

The way to download the raw datasets with TCGA2STAT

Reference: <http://www.liuzlab.org/TCGA2STAT/>

Please notice: after executing “install.package(TCGA2STAT_1.0.tar.gz”.....), you also have to enter:

```
if (!requireNamespace("BiocManager", quietly = TRUE))
  install.packages("BiocManager")

BiocManager::install("CNTools")
```

You can directly download the file into csv file with the following:

```
write.csv(mirseq.ov[["merged.dat"]], "path")
```

If you want to download more datasets and know more details, please click the reference link.