Final output: REMEMBER!!!!! – THIS MUST BE DONE WITH RN6 ONLY

* Whole genome:
  + Dispersion plot or FPKM SCV plot – quality control of how replicates match
  + Density plot - compares 2 conditions
  + Box plot – FPKM distrubutions of 2 conditions
  + Scatter plot of global changes between conditions
  + MAplot
  + Volcano plot
  + Appendix page of basic cuffdiff output
  + Overveiw of significant features:
    - Significant genes graph
    - K-means clustering graph – groups genes with similar expression profile
* Gene sets:
  + Heat map – overall expression changes in colour
  + csScatter – diff between conditions
  + volcano plot
* Individual genes:
  + We want to know:
    - Whole gene expression change between conditions
    - Each isoform expression change between conditions
    - Identify TSS of each isoform
    - Similar genes
  + How to show that:
    - expressionPlot – shows expression changes of both whole gene and all isoforms
    - Bar plot – does same
    - Pie charts – showing individual isoform abundance per condition
    - findSimilar – shos similar genes (9.6 on .pdf)