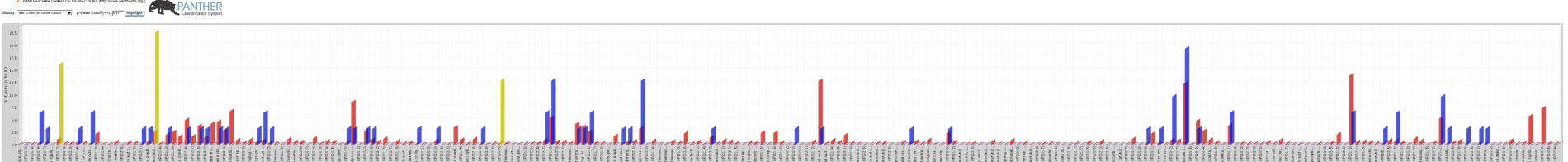
PANTHER BAR CHART OF GENE COUNT (http://www.pantherdb.org/)

Display: Bar Chart of Gene Count p-Value Cutoff (<=): 0.01 Highlight



Click tooltips are read as: File name:Category name (Accession): # genes, Percent of gene list against total # genes in the list, p-value
% of genes list in the category is calculated for each binning list as: # genes for this category/ # total genes in the list * 100.

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