

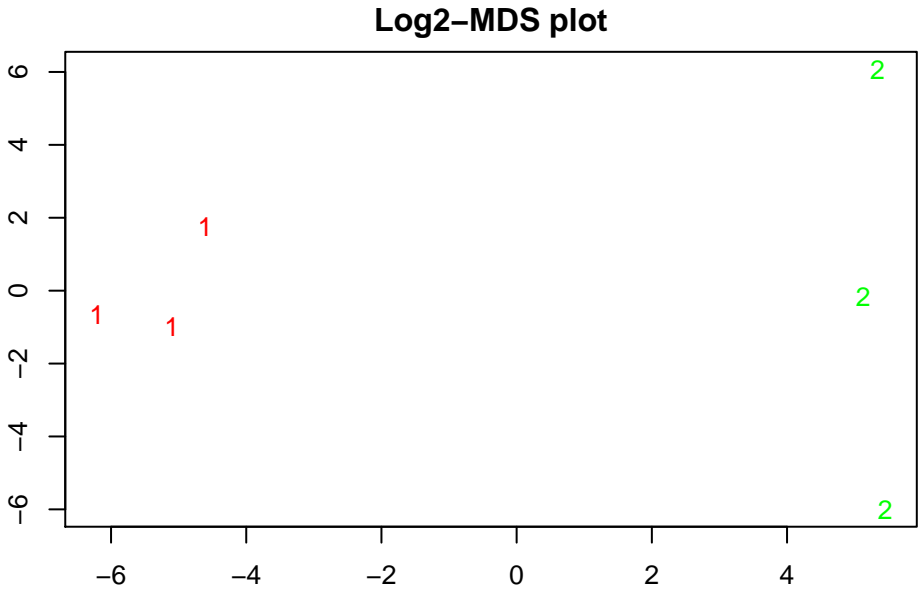
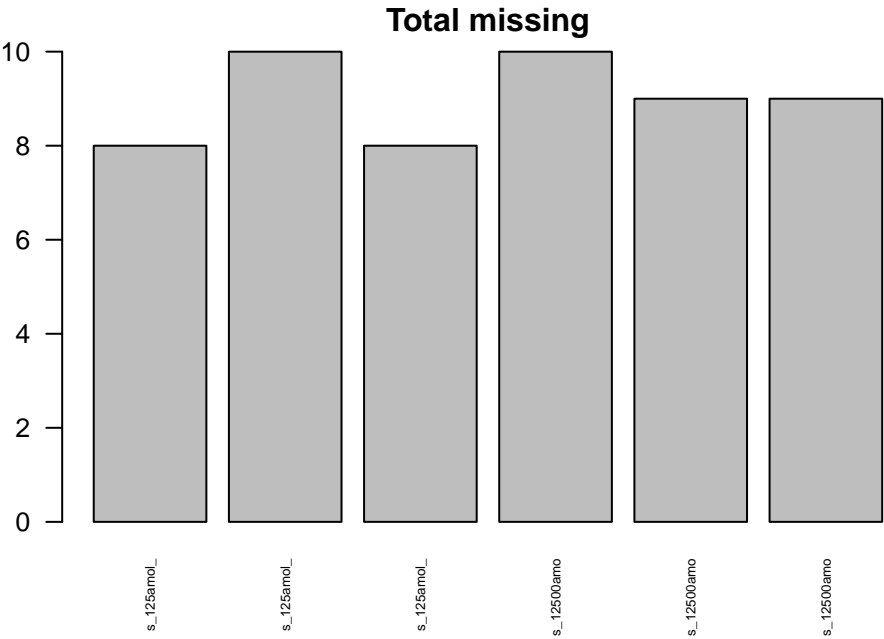
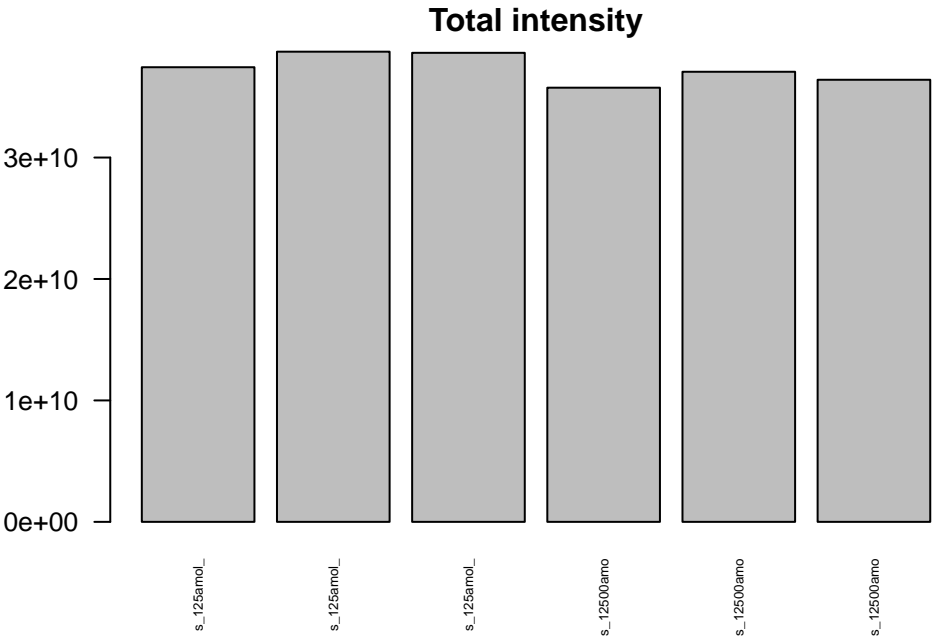
Project Name: unit\_test\_run\_norm

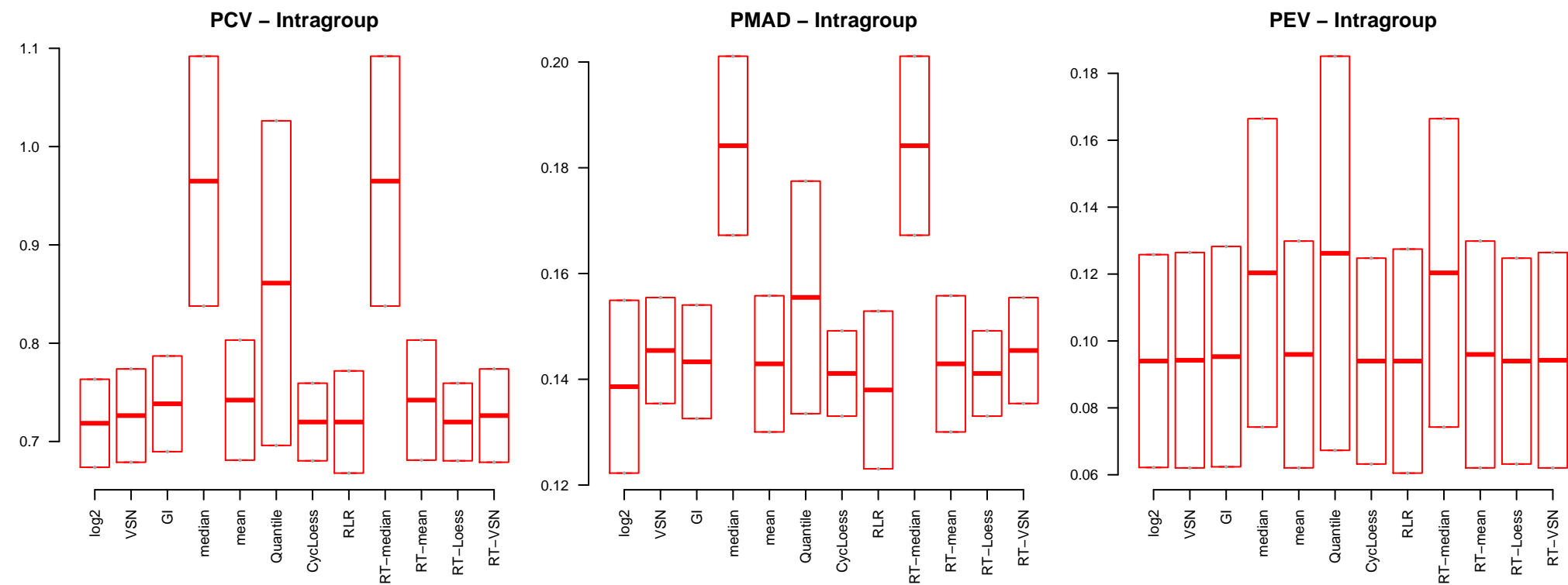
# Normalyzer (ver (version not found) )

Report created on: 2018-12-06

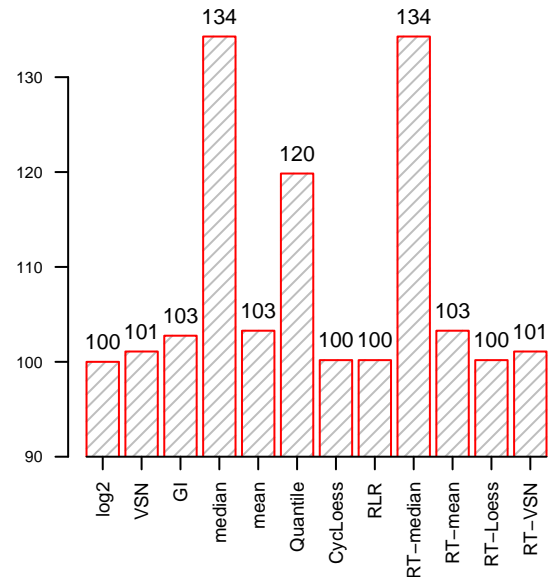
Citation: NormalyzerDE: Online Tool for Improved Normalization of Omics Expression Data and High-Sensitivity Differential Expression Analysis  
Journal of Proteome Research (2018), 10.1021/acs.jproteome.8b00523

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

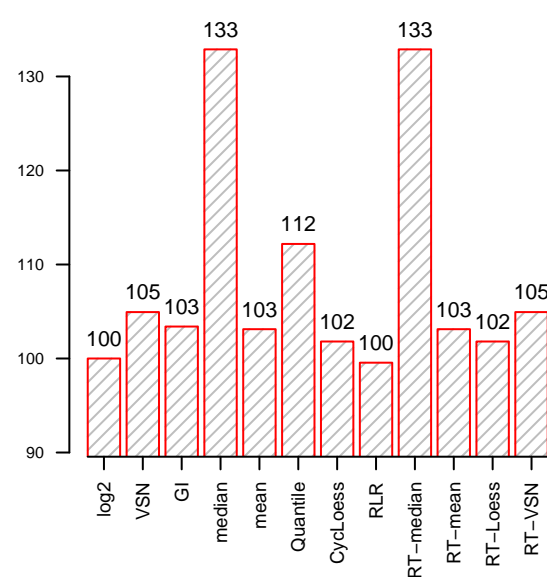




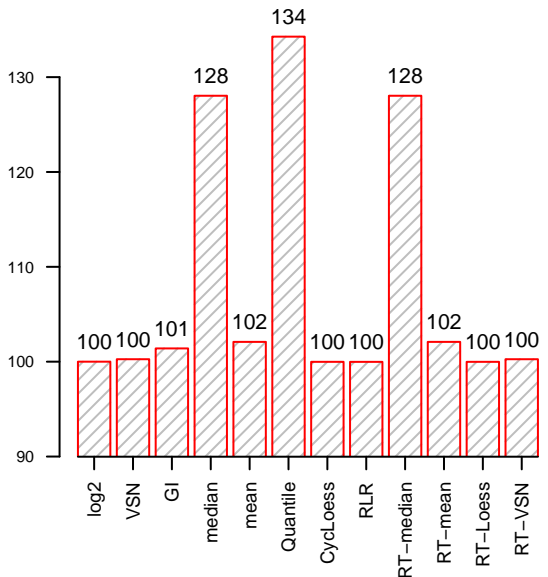
PCV compared to log2



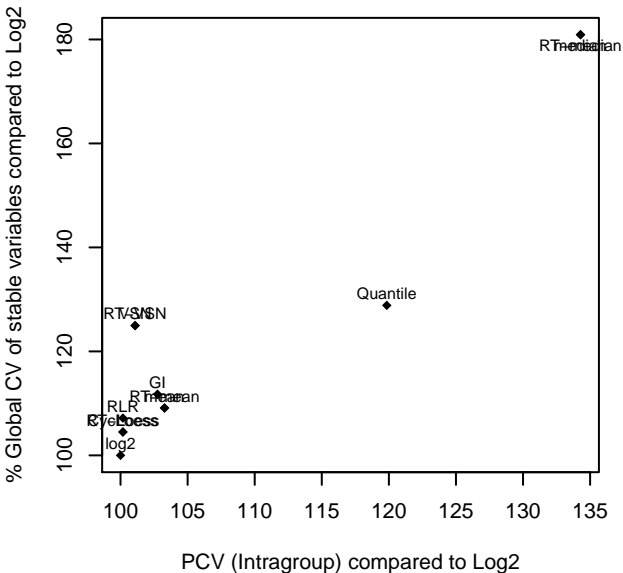
PMAD compared to log2

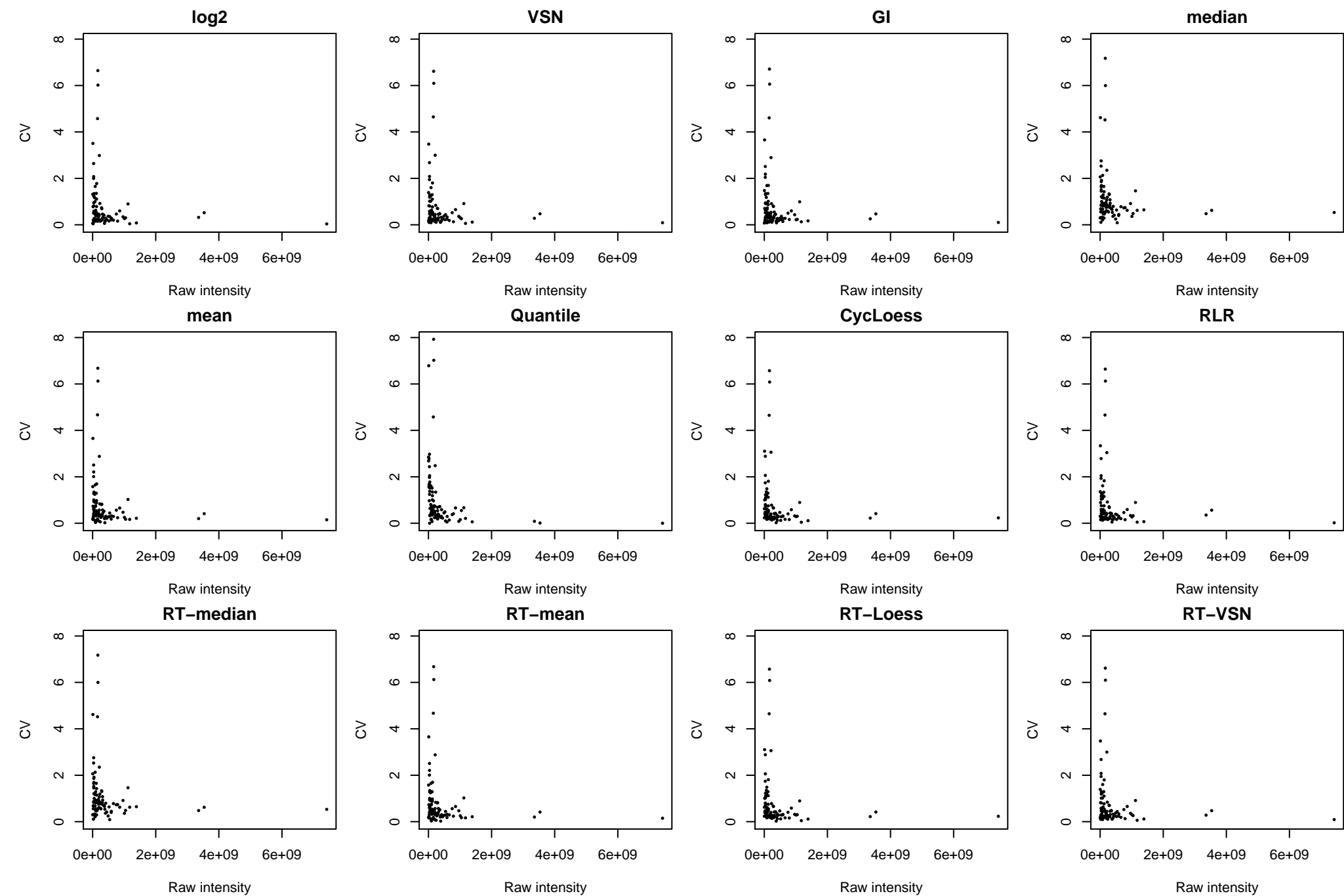


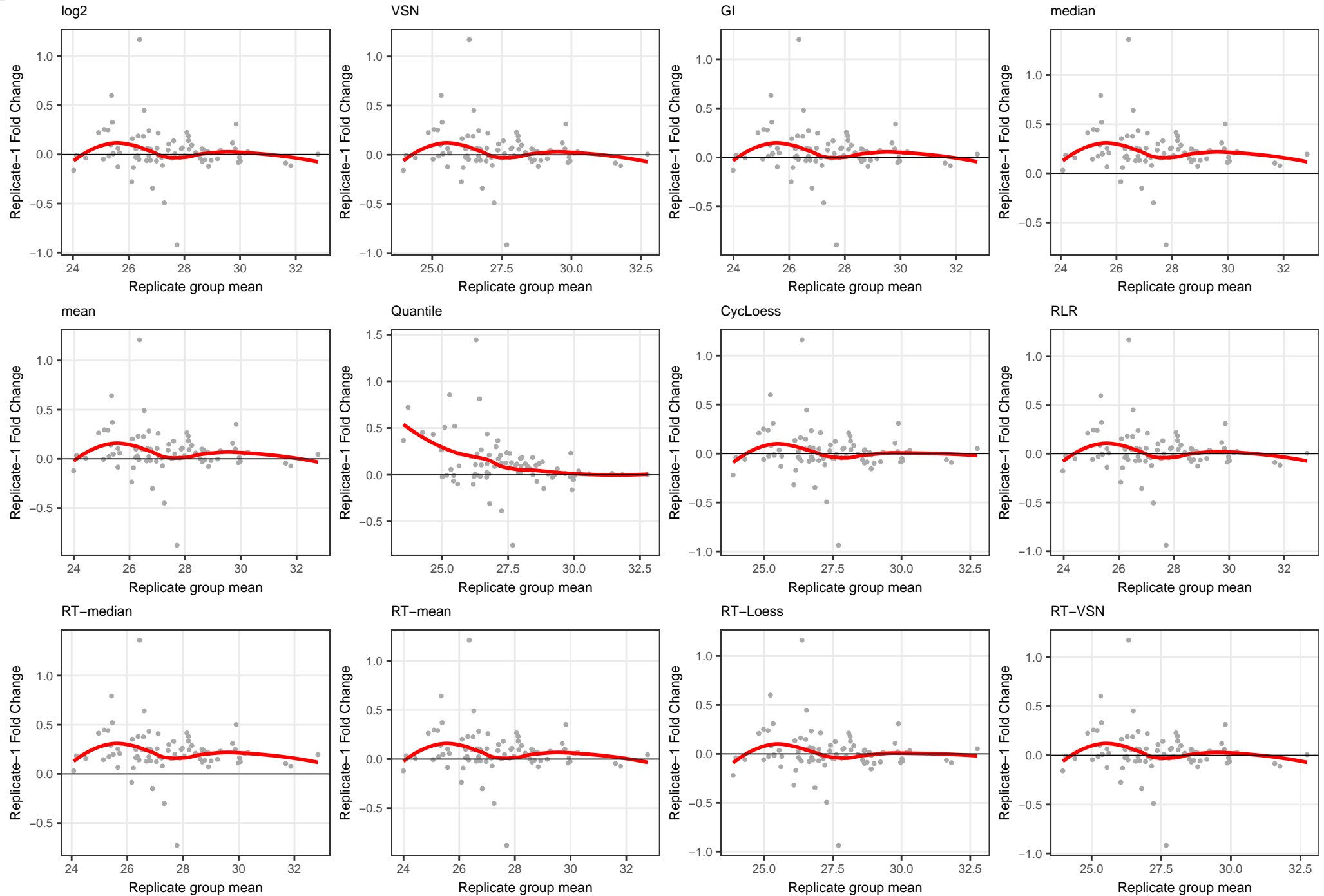
%PEV – compared to log2

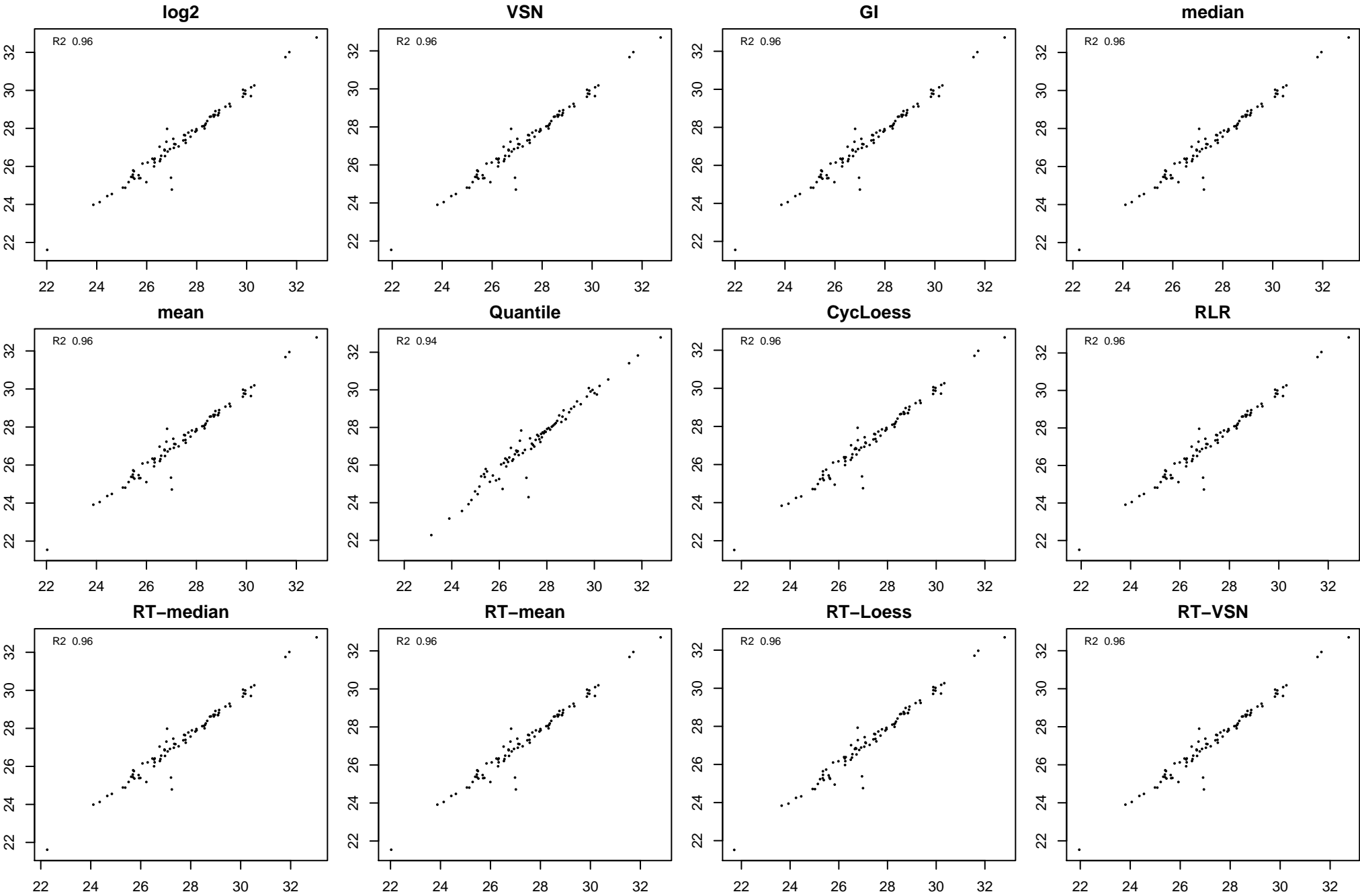


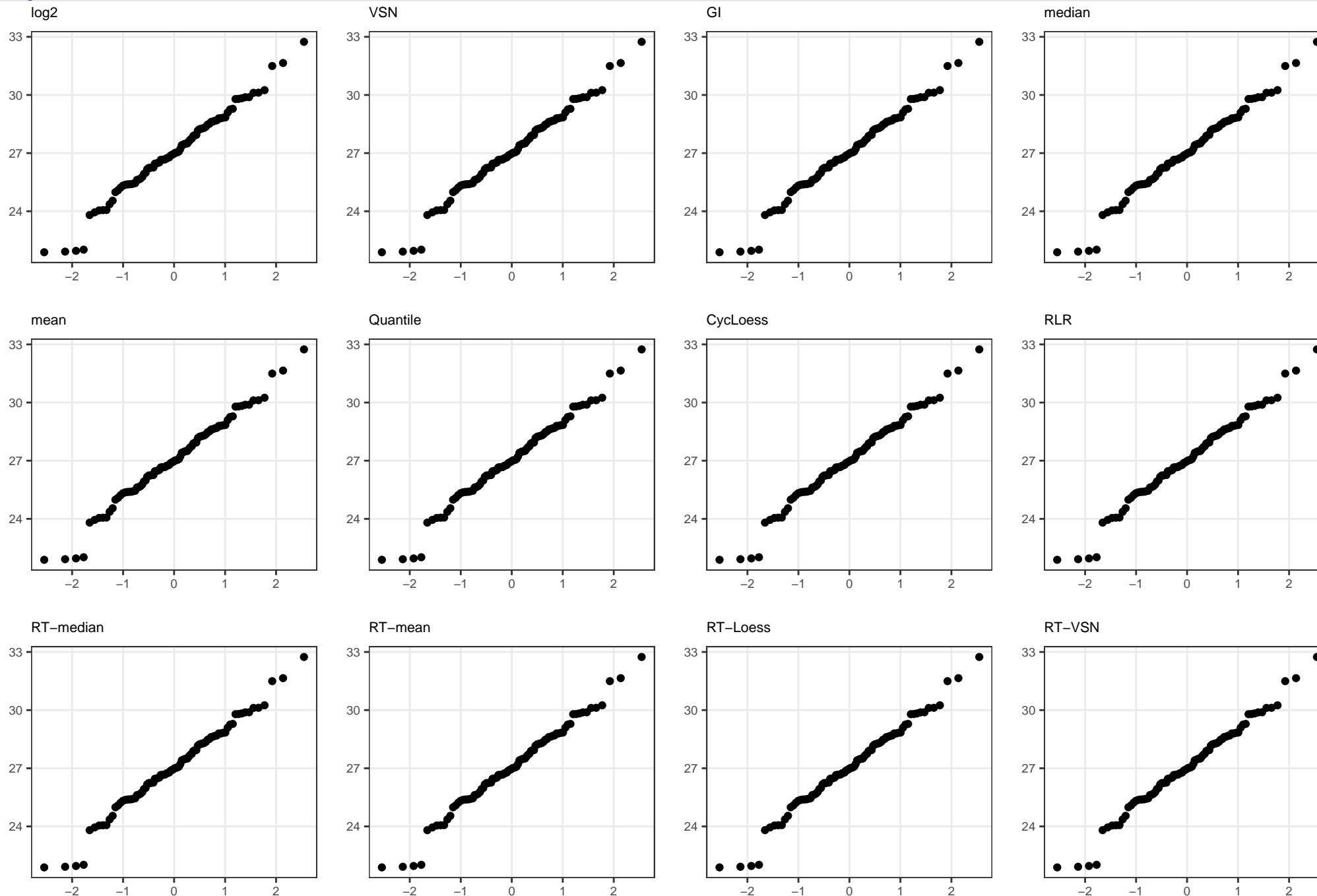
Stable variables plot



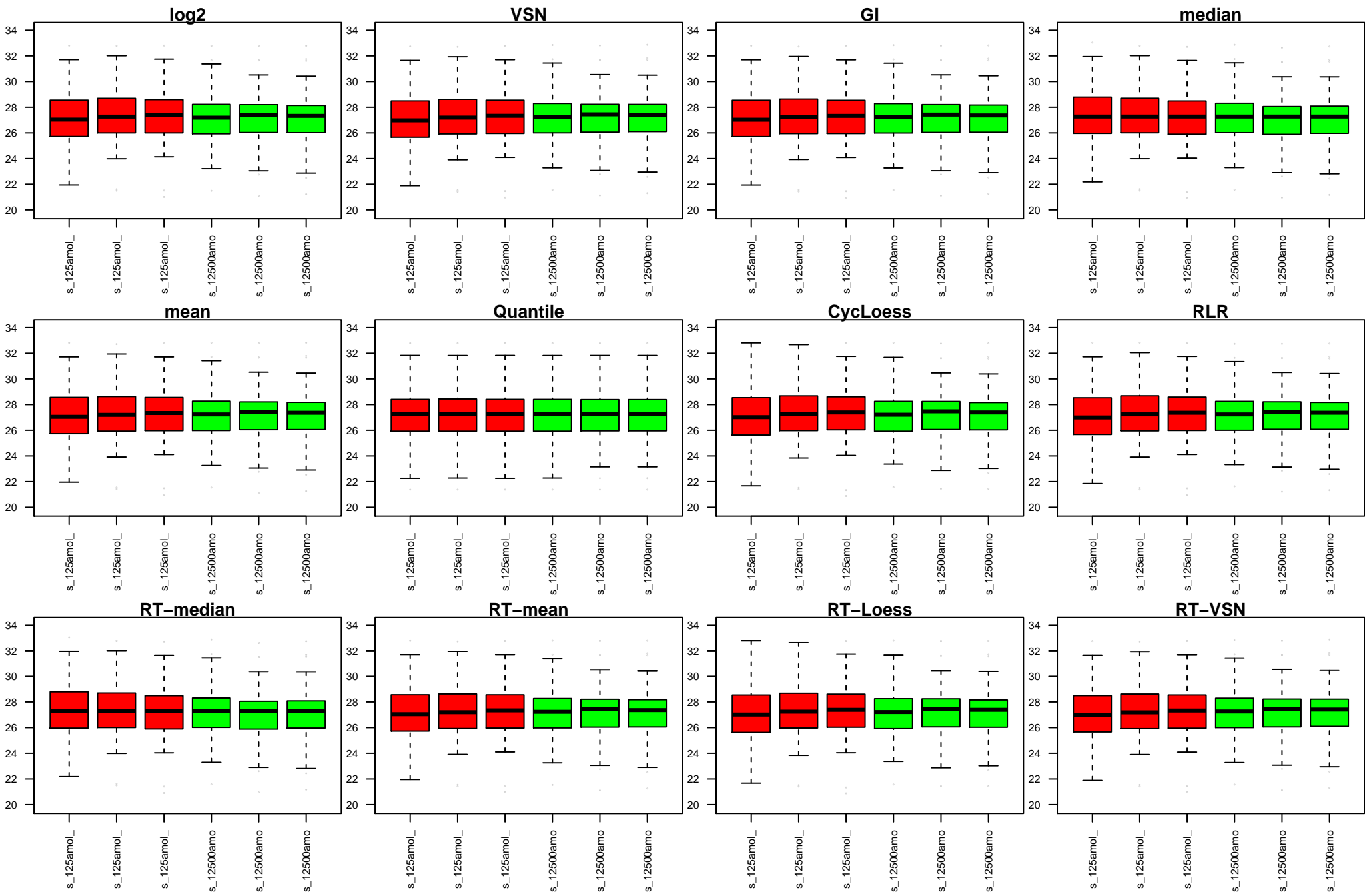


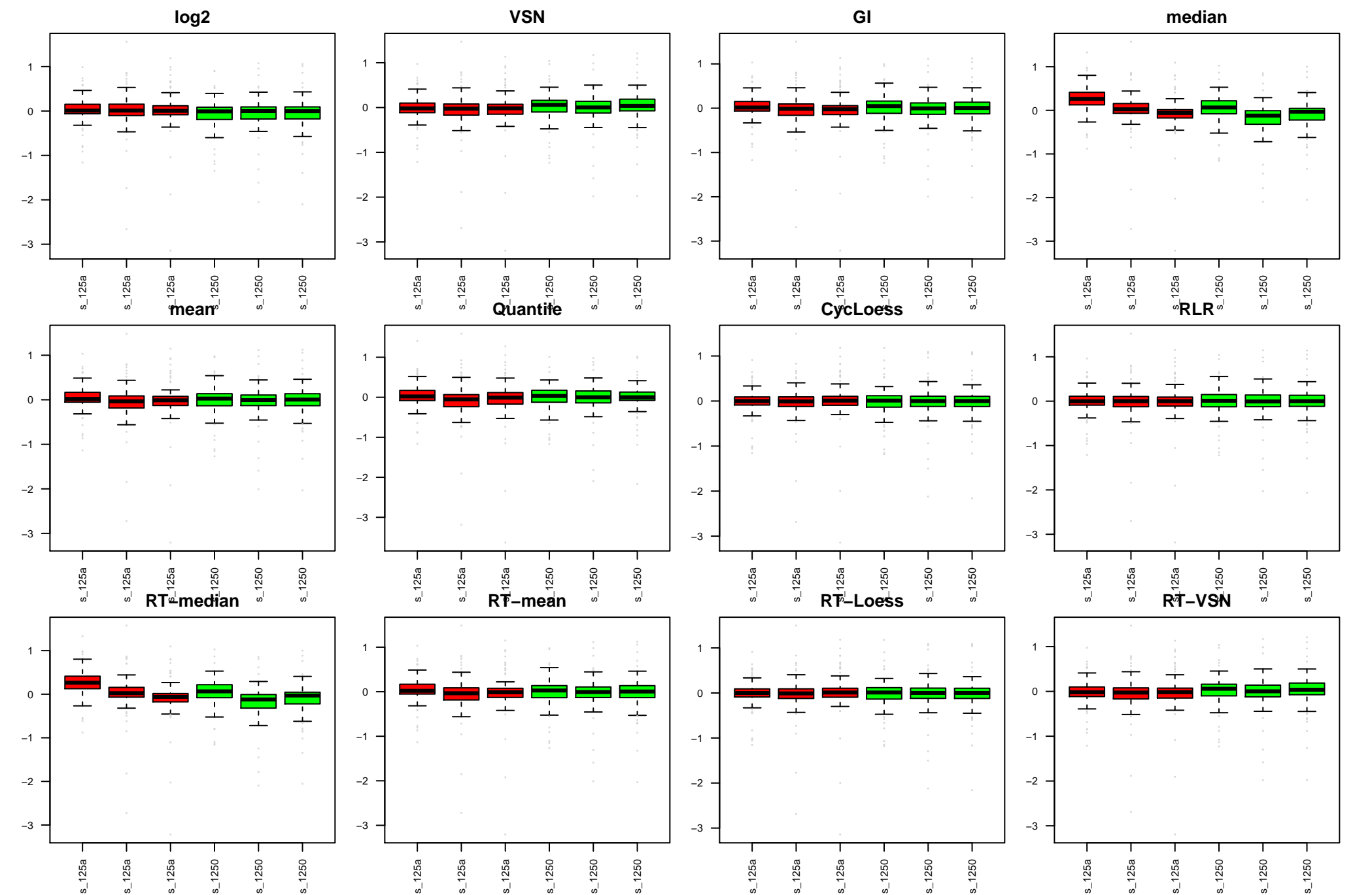




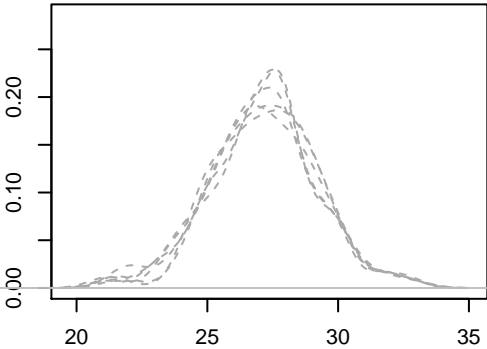




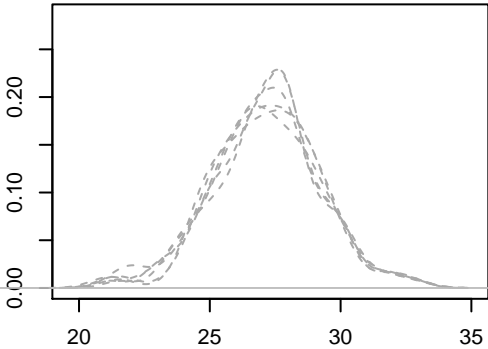




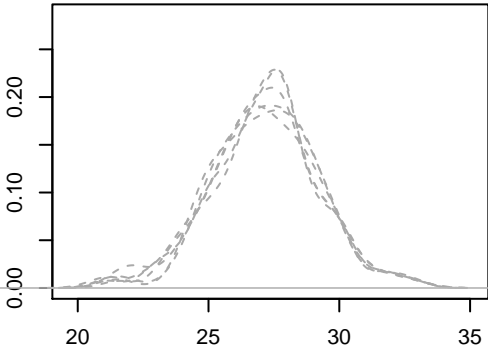
log2



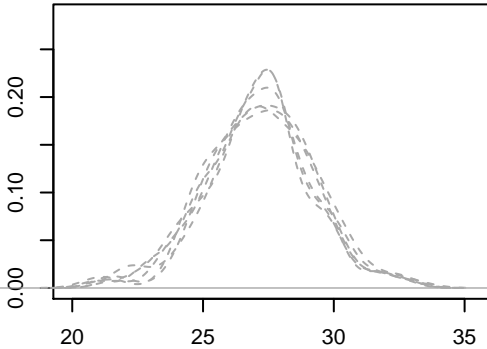
VSN



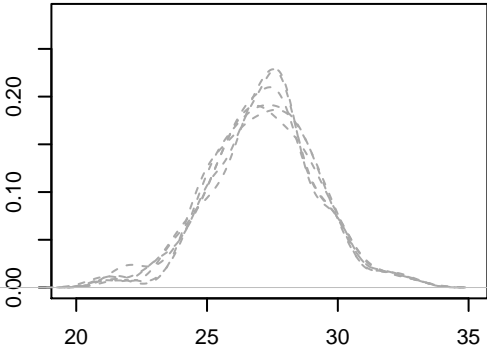
GI



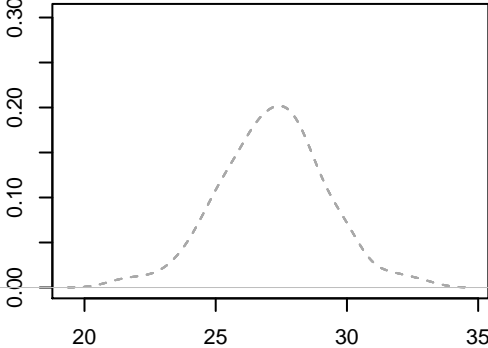
median



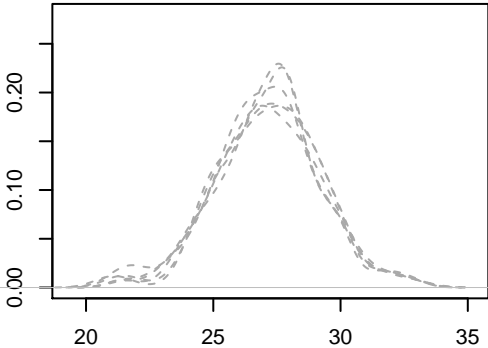
mean



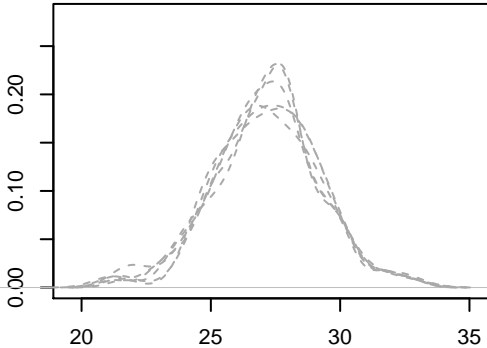
Quantile



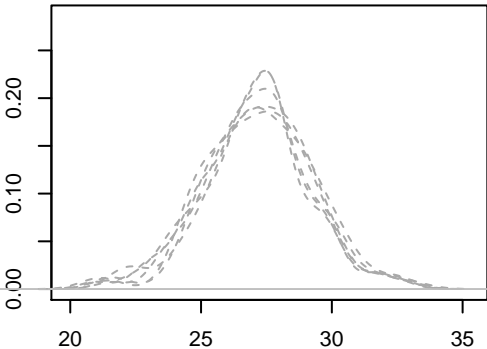
CycLoess



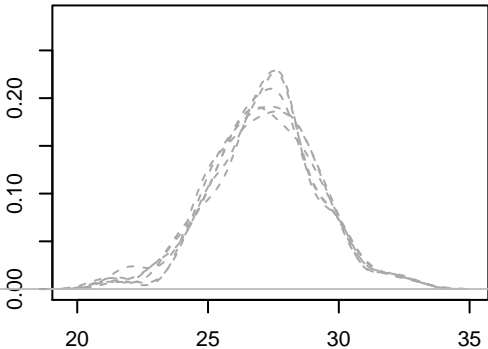
RLR



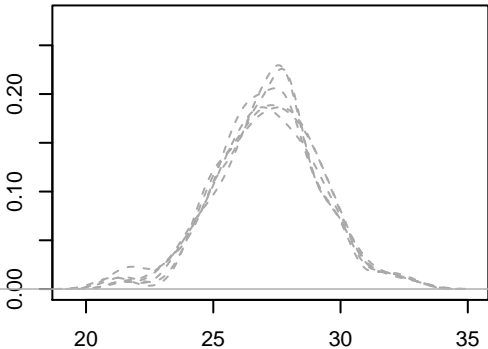
RT-median



RT-mean



RT-Loess



RT-VSN

