

NetID: au198  
Aizhan Uteubayeva

Step 2  
Final Project: Enrichment analysis using EnrichR  
HIDS-7003-01

**Goal:** Use the t-test results and shortlist based on p-value  $< 0.001$  in order to identify various functional groups of the selected genes

**Workflow:**

- 1) Import T-test results for further shortlisting
  - a) File that contains group comparison results
- 2) Filtering/shortlisting
  - a) Short listed the results based in p-value cut off (p-value  $\leq 0.001$ )
- 3) Export the short listed results for reference
- 4) Clean gene names
  - a) Genes name are clean and ready to proceed - 821
  - b) Cleaning the gene list from duplicates and only leaving unique genes - 702 genes were discovered.
- 5) Loading Databases for EnrichrR package
  - a) Use the list of databases for which enrichment analysis will be run
- 6) Calling function to run Enrichment