

Figure 1

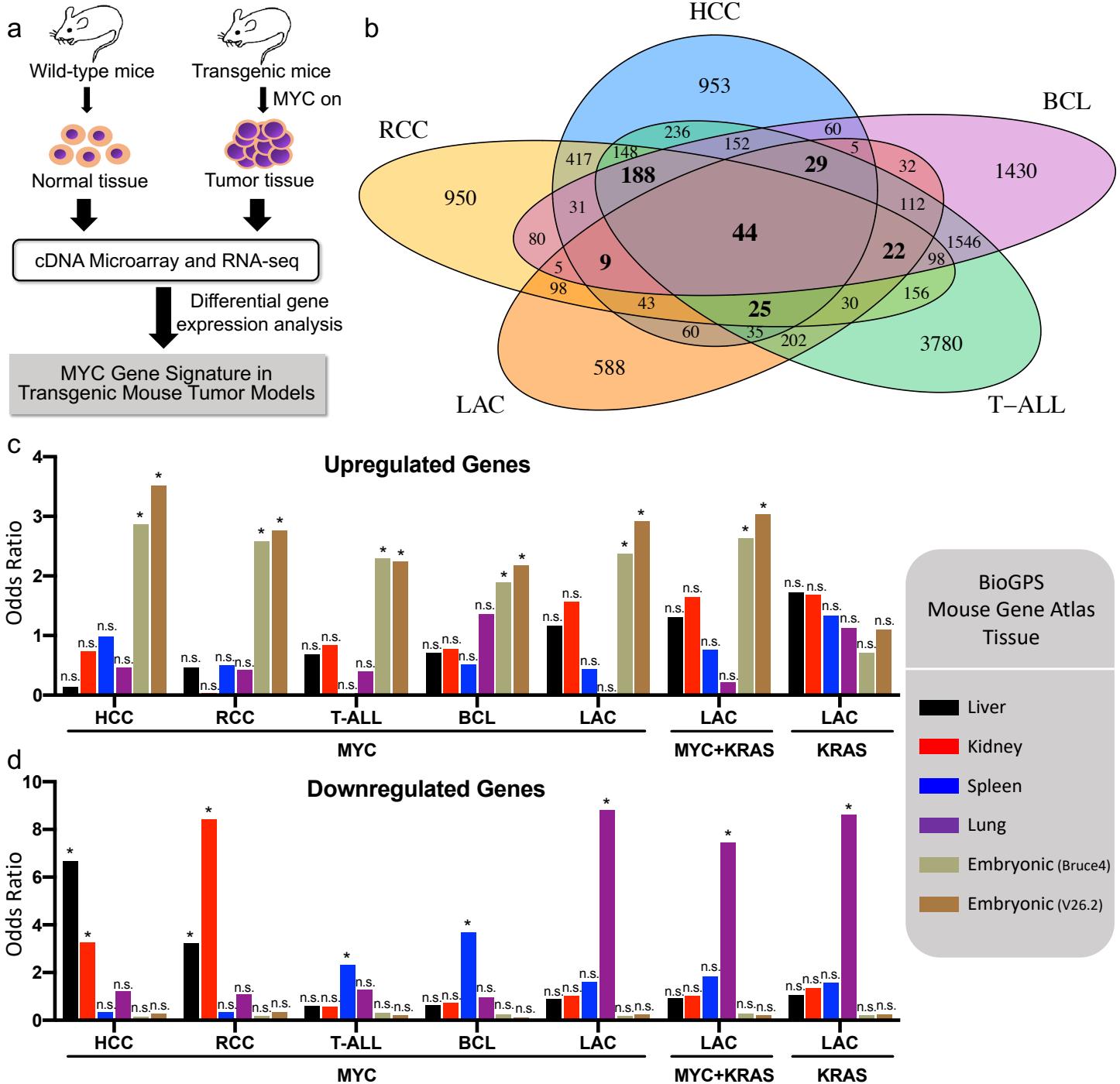


Figure 2

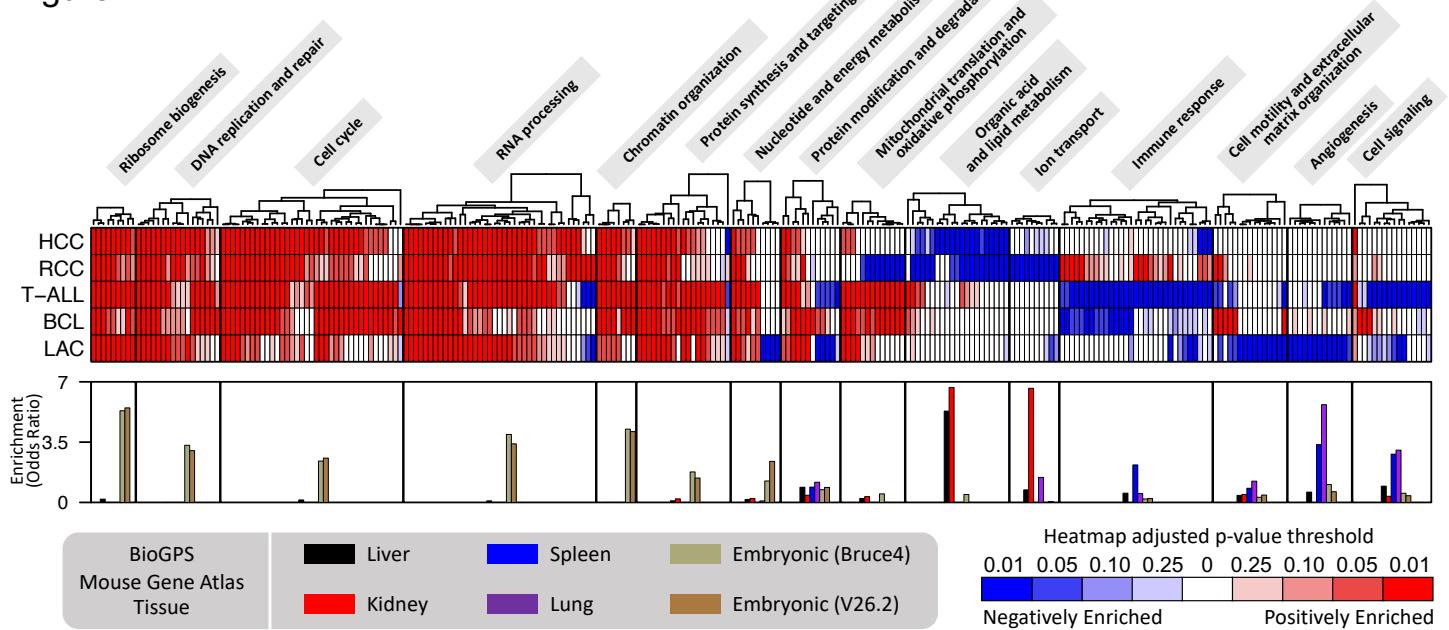


Figure 3

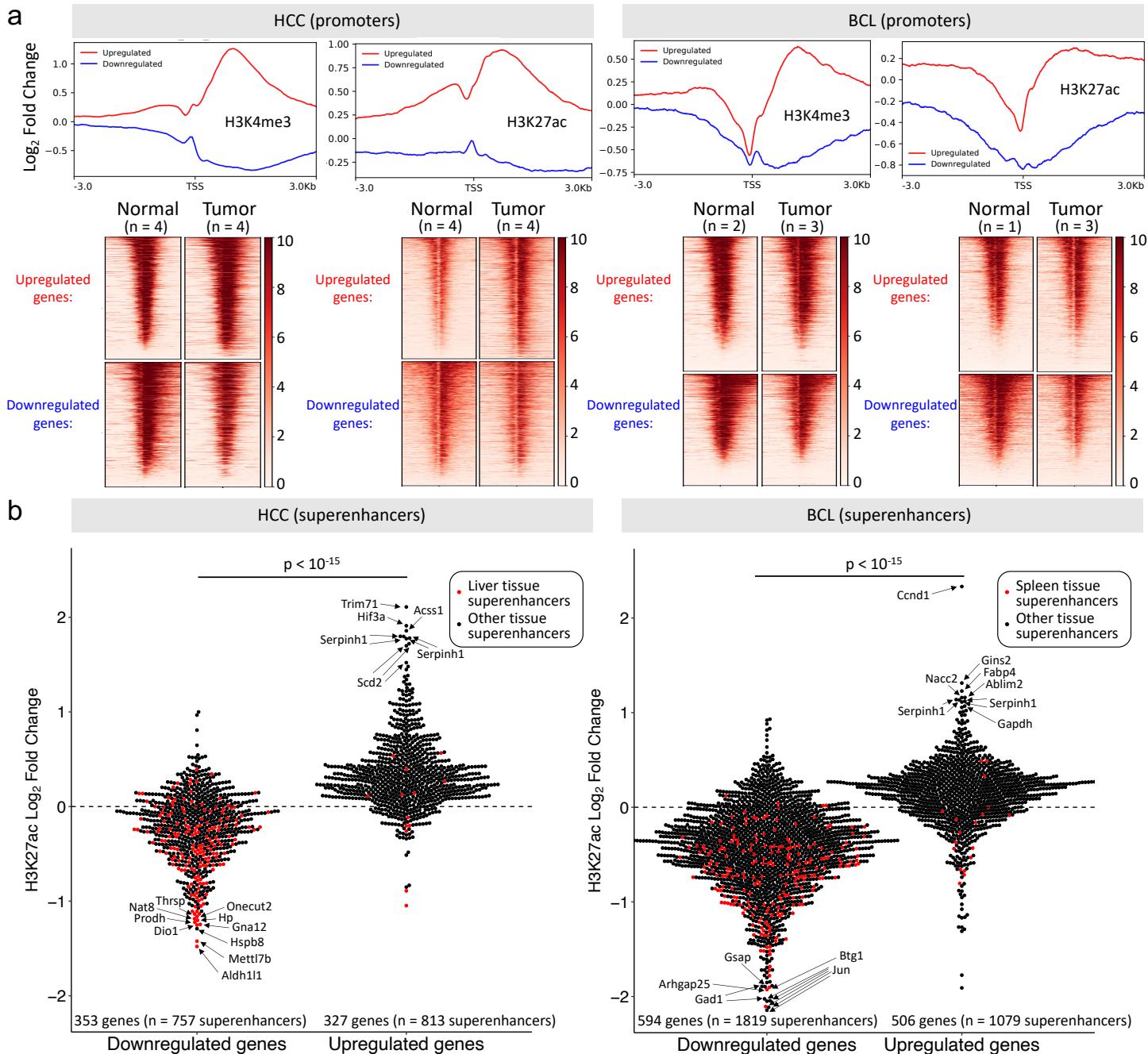


Figure 4

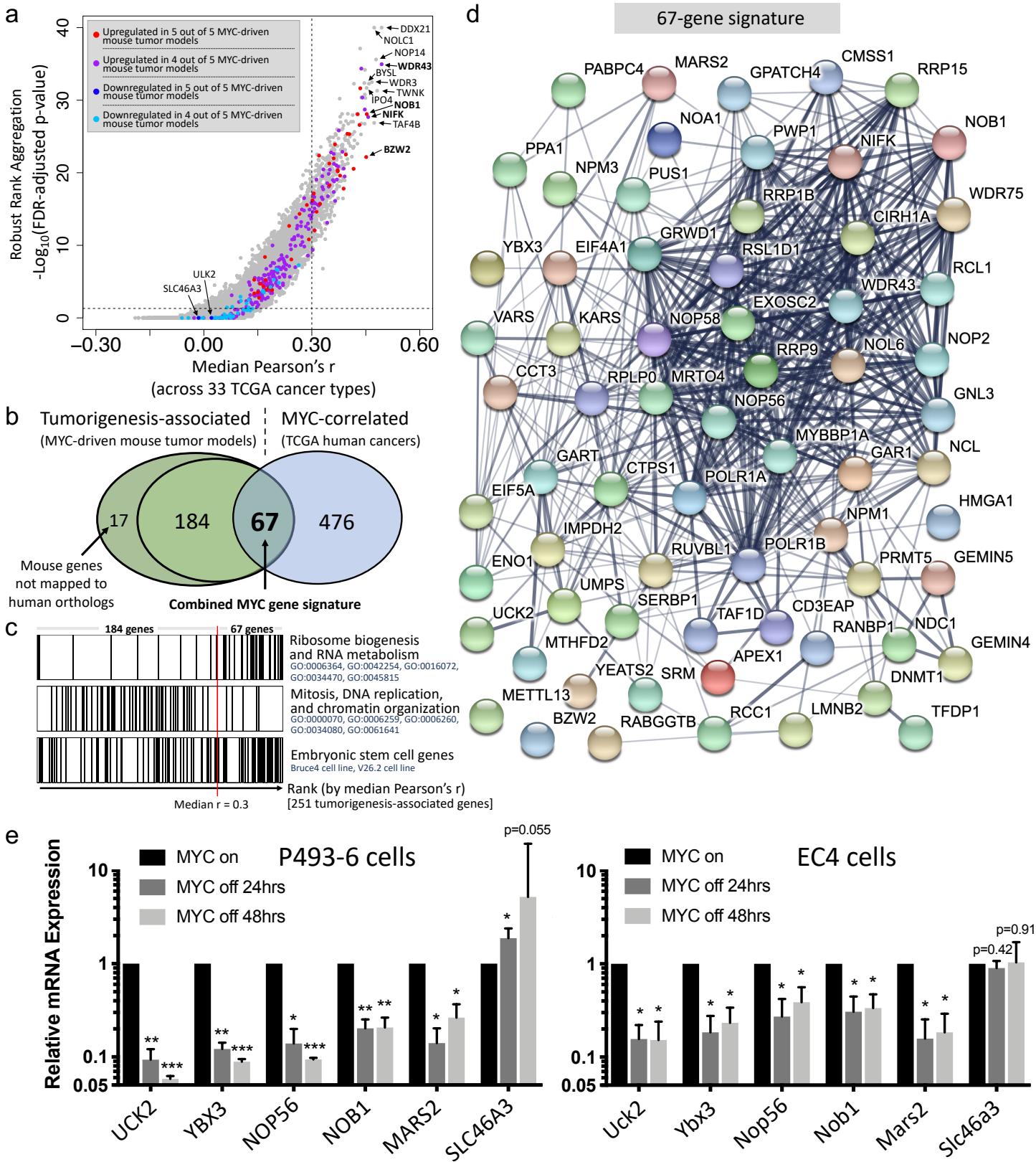


Figure 5

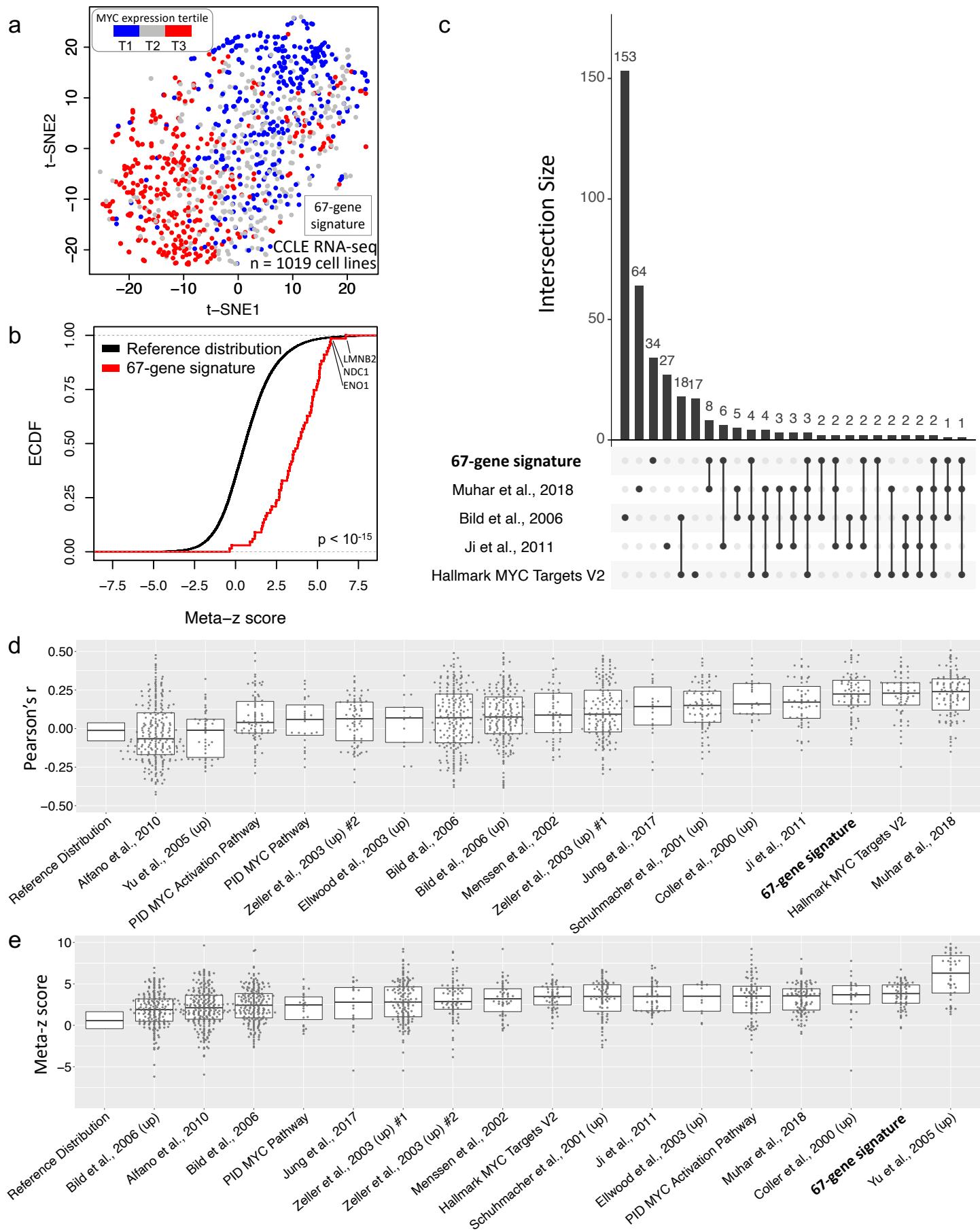


Figure 6

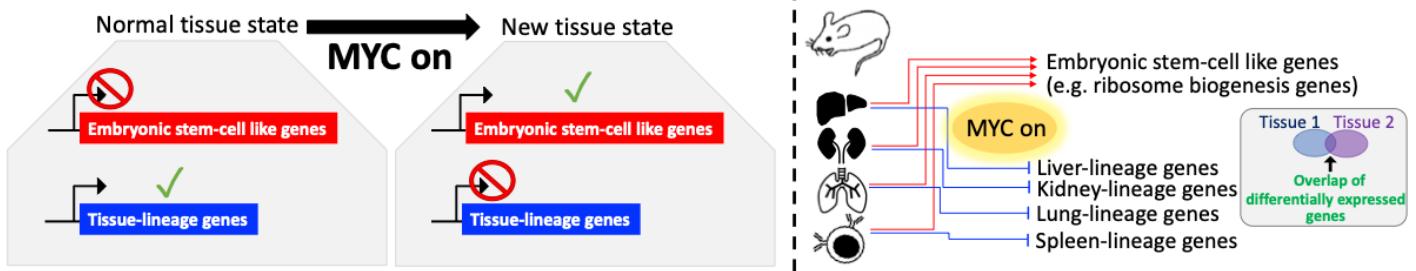


Table 1

| Tumor Type | Mouse Background | Mouse Genotype | Number of Samples | |
|--|------------------|--|-------------------|-------------|
| Hepatocellular carcinoma (HCC) ¹ | FVB/N | LAP-tTA/TetO-MYC | n = 4 liver | n = 6 tumor |
| Renal cell carcinoma (RCC) ¹ | FVB/N | GGT-tTA/TetO-MYC | n = 3 kidney | n = 3 tumor |
| Lung adenocarcinoma (LAC) ¹ | FVB/N | CCSP-rtTA/TetO-MYC | n = 2 lung | n = 5 tumor |
| | FVB/N | CCSP-rtTA/TetO-KRAS ^{G12D} | | n = 5 tumor |
| | FVB/N | CCSP-rtTA/TetO-MYC/TetO-KRAS ^{G12D} | | n = 4 tumor |
| T-cell acute lymphoblastic lymphoma (T-ALL) ² | FVB/N | E μ SR α -tTA/TetO-MYC | n = 3 spleen | n = 3 tumor |
| B-cell lymphoma (BCL) ³ | C57/Bl6 | E μ -Myc | n = 4 spleen | n = 3 tumor |

¹Microarray data (GEO accession number: GSE143254)
²RNA-seq data (GEO accession number: GSE106078)
³RNA-seq data (GEO accession number: GSE51008)