

# Differential Analysis

FDRdb allows you to upload your own RNA expression profile data for calculation and analysis. You should prepare two kinds of profiles:

Expression profile ([Download example file](#))

Sample class profile ([Download example file](#))

③

Both of the profile must be uploaded in .txt format.

As uploading the data, you can select the species for which the data correspond to get the fibrosis-related RNA results. Point the “Apply” button, the FDRdb will analyze the data you upload and visualize the results. **Please pay attention! If you want to get fibrosis related RNA results, the RNA label in your profiling data must follow the rules.**

See the "Help" page for more details.

*\*Please upload the **expression profile** here.*

①

Browse...

*\*Please upload the **sample class file** here.*

②

Browse...

*You can select the species that corresponds to your file sample to obtain fibrosis related results.*

④

None



⑤

Apply