

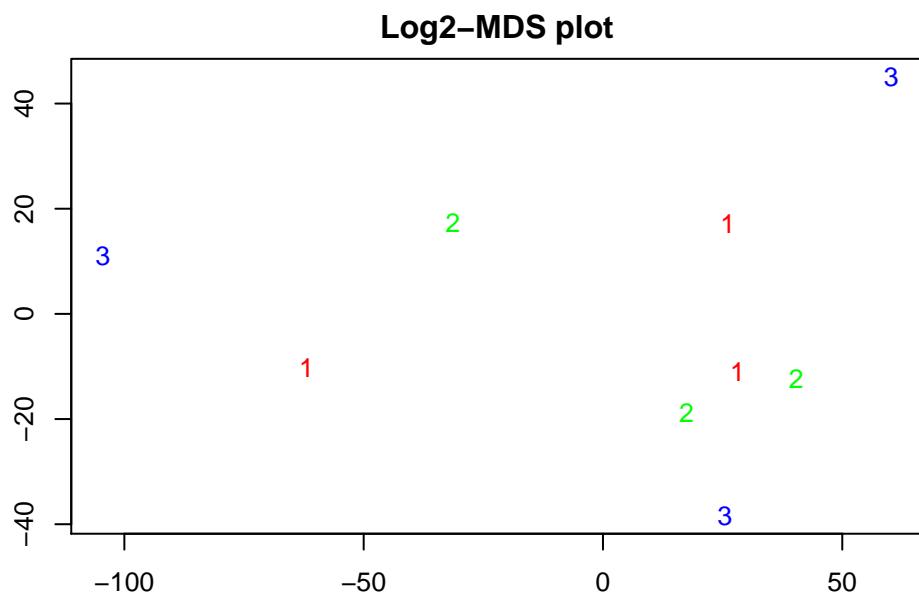
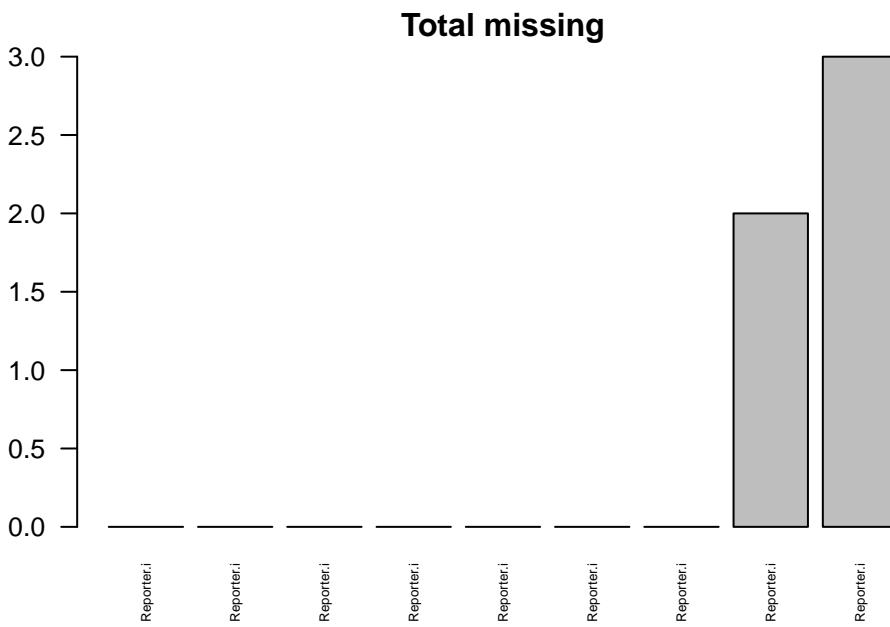
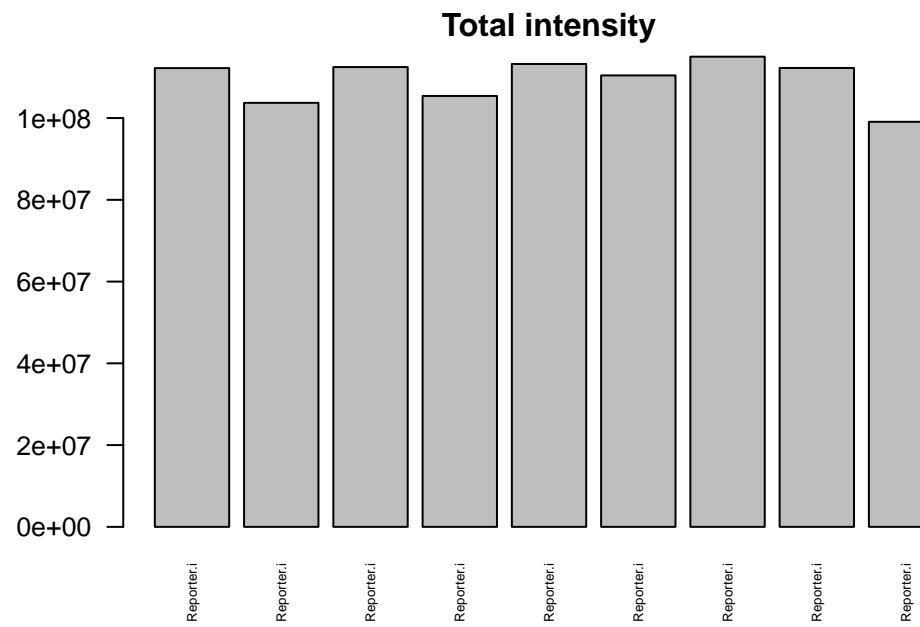
Project Name: sumExpRun

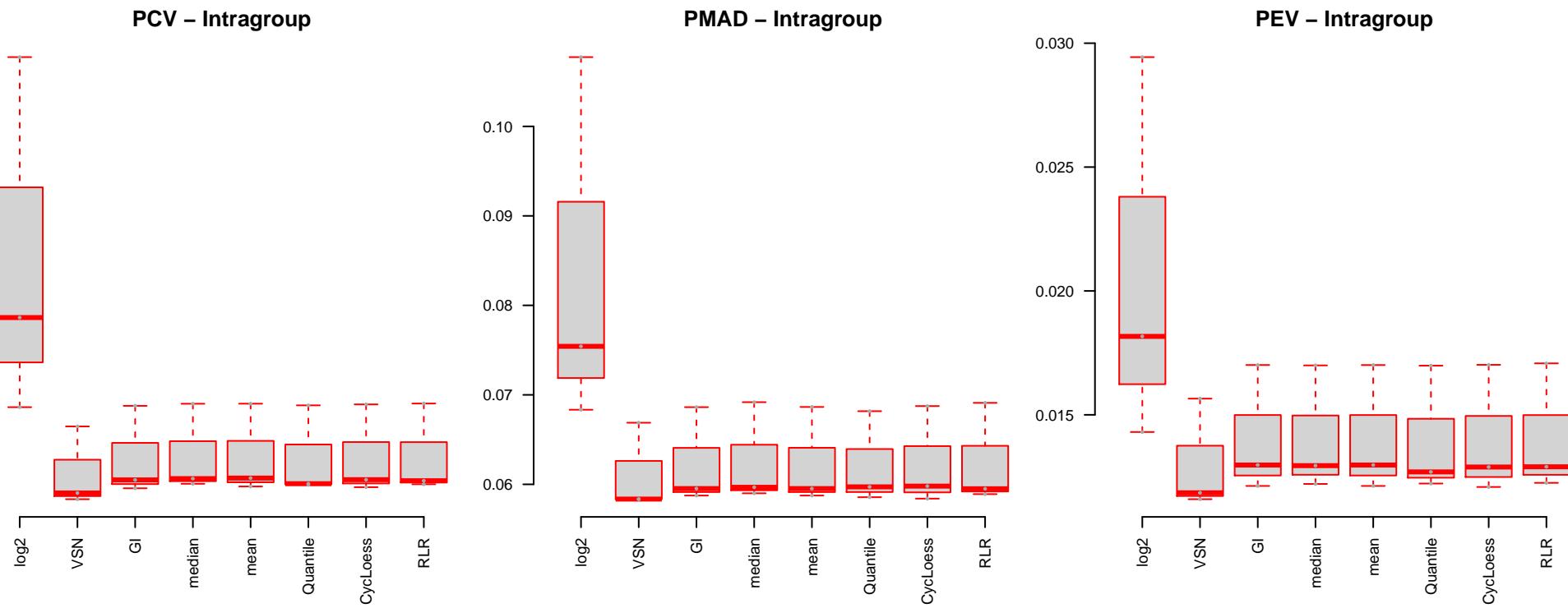
# Normalizer (ver 1.8.0 )

Report created on: 2022-04-06

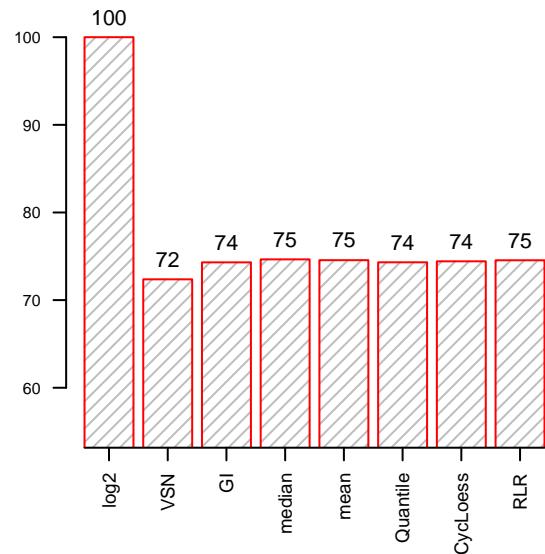
Citation: NormalizerDE: Online Tool for Improved Normalization of Omics Expression Data and High-Sensitivity Differential Expression Analysis  
Journal of Proteome Research (2018), 10.1021/acs.jproteome.8b00523

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

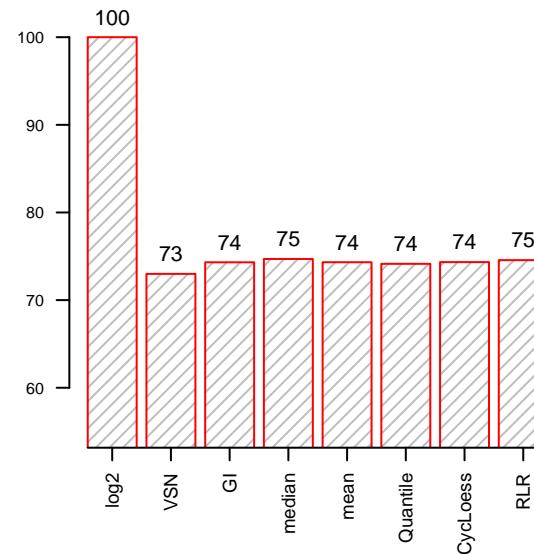




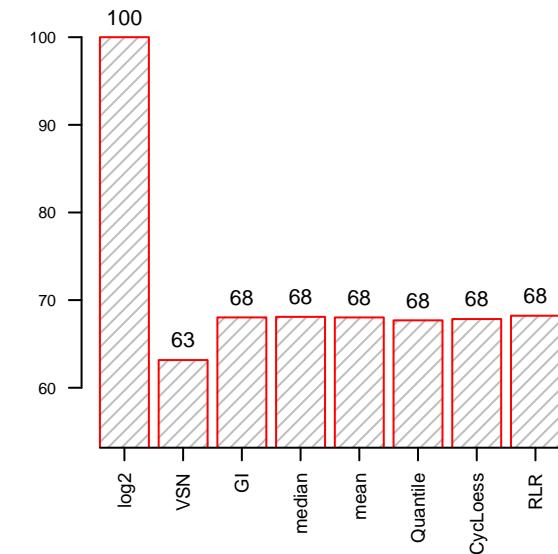
PCV compared to log2



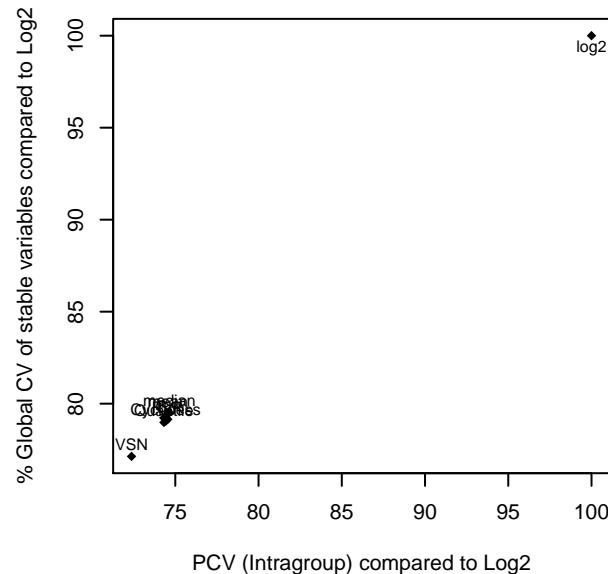
PMAD compared to log2

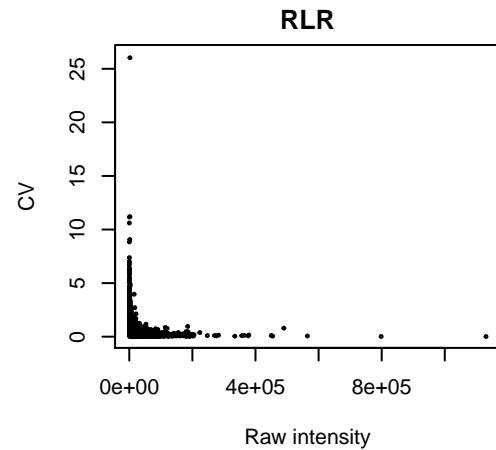
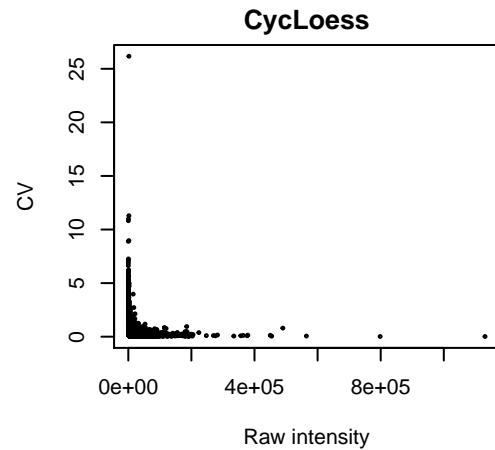
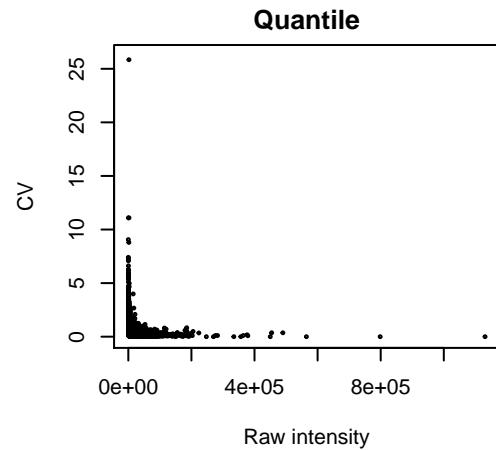
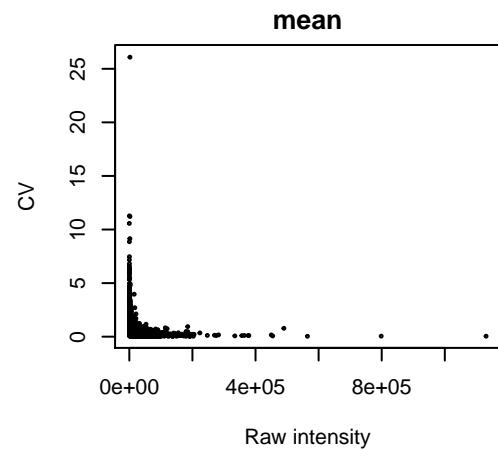
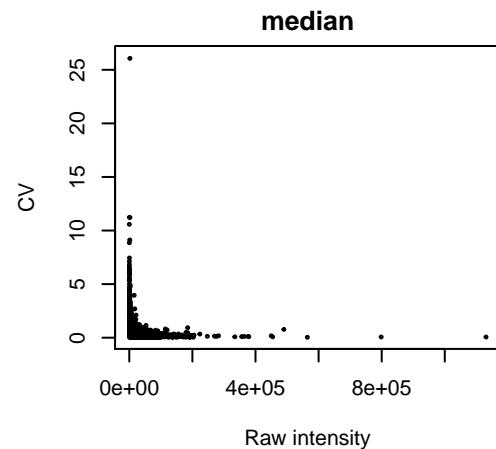
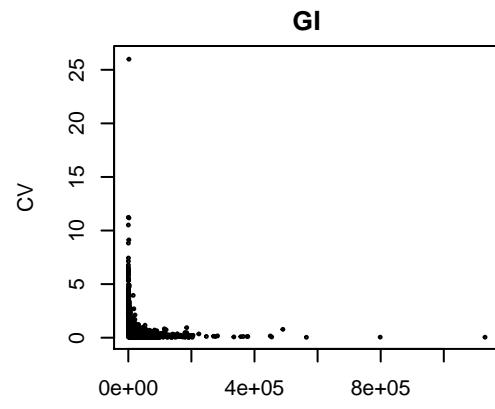
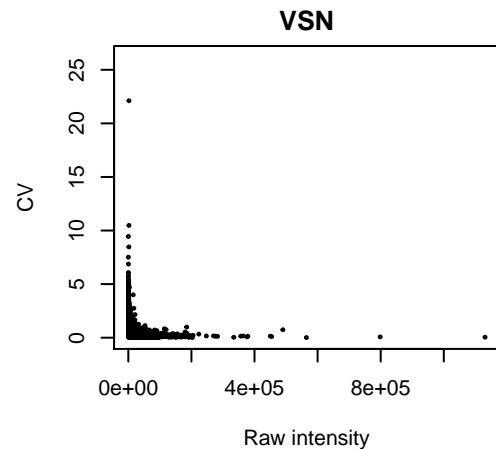
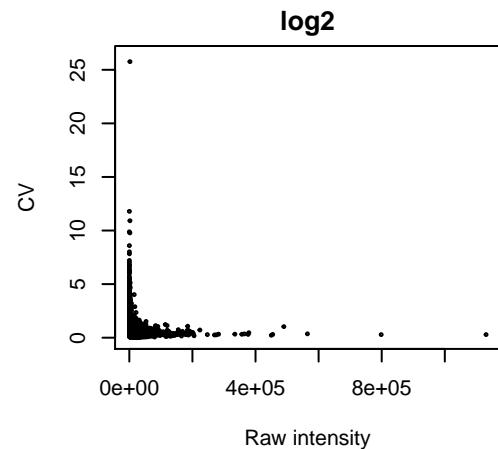


%PEV – compared to log2



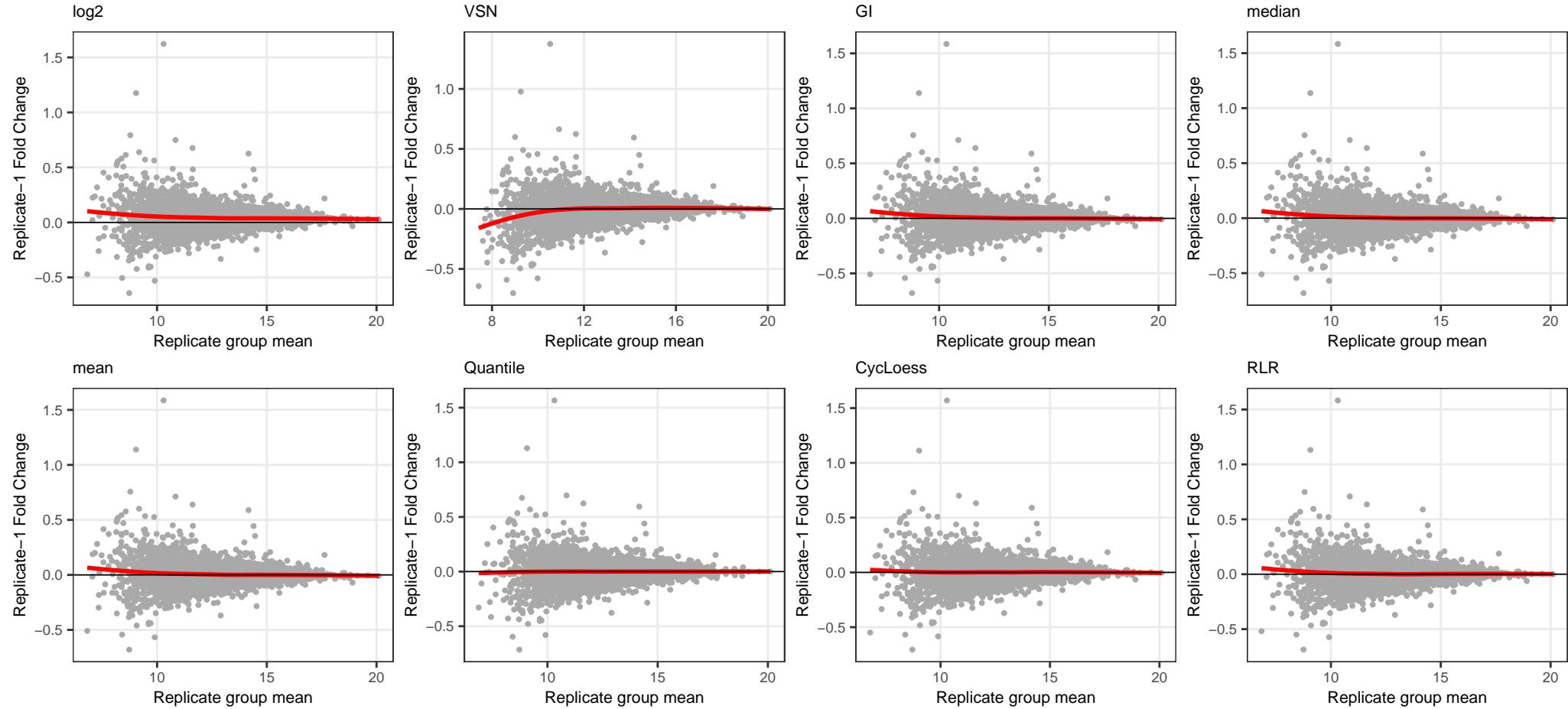
Stable variables plot

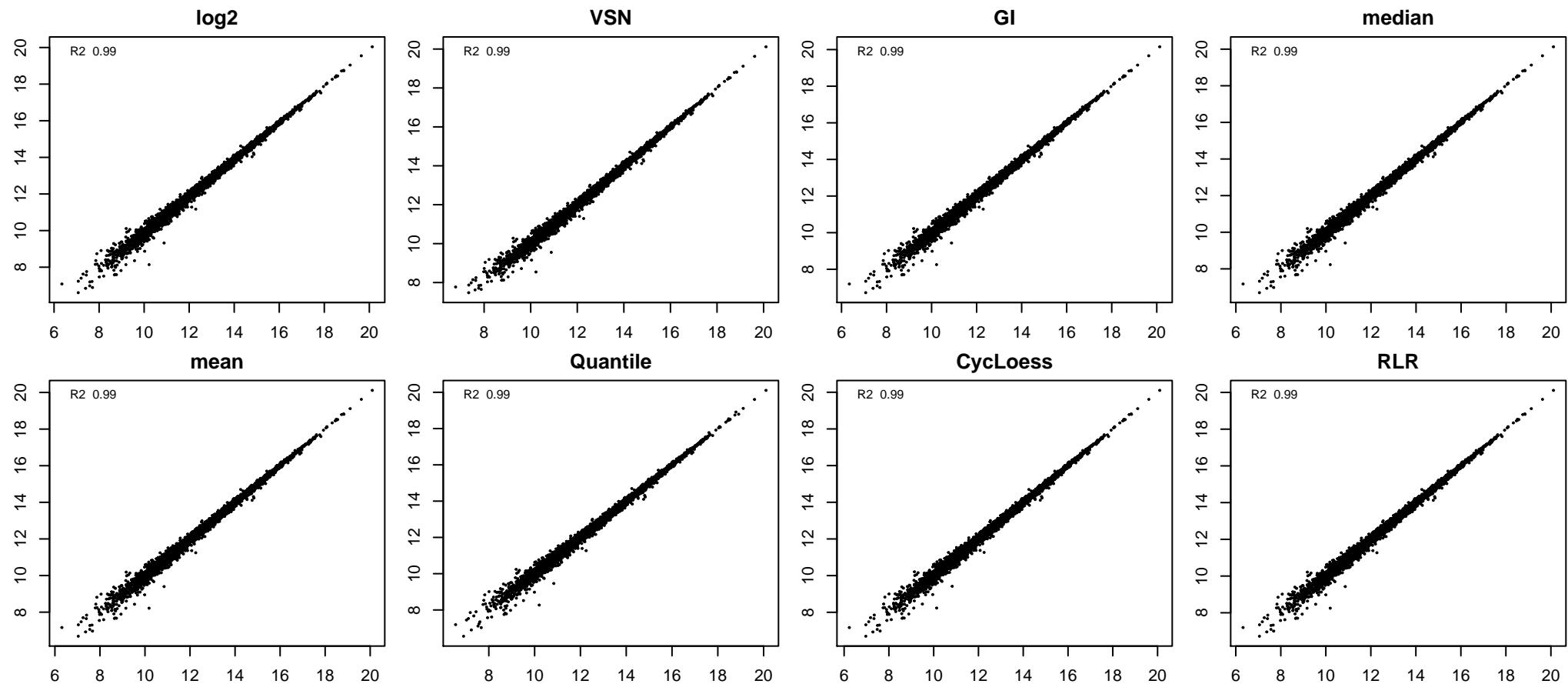




## MA plots

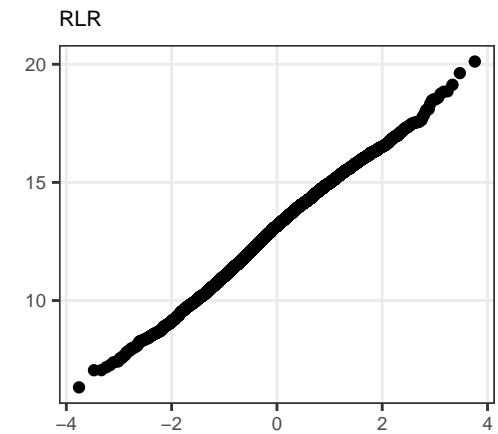
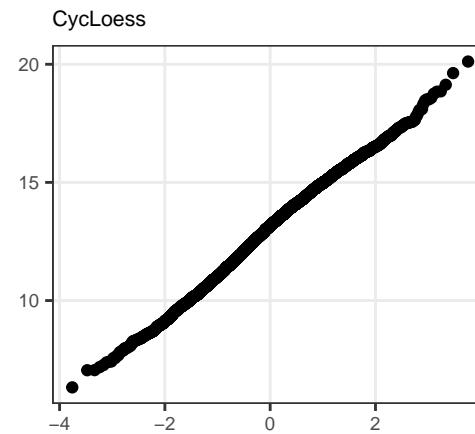
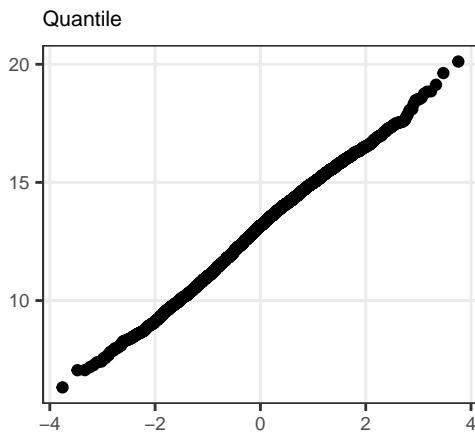
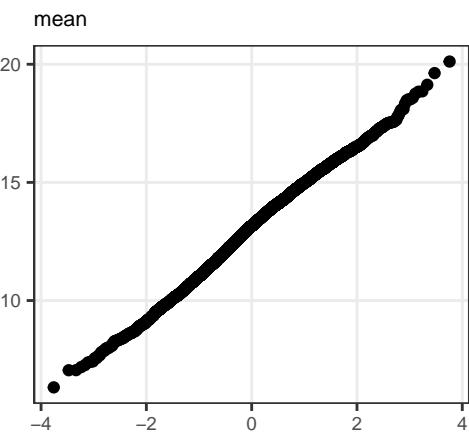
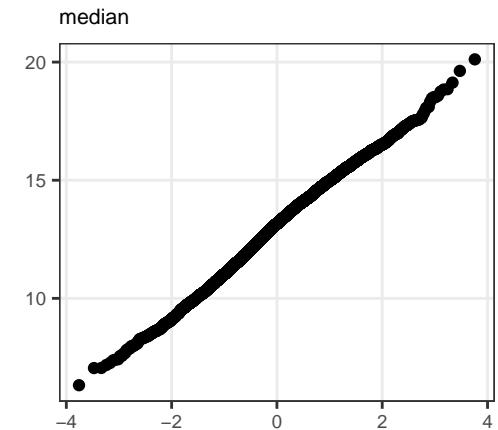
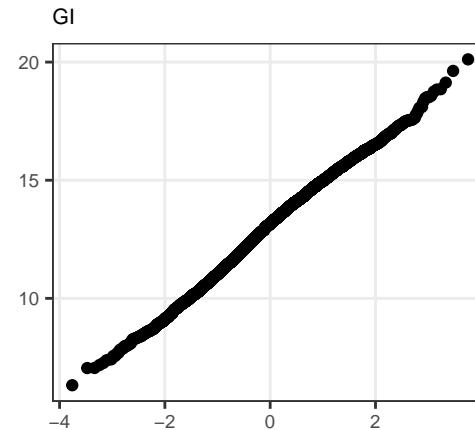
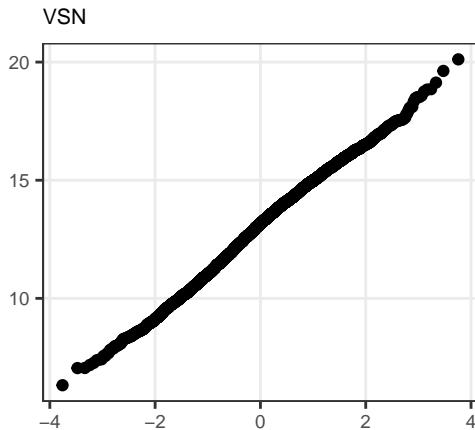
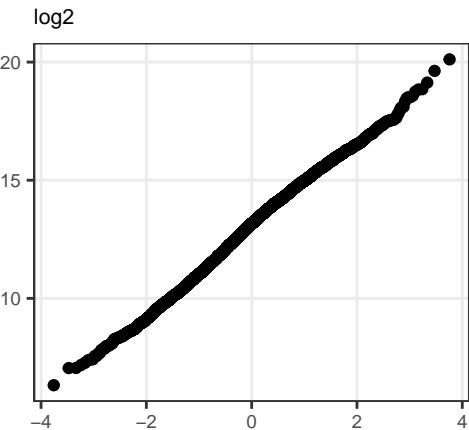
## NormalizerDE Report





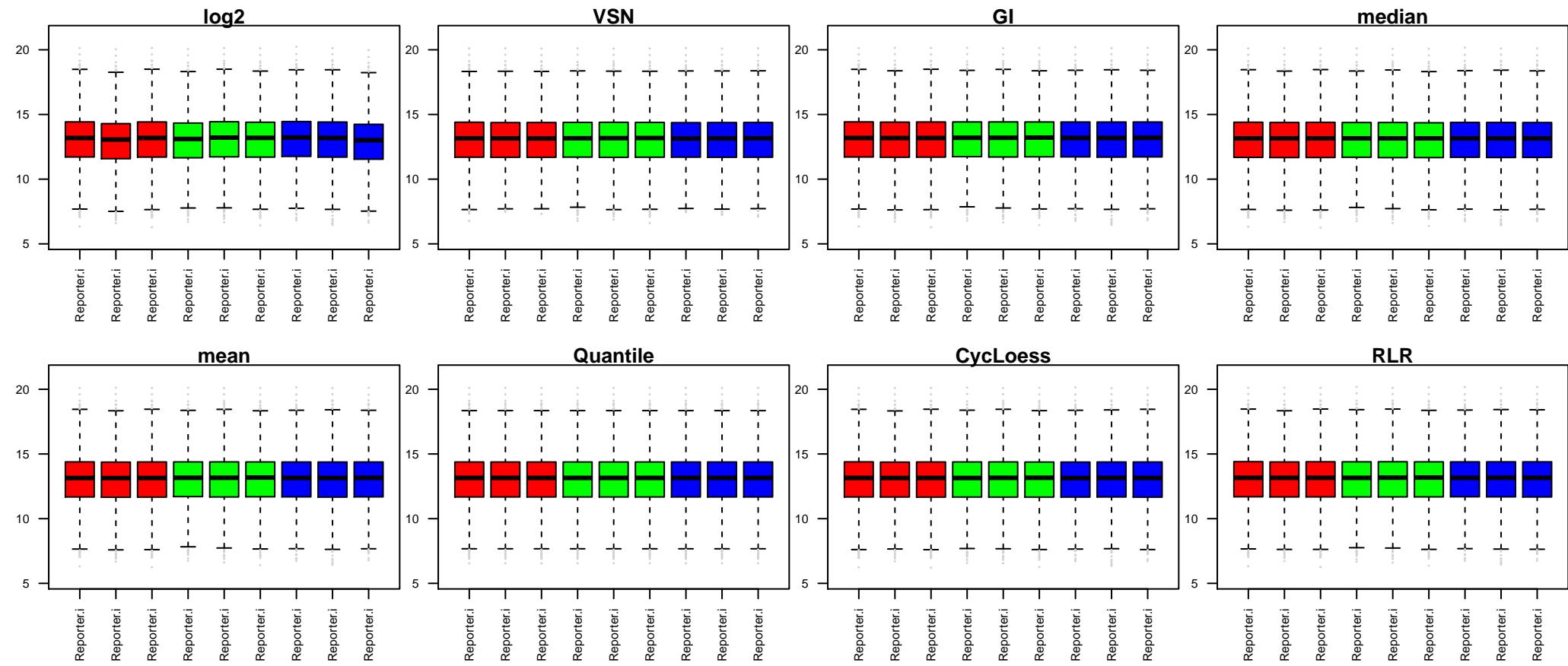
## Q-Q plots

## NormalizerDE Report



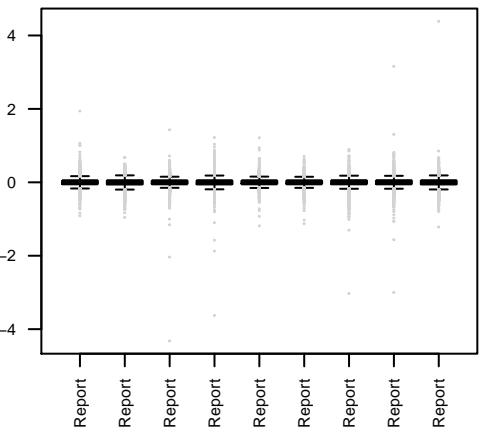
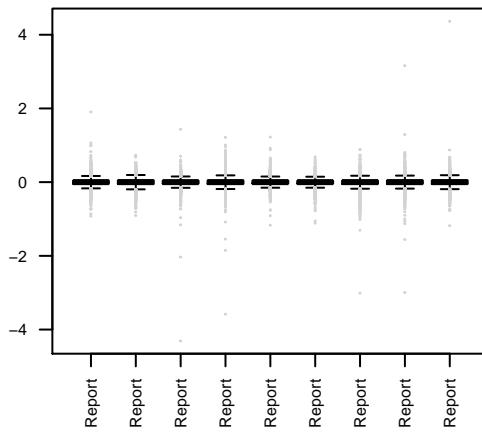
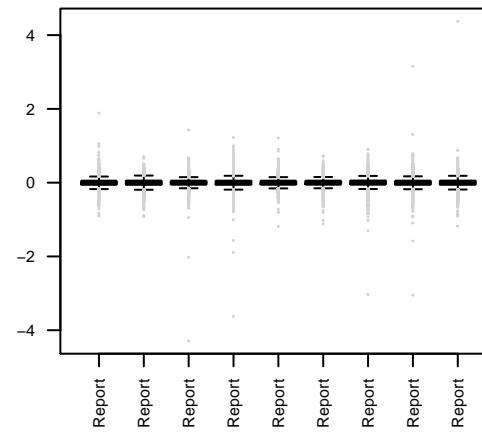
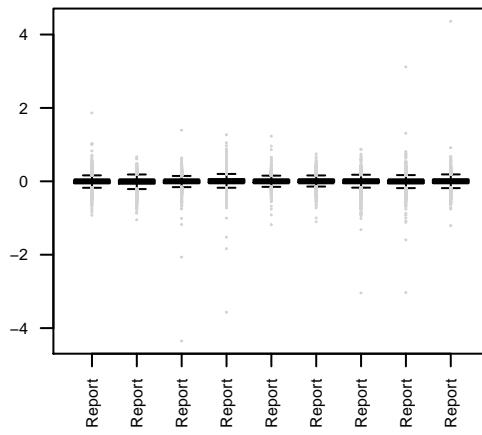
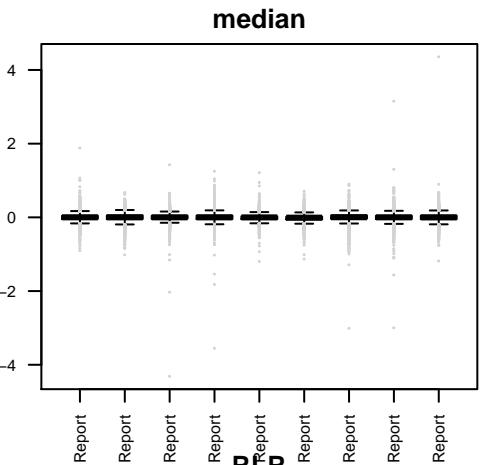
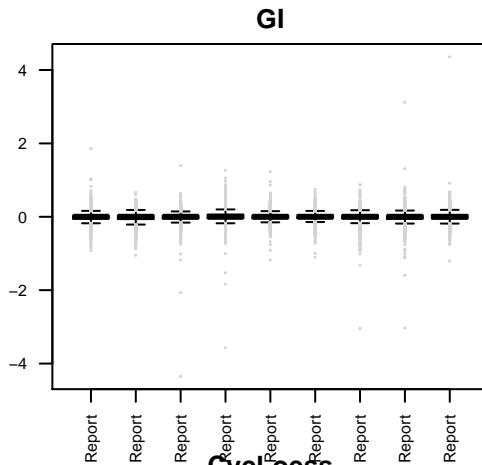
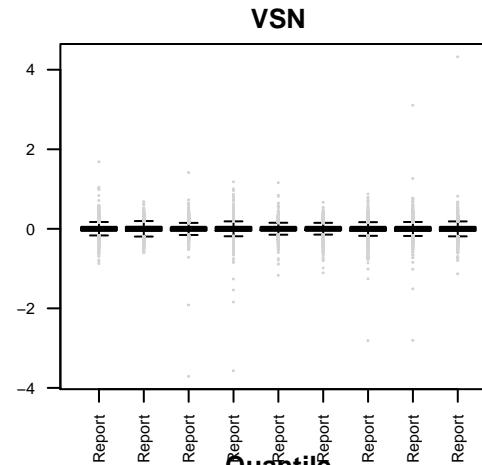
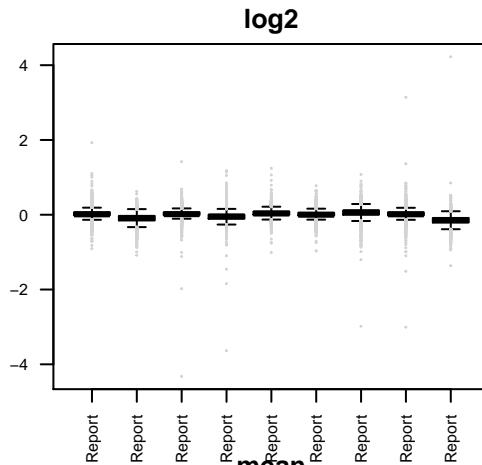
# Boxplots

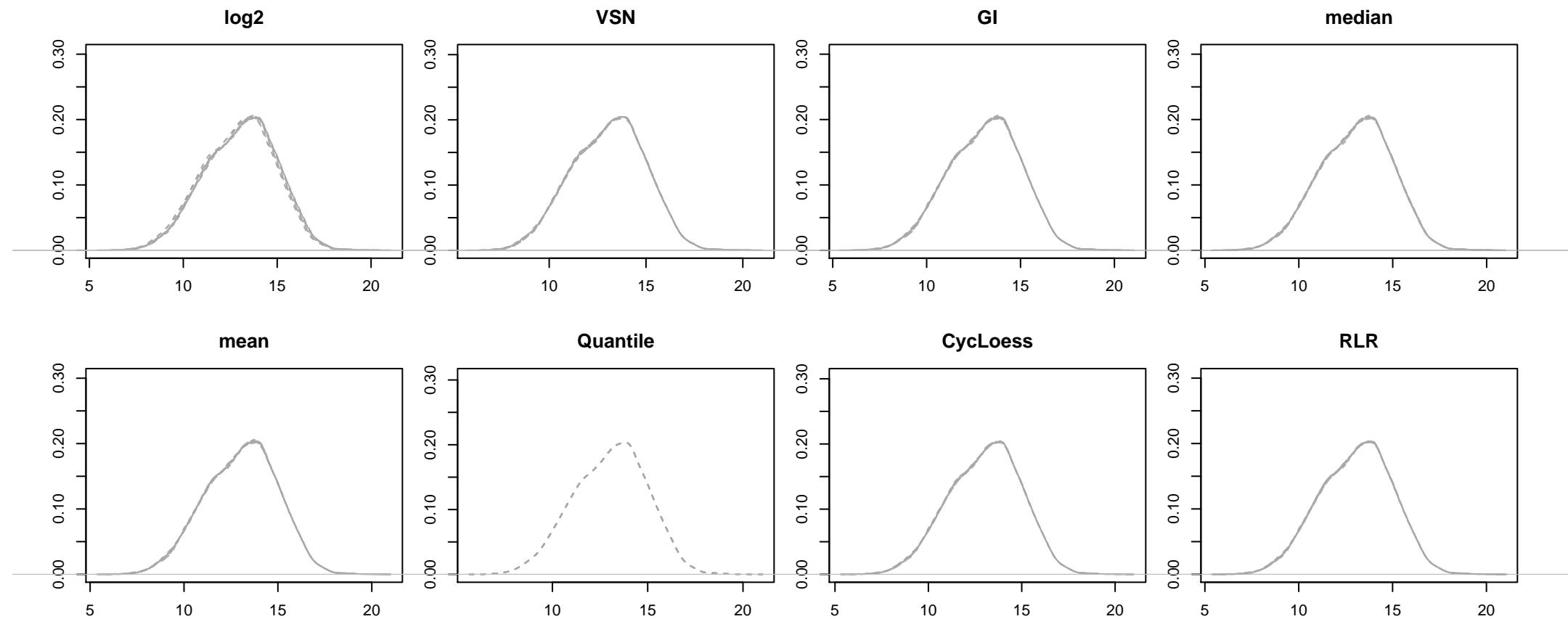
# Normalizer Report

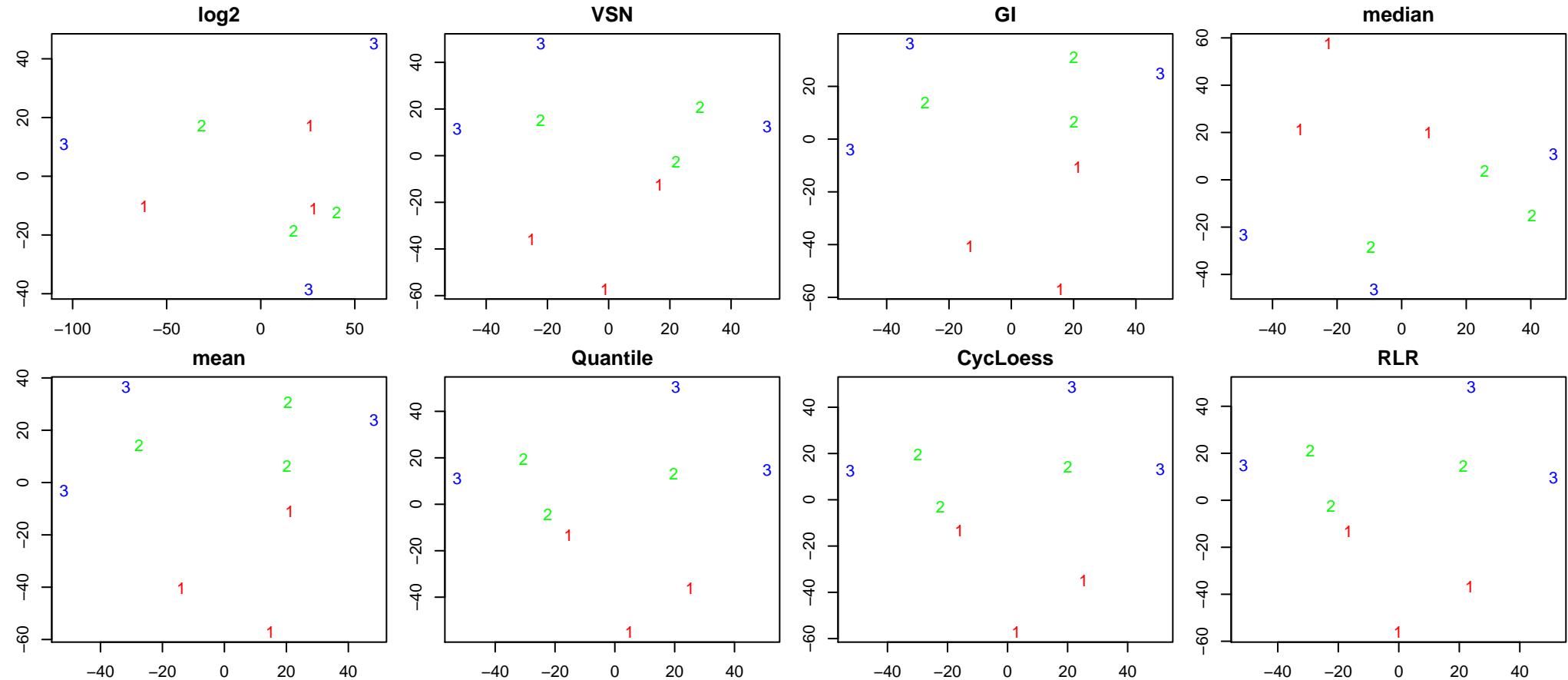


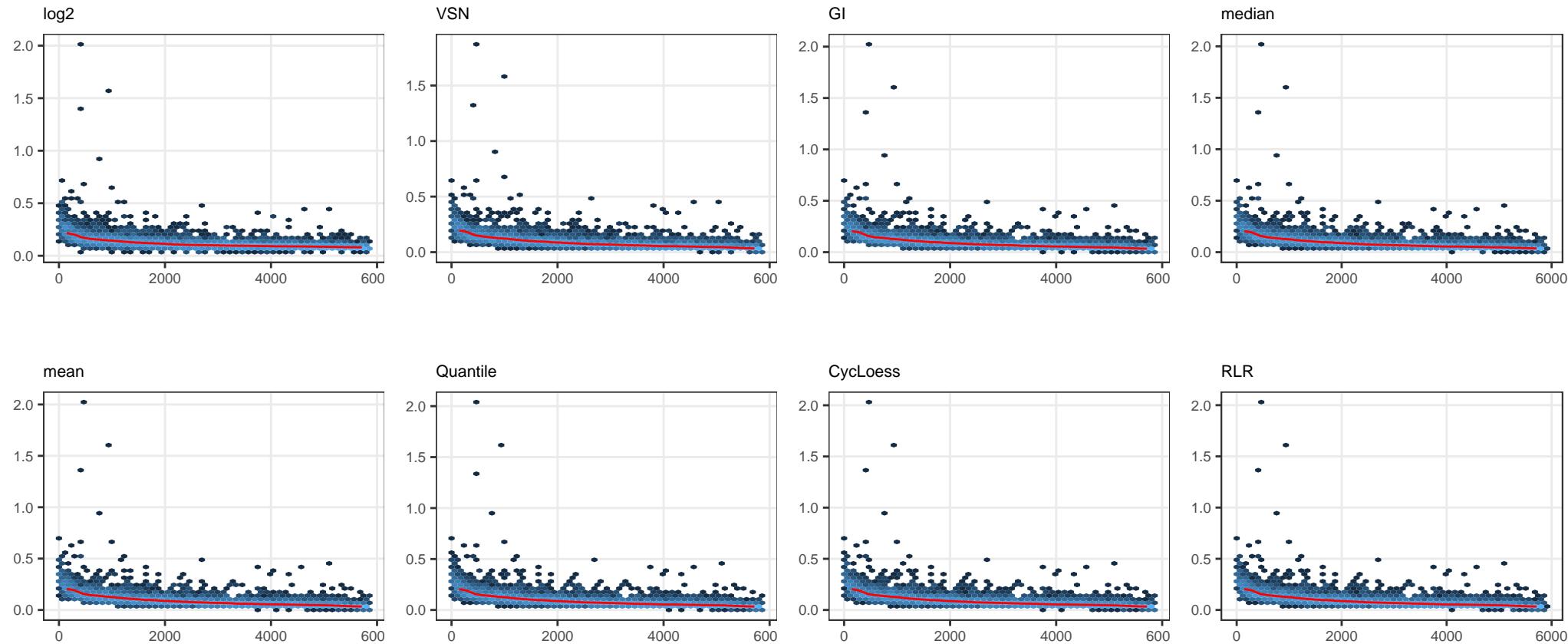
## Relative Log Expression (RLE) plots

Normalizer Report

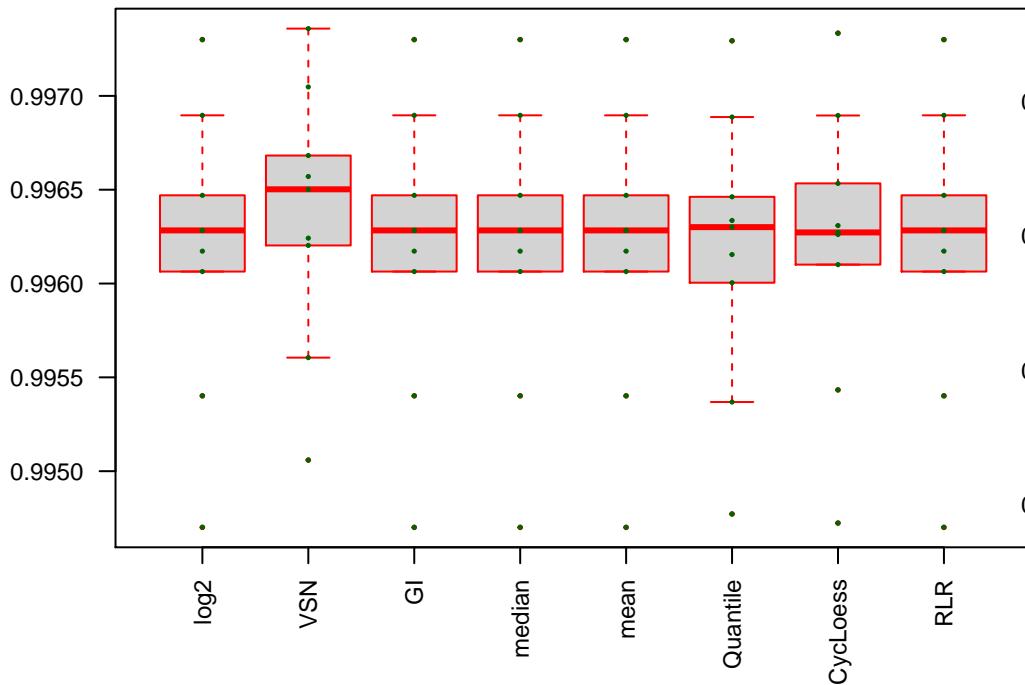




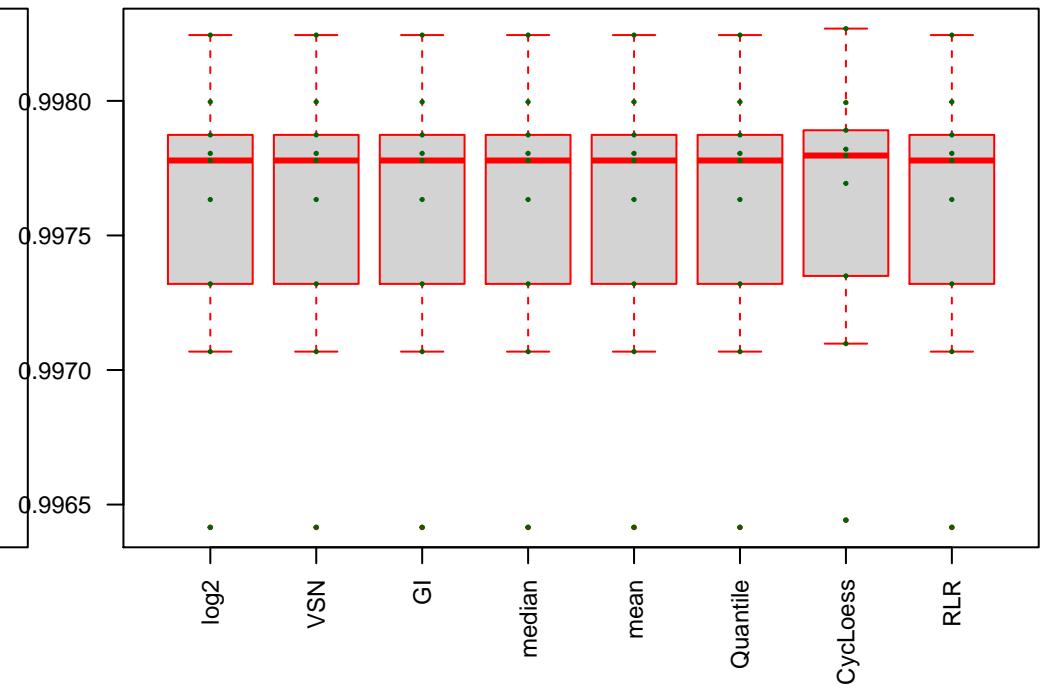


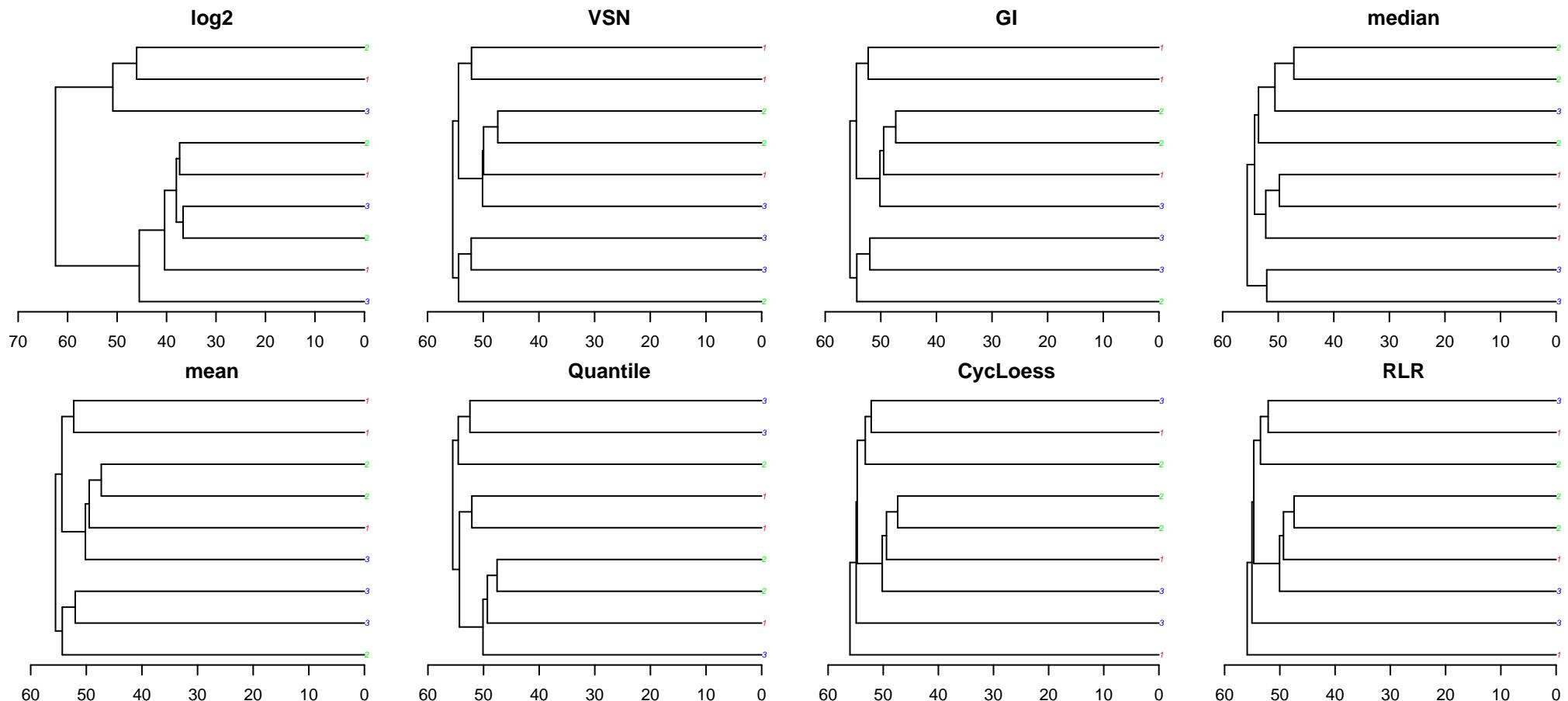


Pearson correlation – Intragroup



Spearman correlation – Intragroup





## HistPlots

## NormalizerDE Report

