

**DDS document:**

This package will only accept Ct values in .csv files. Example datasets can be found in the /data folder of this repository. Please convert your Ct values to the appropriate format and note for polysome profiling and pfaffl analysis both technical and experimental replicates are required for input. This is to ensure rigor and reproducibility of science. Pfaffl method also requires a primer efficiency calculation to ensure accuracy of data.

**Updates:**

Will continue to work on the design by incorporating the ability to perform statistics. These include ANOVAs and t-tests on the data. Also will incorporate the ability to save your csv files. Hojae added functionality to save plots, so I will update the tutorial to show that. Finally will try to improve data importing by allowing for more input by the users