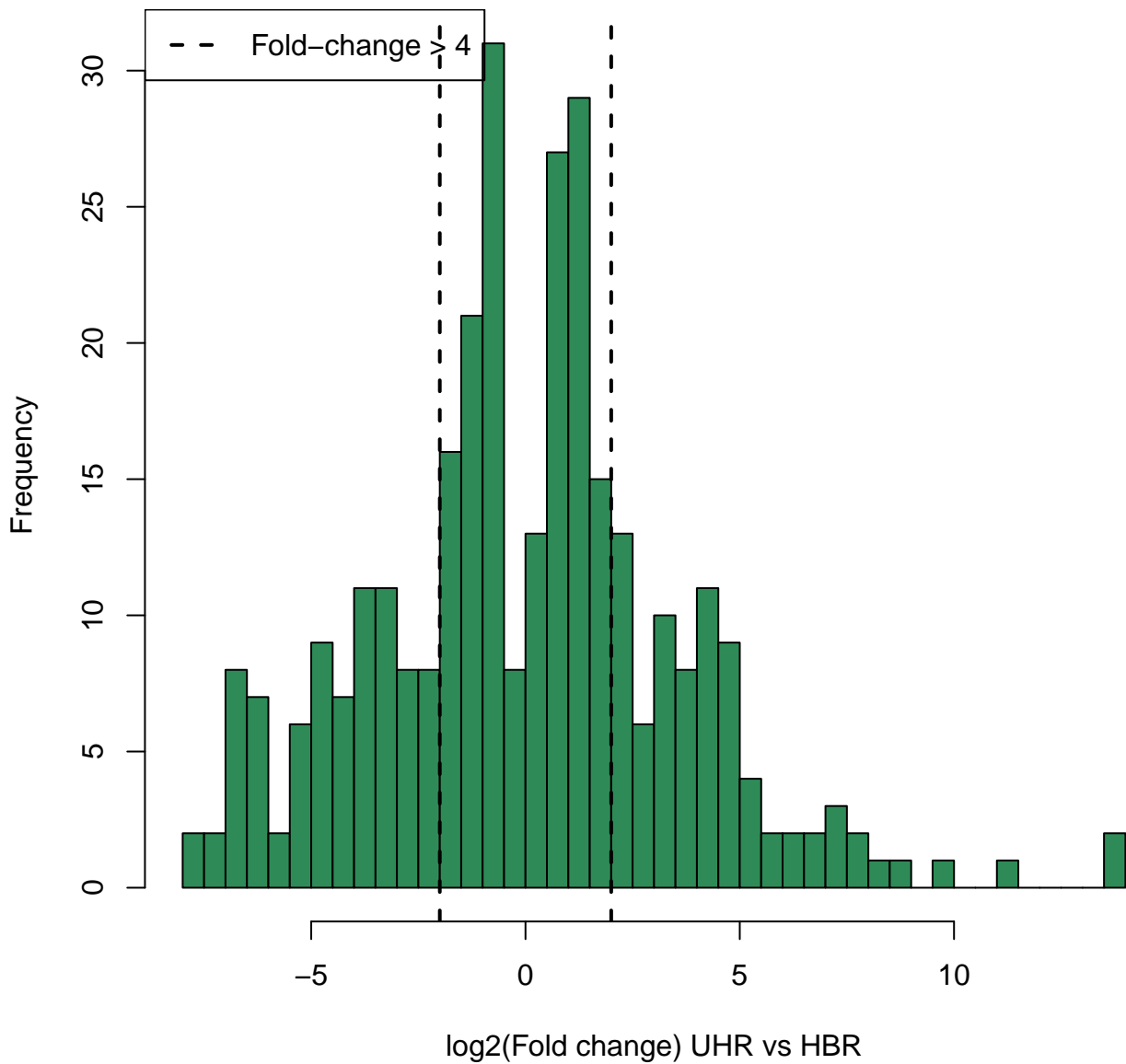
UHR_2 vs HBR_2



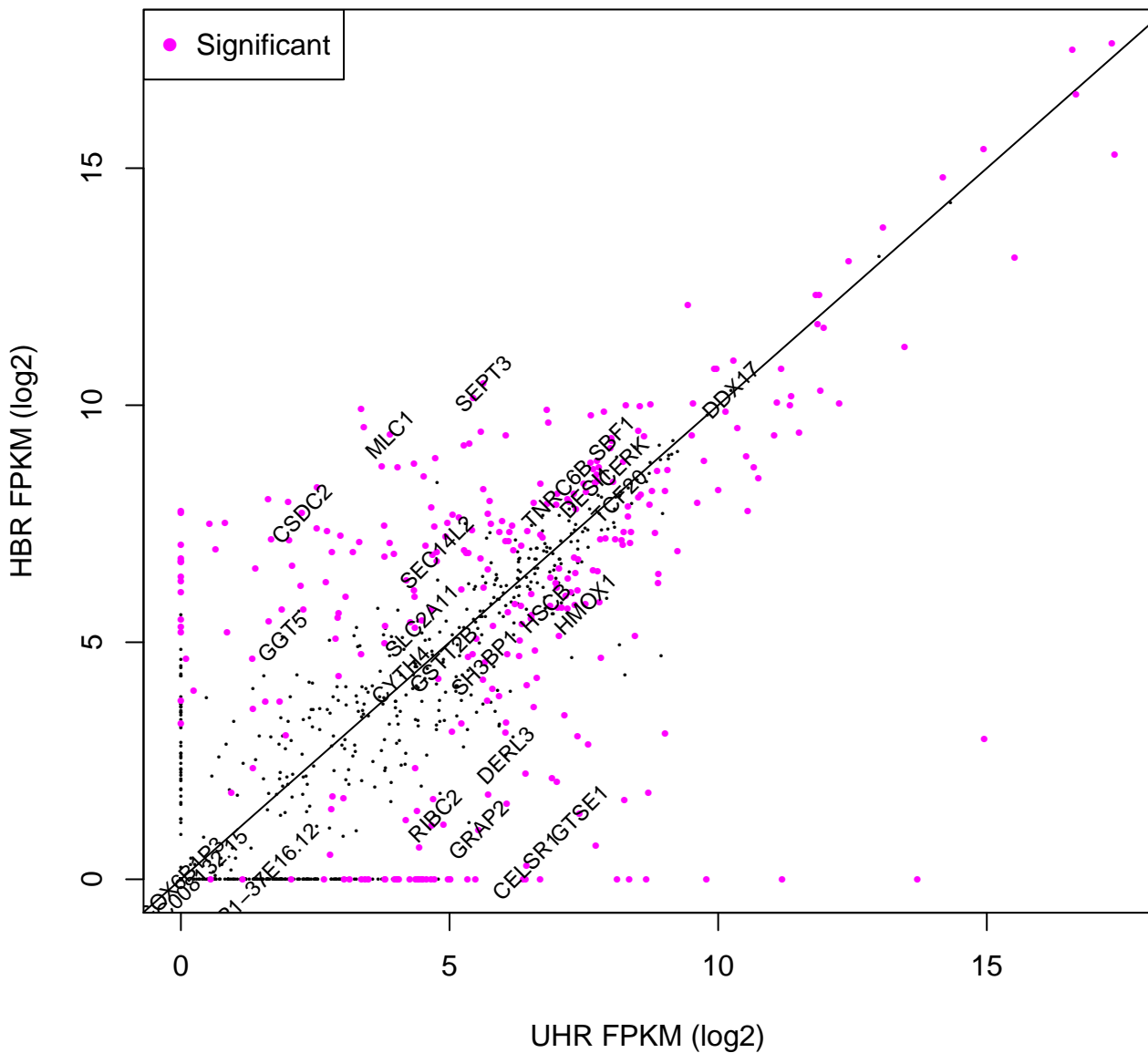
MDS distance plot (all non-zero genes)



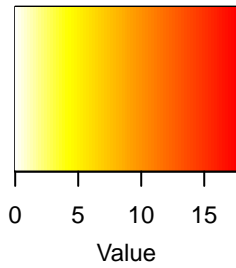
Distribution of differential expression values



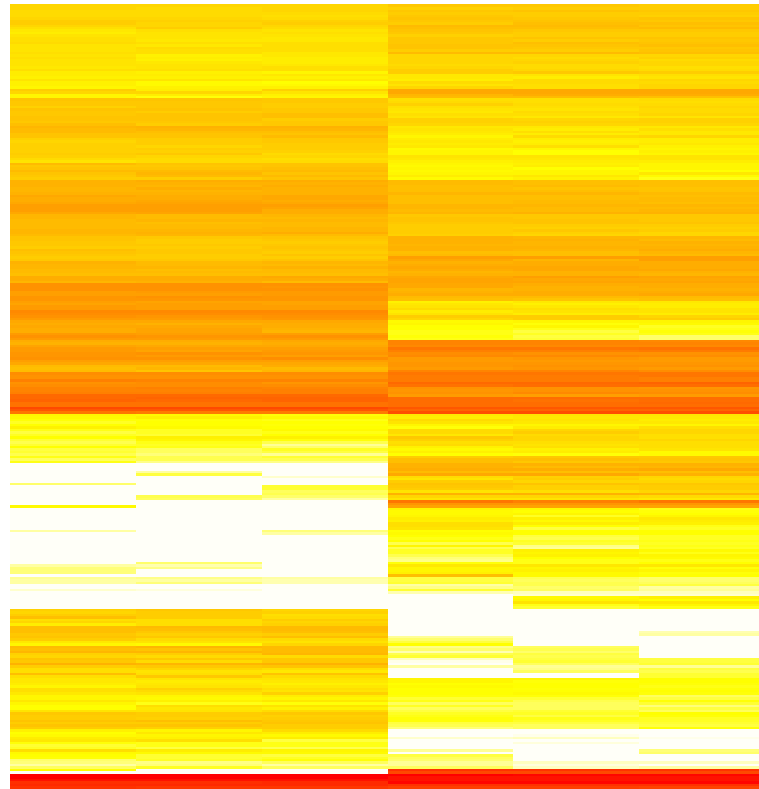
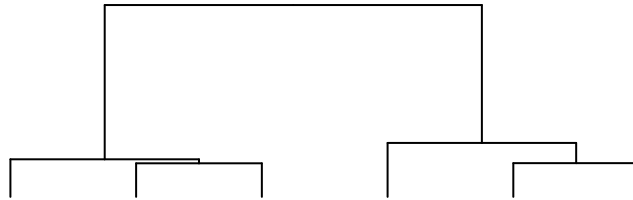
UHR vs HBR FPKMs



Color Key



sig DE Transcripts



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