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| **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-KRASG12D | n = 5 tumor |
| FVB/N | CCSP-rtTA/TetO-MYC/TetO-KRASG12D | n = 4 tumor |
| T-cell acute lymphoblastic lymphoma (T-ALL)2 | FVB/N | EμSRα-tTA/TetO-MYC | n = 3 spleen | n = 3 tumor |
| B-cell lymphoma (BCL)3 | C57/Bl6 | Eμ-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) | | | | |