1. https://cran.r-project.org/web/packages/dnet/index.html

   Paper: Transfer Learning for Survival Analysis via Efficient L2,1 -norm Regularized Cox Regression ([3] in Asymmetric Graph…paper)

2. https://portal.gdc.cancer.gov/

   Medical data query platform (official)

3. https://www.bioconductor.org/packages/release/data/experiment/vignettes/RTCGA.mRNA/inst/doc/downloading_mRNA_datasets.html

   This tool showed up in one paper. I didn't check this one carefully. But it can be an alternative.

4. https://github.com/compgenome365/TCGA-Assembler-2

   Many papers used this tool (with detailed tutorial inside). There are many tools available in Github that help us download datasets and do some preprocessing work on them. You can just google "TCGA download Github". But the one mentioned here appears in several papers which shows that it's a useful tool. I don't remember which papers used it. When I find out later, I will add the papers.