

Read count deviation from samples of an equimolar multiplexed library

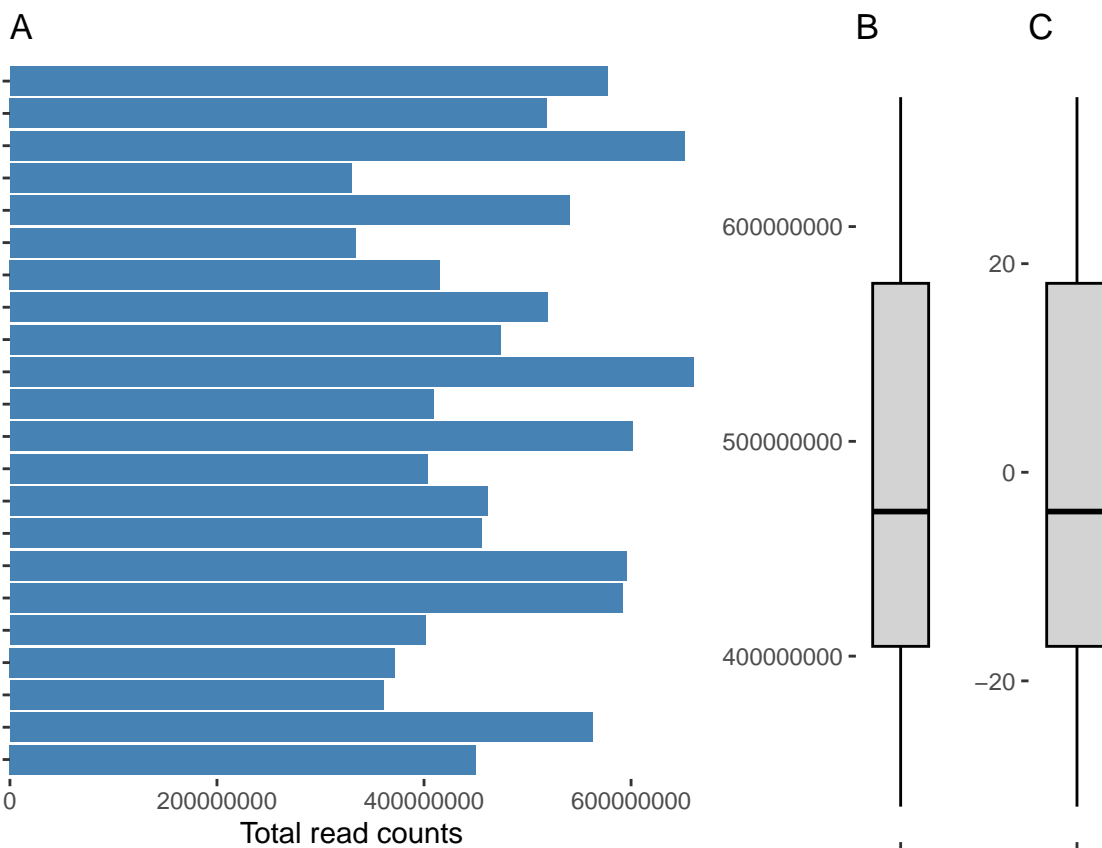
VIB Nucleomics Core

Thu Jul 13, 2023

Input file: 4490_230627_A01044_0335_AH2M7CDSX7-poolingQC-FinalTable.csv

Experiment: 4490

Centering on the mean



Legend:

A,B Total read counts

C: Boxplot of count difference from the mean

Sample count: 22

Total read count: 10683373649

Mean read count: 485607893

mean %Difference from mean (Absolute Value): 18.01 %