

## Submission Instructions

1. Please fill out a survey for every different condition that you use. E.g, if you ran a Taqman Assay and a SYBR green assay. It is easy to submit more than 1 assay. After submitting the first assay, use the BACK button on your browser. It will take you back to the form that you just filled out and then you need only change the variable(s) which might be different. If you click on the return link on the thank you page, then it will return you to a blank survey and you will have to refill all the answers.
2. Please e-mail the following to "Gregory L Shipley" [Gregory.L.Shipley@uth.tmc.edu](mailto:Gregory.L.Shipley@uth.tmc.edu)
  - A copy of you exported data labeled with your 4 digit ID code and exportdata. E.g, 1234exportdata. The export data should contain (minimally) the Ct for each well, whether there was DNA or RNA in the well, the amount of DNA or RNA in the well
  - A screen capture( pdf or jpg) of the amplification curves for DNA labeled 1234DNAamp
  - A screen capture( pdf or jpg) of the amplification curves for RNA labeled 1234RNAamp
  - A screen capture( pdf or jpg) of the Standard curve for DNA labeled 1234DNAstd
  - A screen capture( pdf or jpg) of the Standard curve for RNA labeled 1234RNAstd

A copy of these instructions can be downloaded from

[http://www.abrf.org/index.cfm/group.show/NucleicAcids.32.htm#R\\_4](http://www.abrf.org/index.cfm/group.show/NucleicAcids.32.htm#R_4).

They are also available on the “ View Questions to be Answered” which is available from the same site.