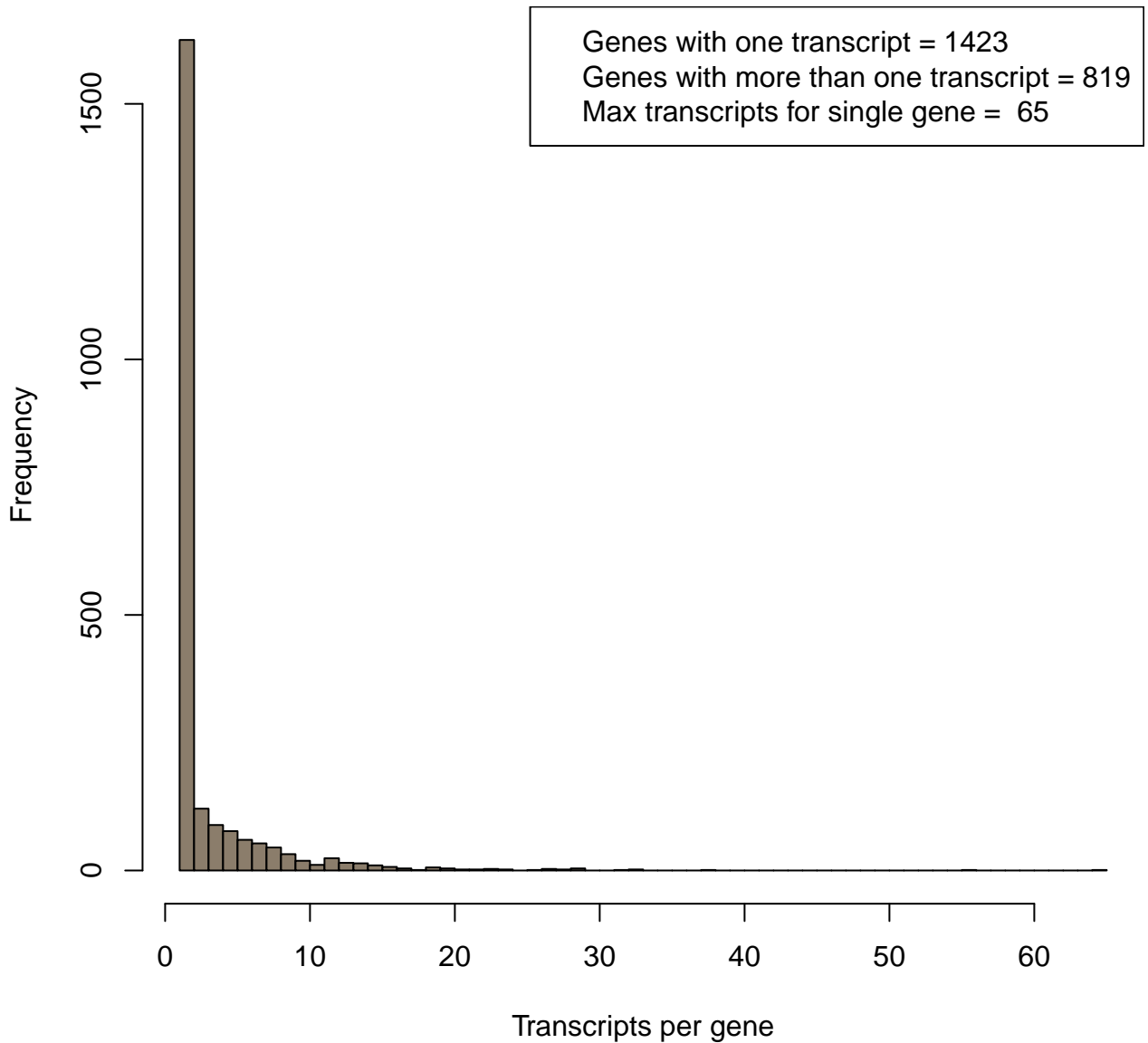
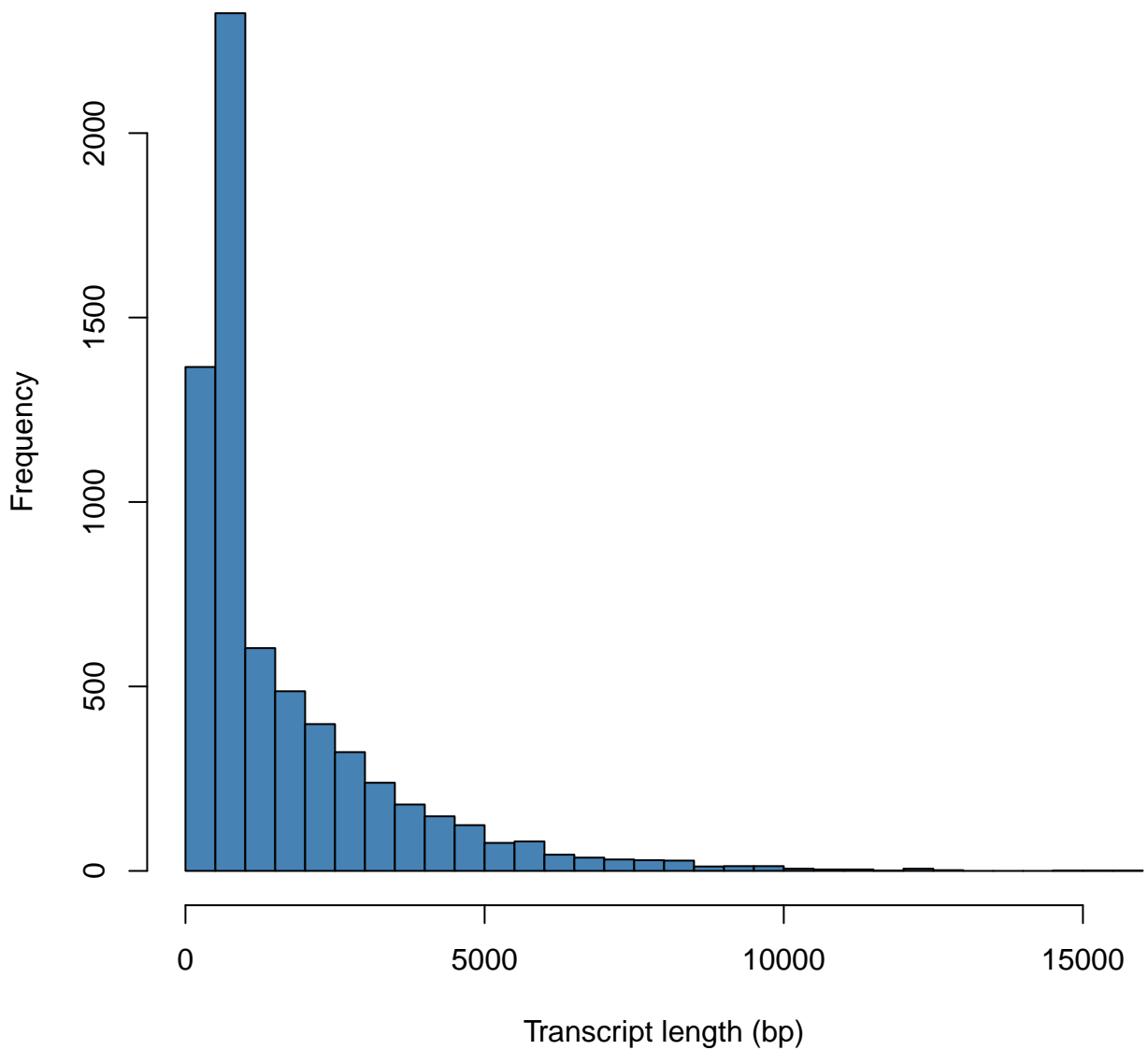


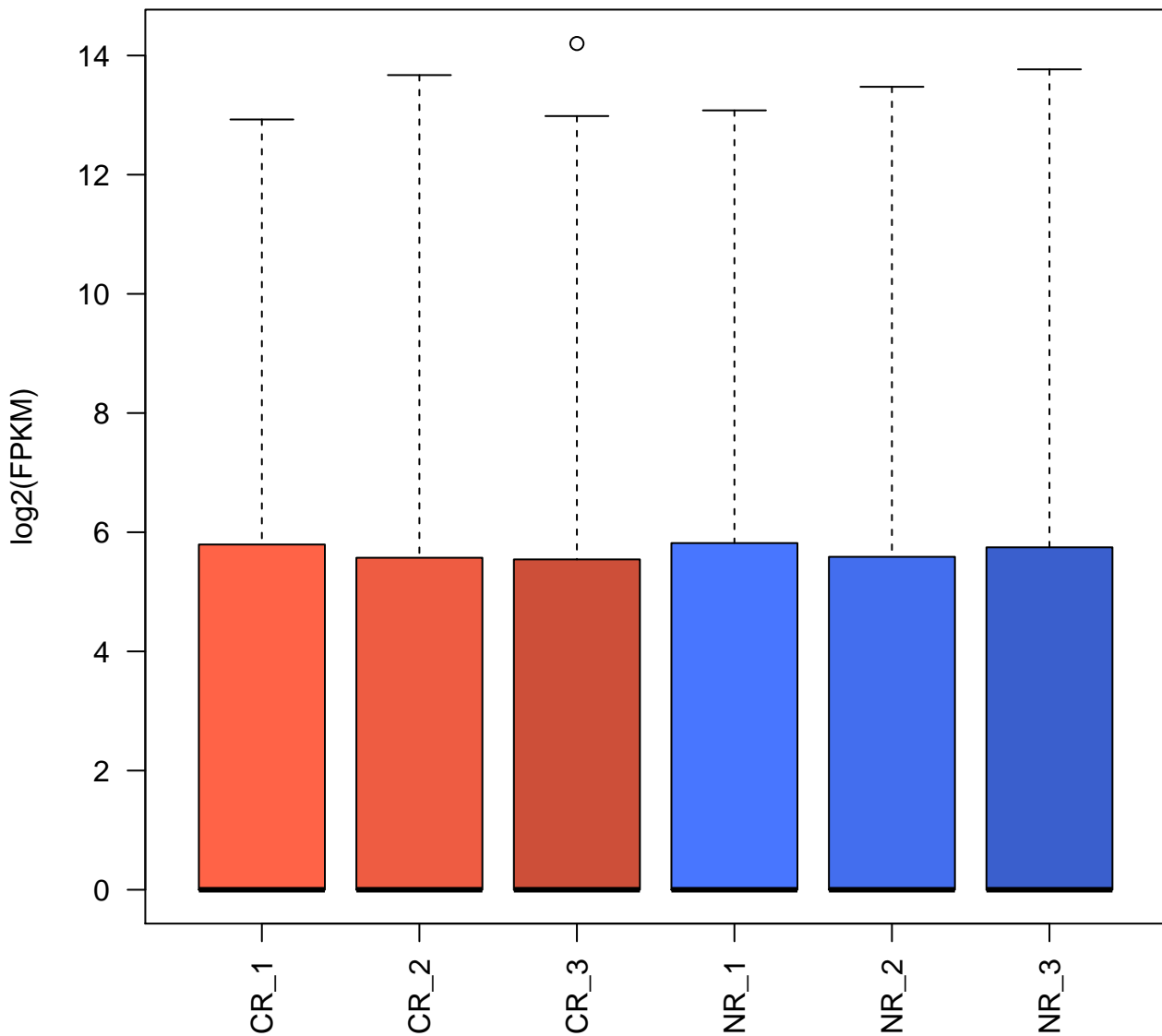
## Distribution of transcript count per gene



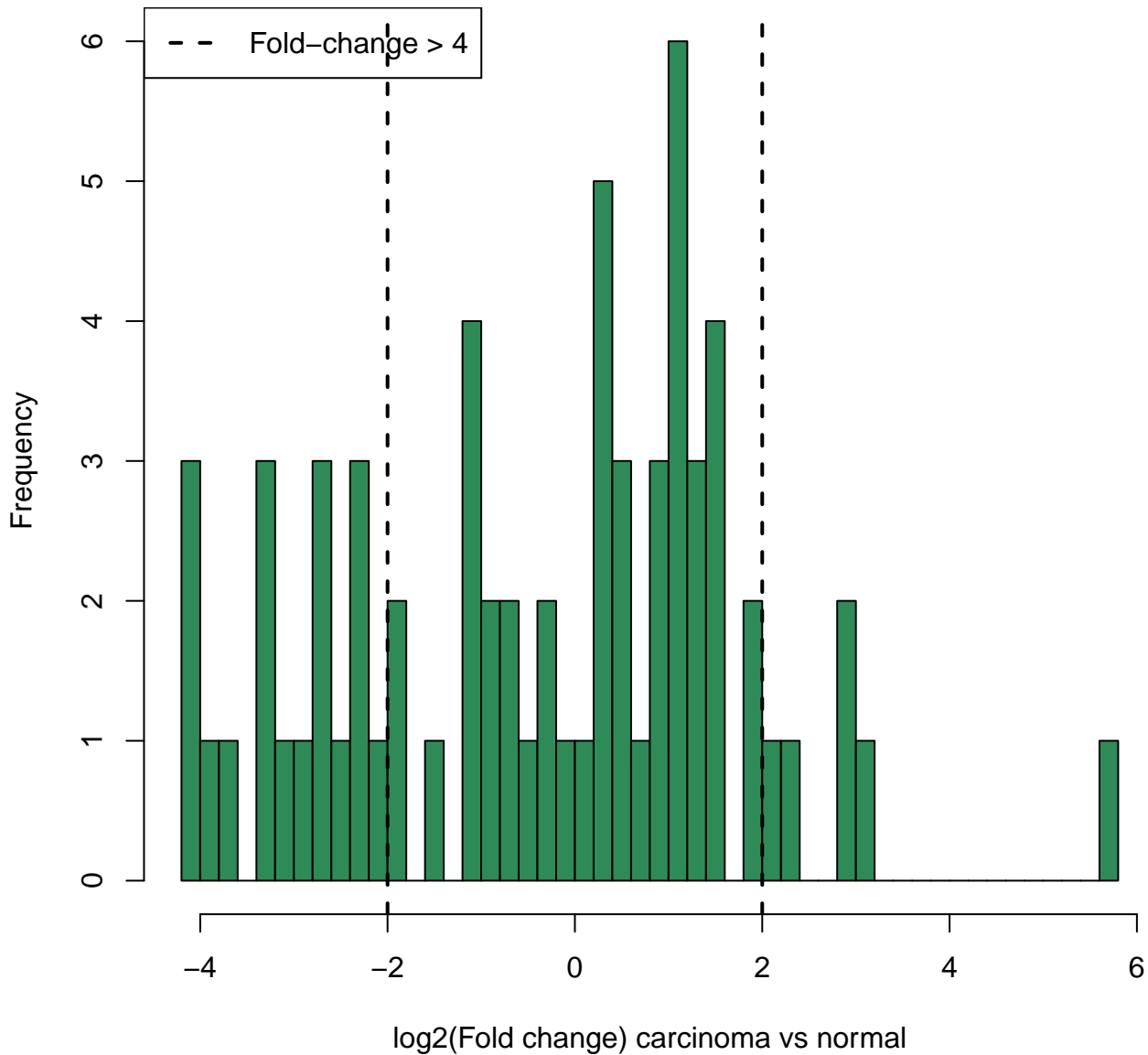
# Distribution of transcript lengths



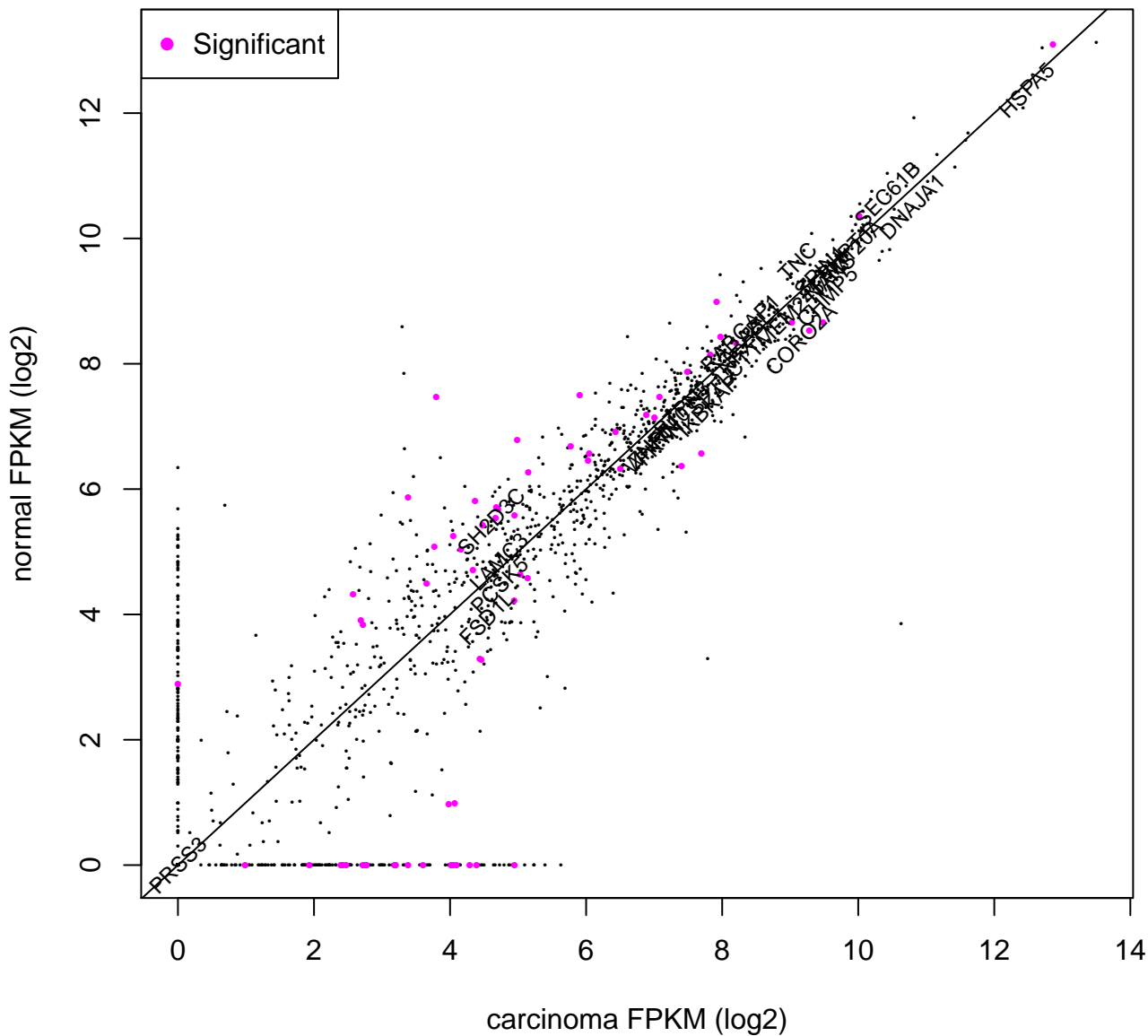
## Distribution of FPKMs for all 6 libraries



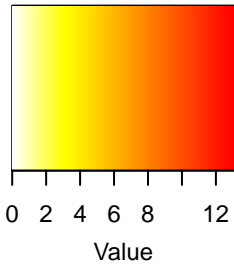
# Distribution of differential expression values



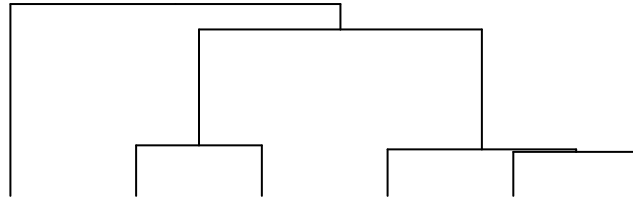
# carcinoma vs normal FPKMs



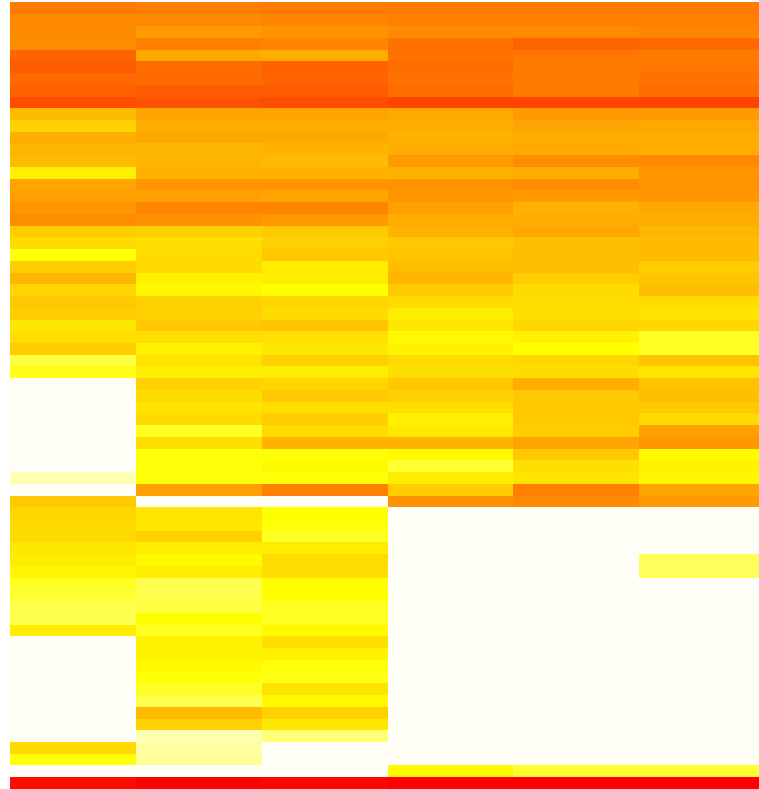
Color Key



# sig DE Transcripts



- RALGDS
- CIZ1
- BAG1
- SNORD36C
- TRM14
- RASEF
- RNF20
- SH3GLB2
- ANF22B
- RP11-172F4.2
- EGFL7
- SLC25A25-AS1
- ANKK6
- PTGES
- CARD19
- MIGA2
- MISANTD3
- GALT
- DENND4C
- ATP9HP2
- WDX21
- NR6A1
- GGTAP1
- SNORD36A
- FGD3
- CEP78
- CBWD1
- ZNF169
- FKBP4P8
- RP11-70L8.4
- FRMD3
- LIX6
- FANCG
- DBH-AS1
- LCN10
- SPK8
- PLPP7
- RP11-92C4.6
- RP11-39B21.11
- C6orf198
- ANKRD19P
- RPL10P3
- SNRPEP2
- RP11-216B9.6
- CH17-29N19.1
- SETP5
- CNTNAP3P1
- RP11-452D2.2
- RP11-216M21.1
- SNX18P4
- SNX18P8
- RP11-149F8.3
- C2orf27AP3
- TME1F1
- FAM166A
- RP11-537H15.4
- CTSV
- FIBCD1
- PAPPA-AS2
- ARMCBP1
- RP11-311H10.7
- TME1G10
- RP11-47924.8
- SLC31A2
- CR2
- PES1P2
- EEF1A1P5



rcinoma\_C03

rcinoma\_C06

rcinoma\_C02

1.normal\_N03

1.normal\_N06

1.normal\_N02