REPORT:

1) Need to analyze gene expression values grouped by disease type. MongoDB has well-documented table import and basic-statistics functions that makes it a suitable choice.

2) The documents have PATIENT\_ID, DIAGNOSIS, and all entrez\_ids as their field. This allows to avoid costly updates that would be cause by smaller documents (e.g., every gene value as a document)

3) Just enter an entrez id of the gene of interest.

4) 