

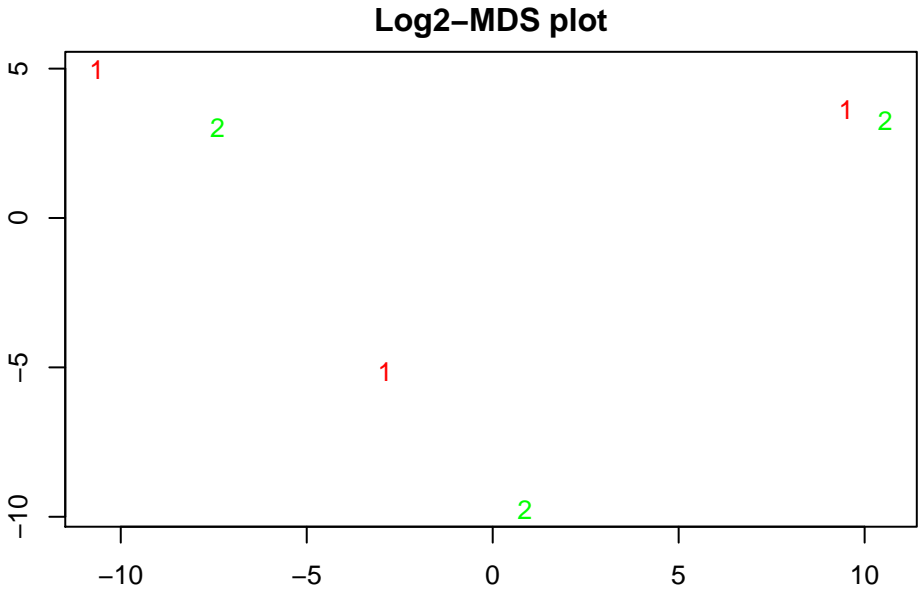
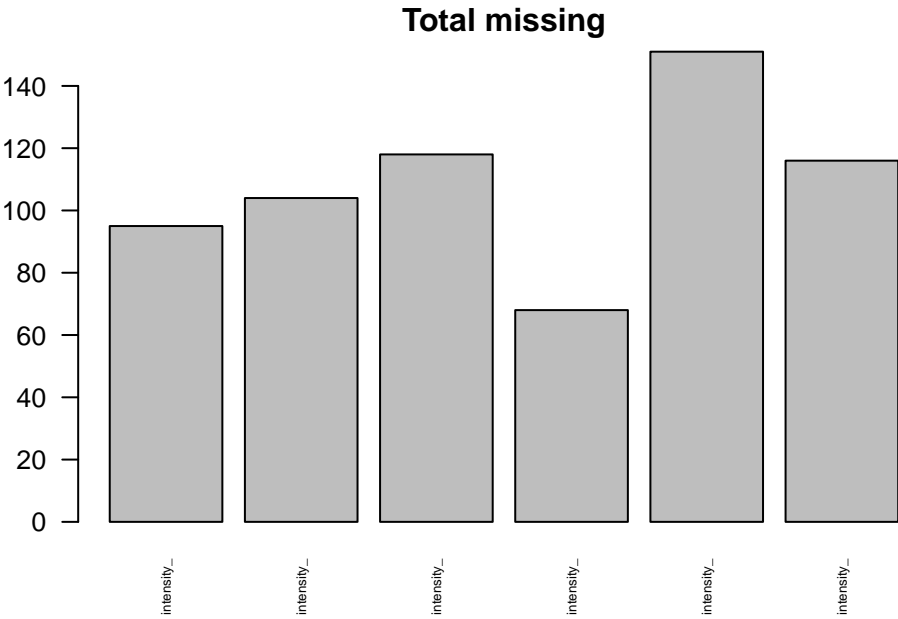
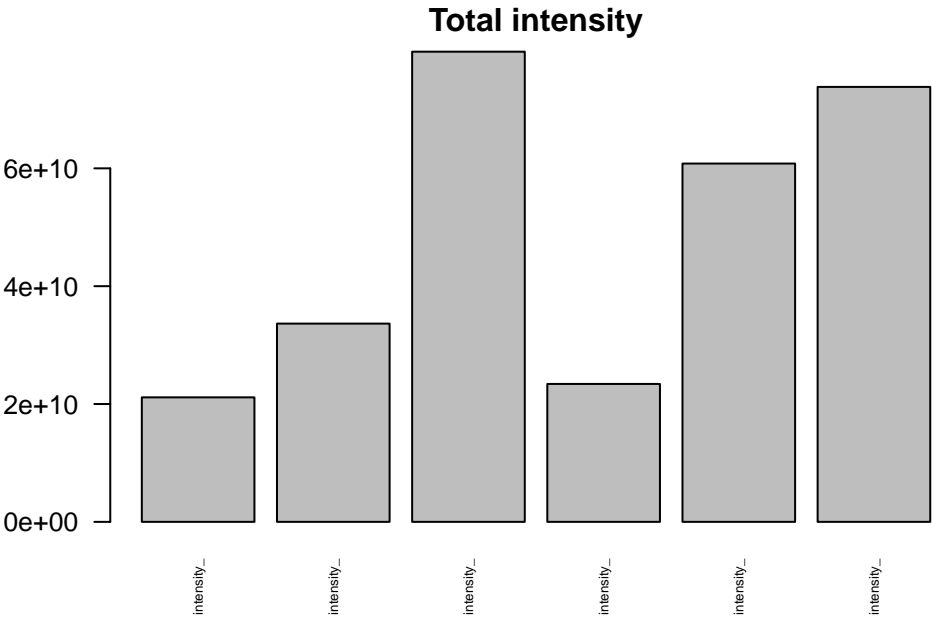
Project Name: junc.exons\_protein\_groups

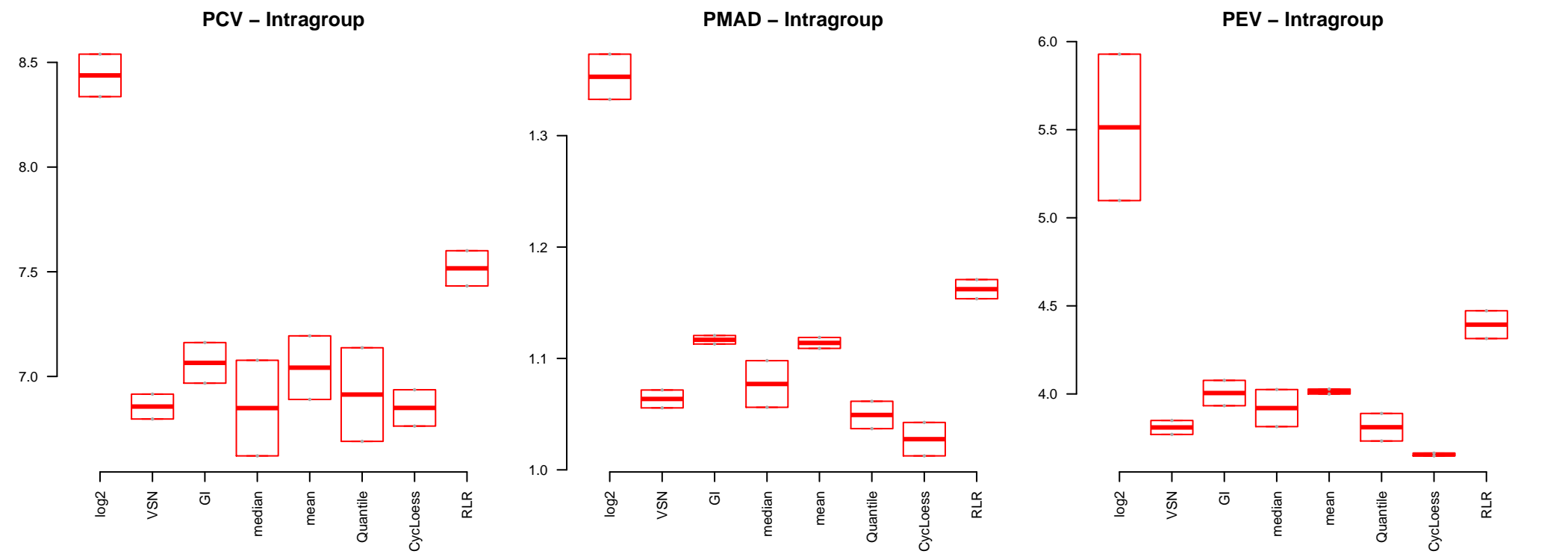
# Normalyzer (ver 1.4.0 )

Report created on: 2020-05-16

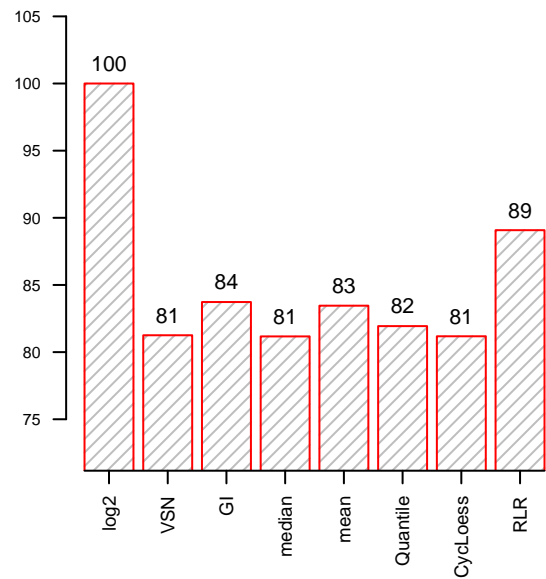
Citation: NormalyzerDE: Online Tool for Improved Normalization of Omics Expression Data and High-Sensitivity Differential Expression Analysis  
Journal of Proteome Research (2018), 10.1021/acs.jproteome.8b00523

Documentation for analyzing this report can be found at <http://quantitativeproteomics.org/normalyzer/help.php>

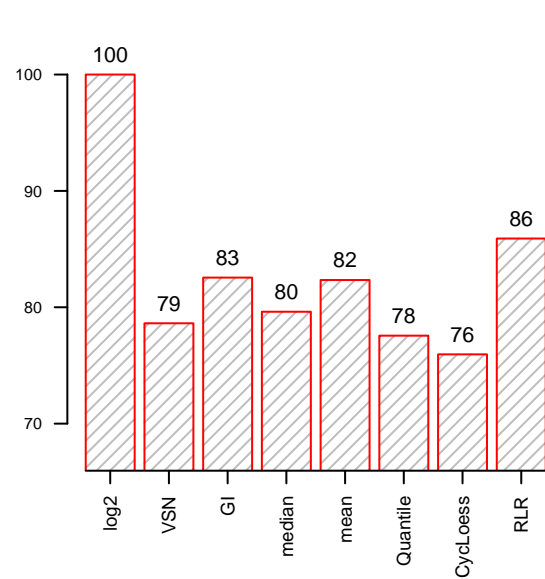




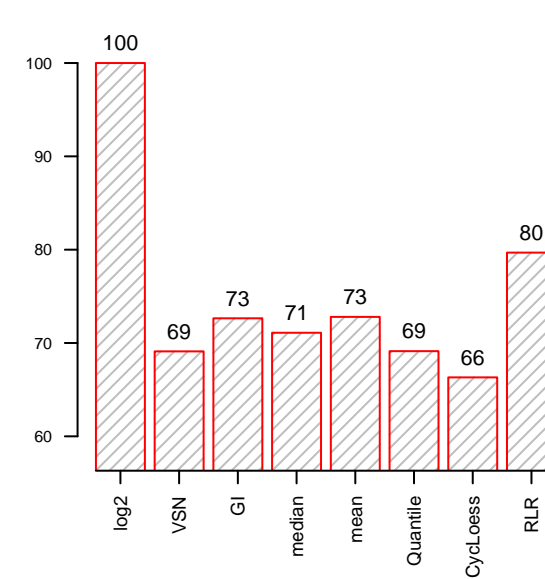
PCV compared to log2



PMAD compared to log2



%PEV – compared to log2



Stable variables plot

