

Project Name: `junc.exons_phosphosites`

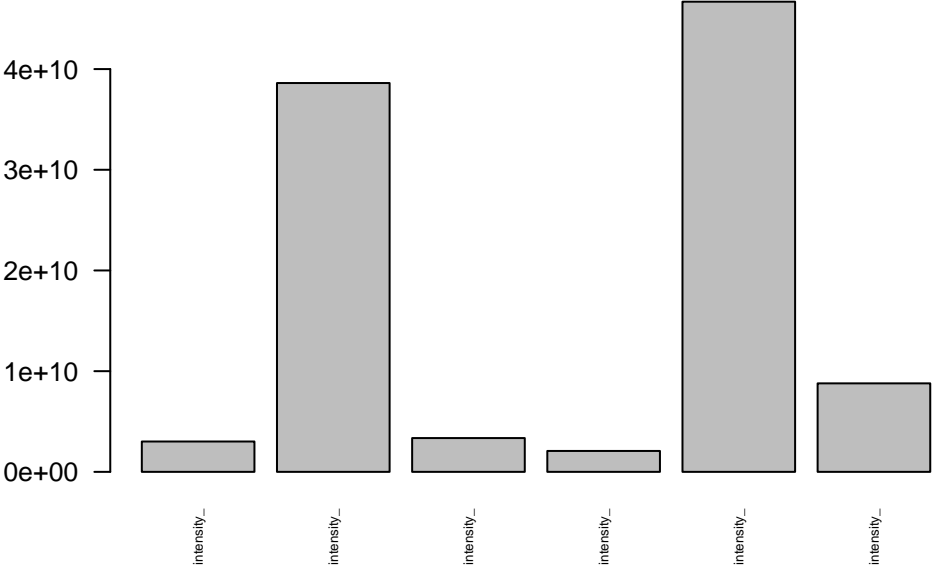
Normalyzer (ver 1.4.0)

Report created on: 2020-05-08

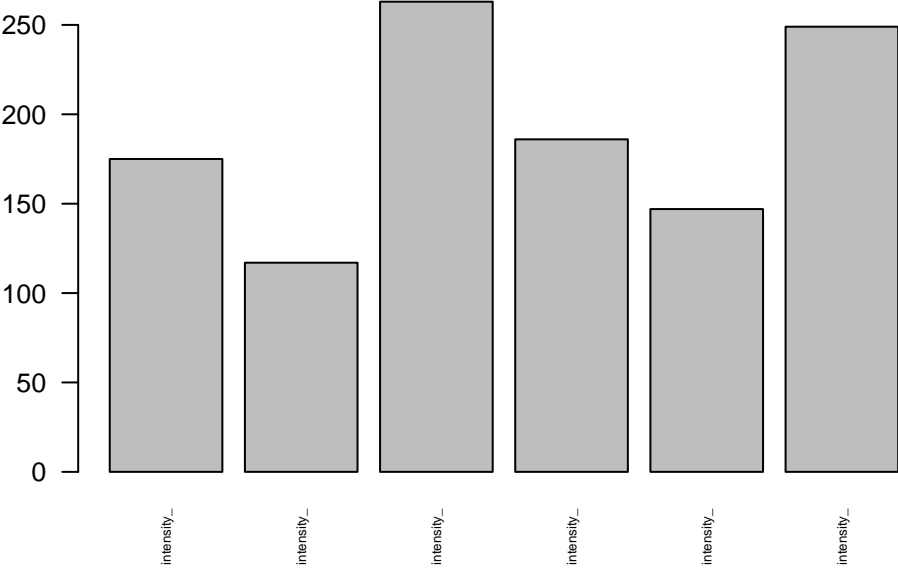
Citation: NormalyzerDE: Online Tool for Improved Normalization of Omics Expression Data and High-Sensitivity Differential Expression Analysis
Journal of Proteome Research (2018), 10.1021/acs.jproteome.8b00523

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

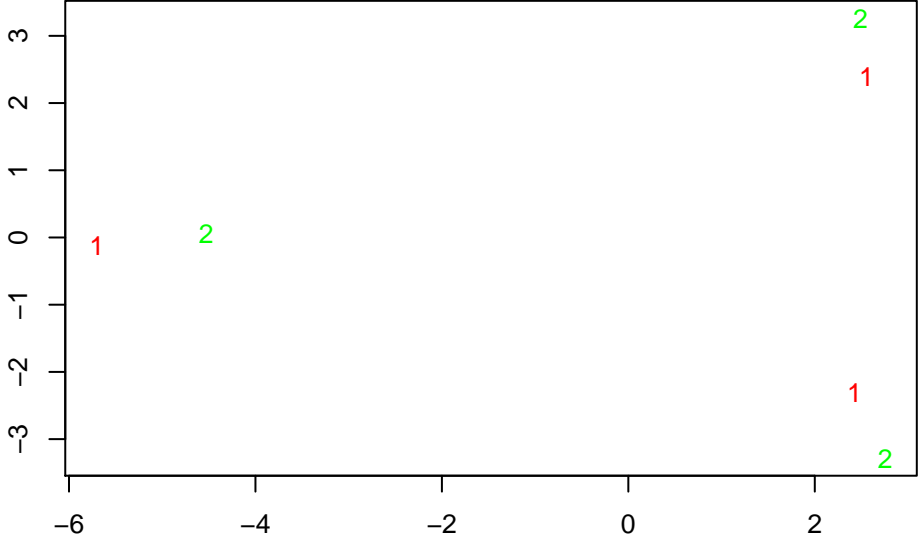
Total intensity



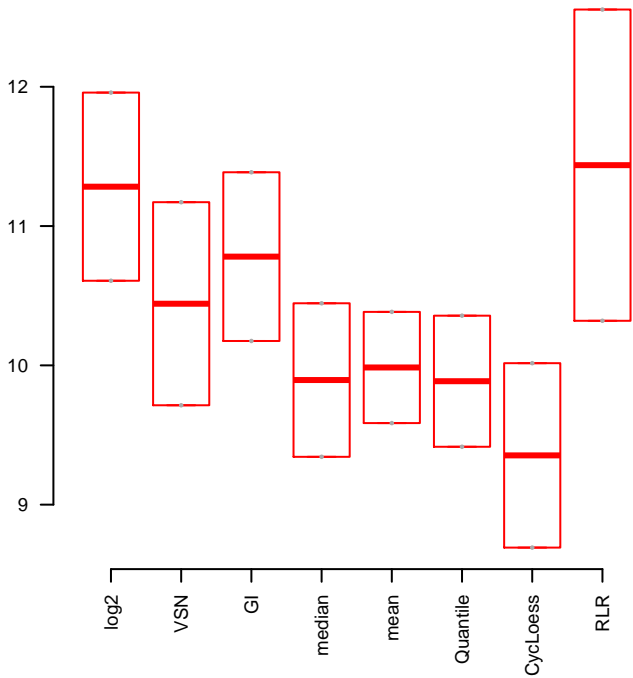
Total missing



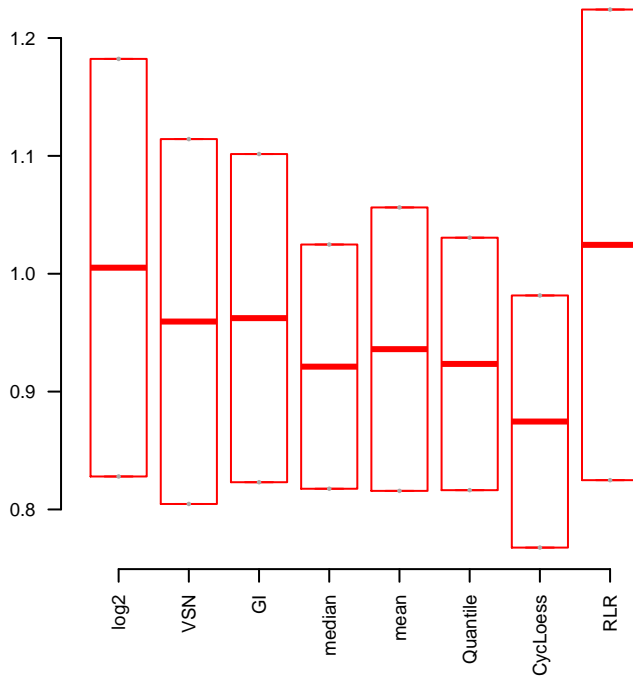
Log2-MDS plot



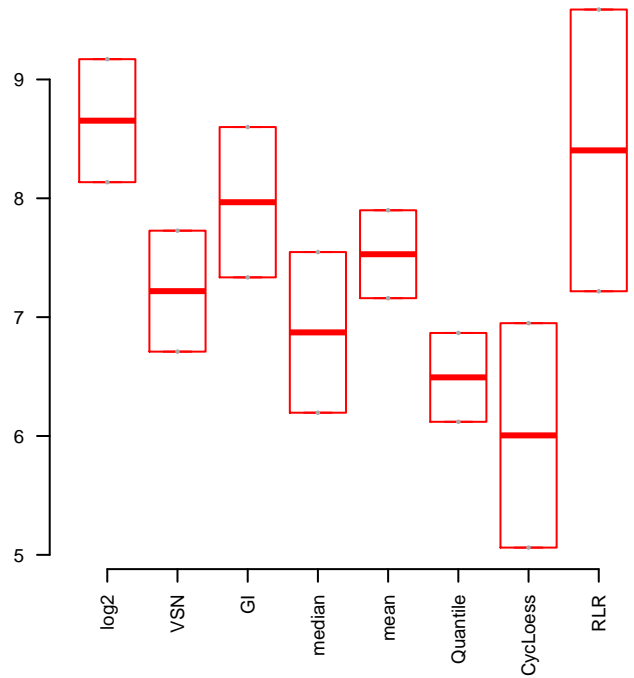
PCV – Intragroup



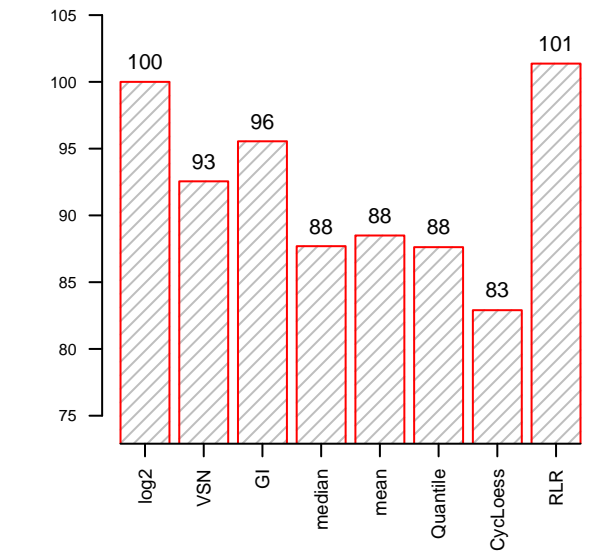
PMAD – Intragroup



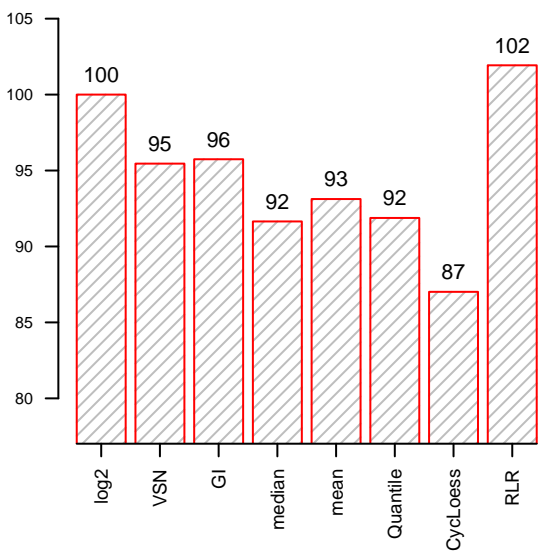
PEV – Intragroup



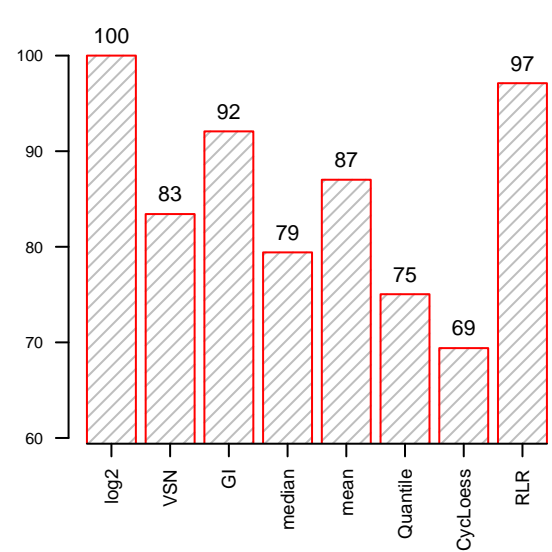
PCV compared to log2



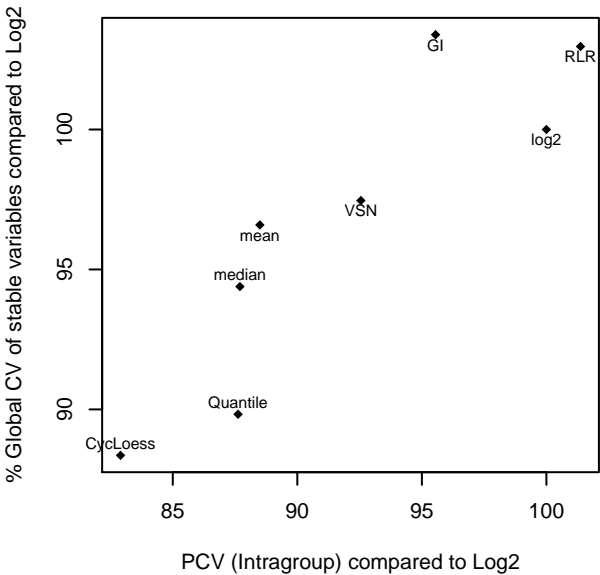
PMAD compared to log2

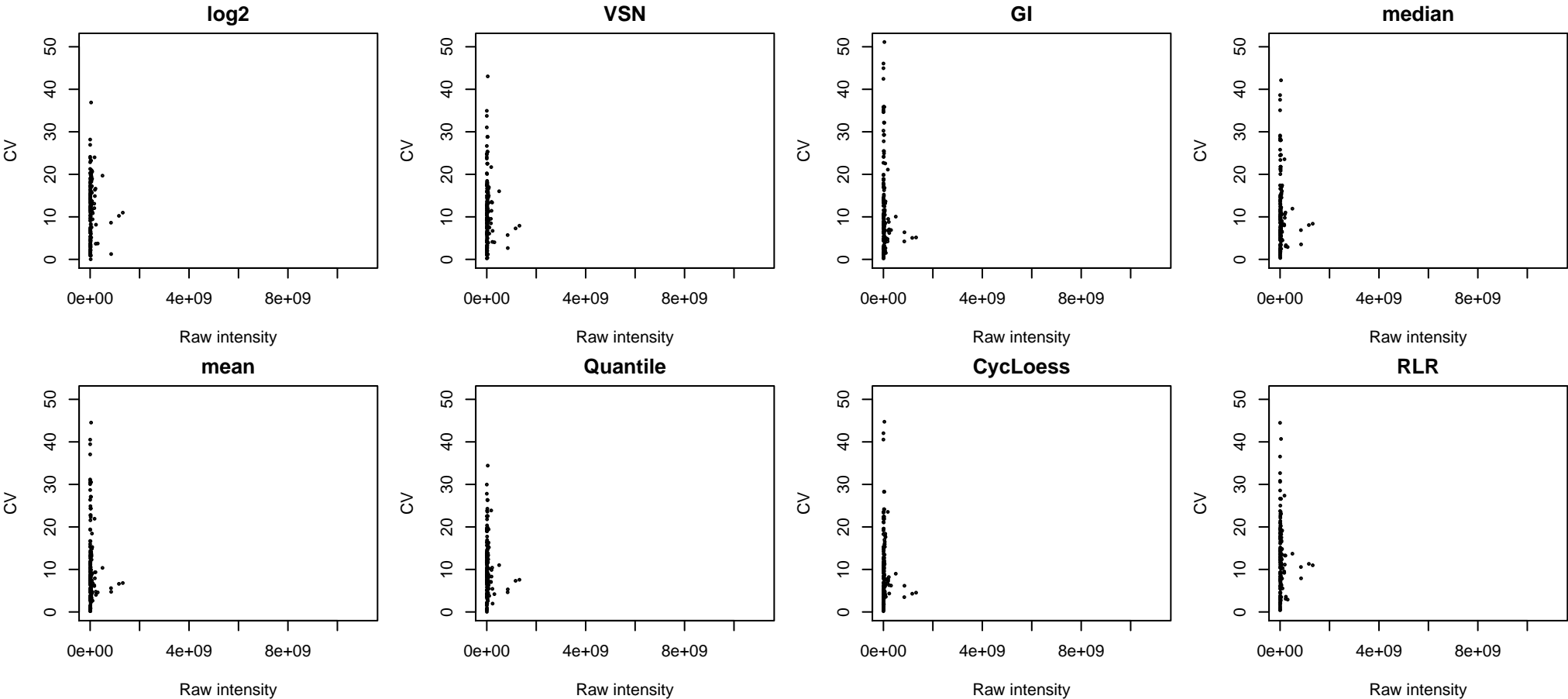


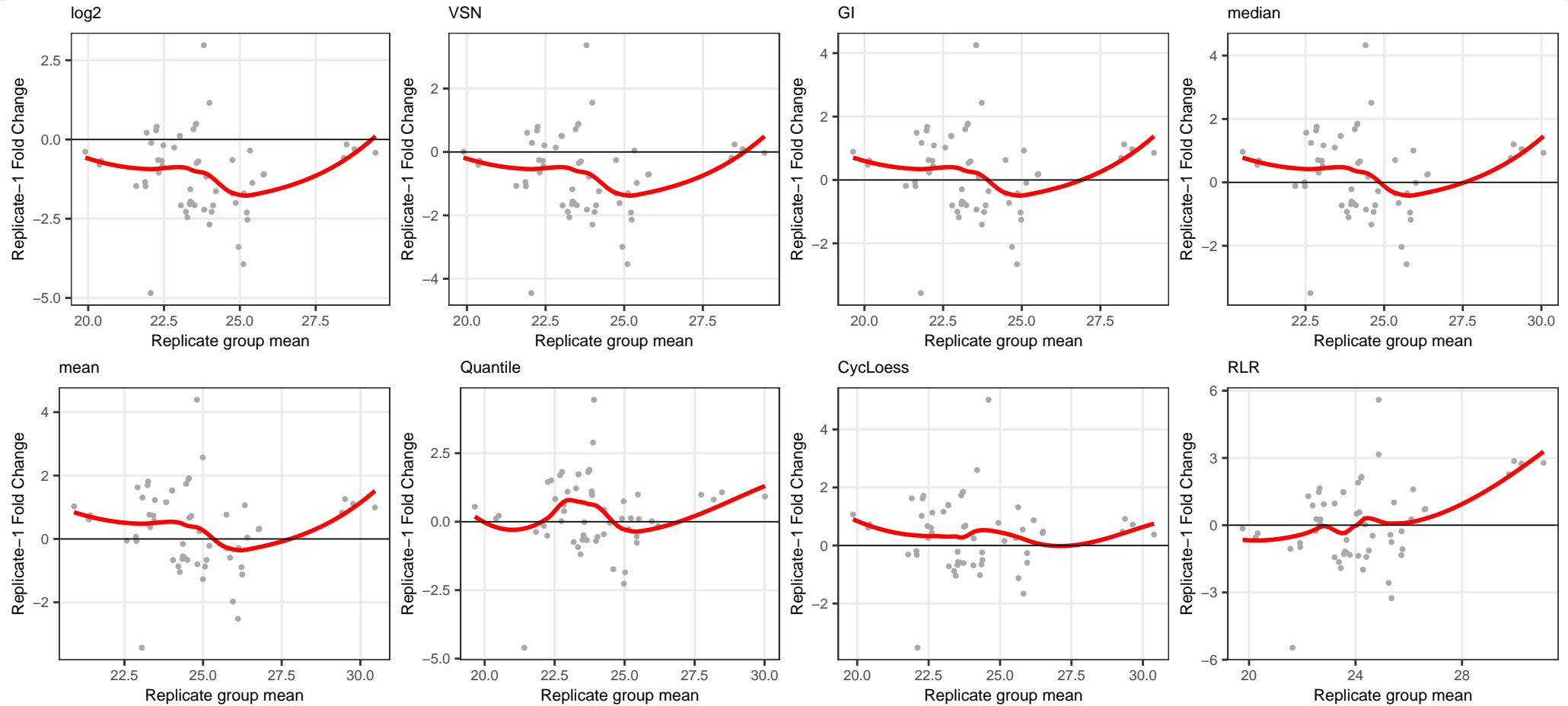
%PEV – compared to log2

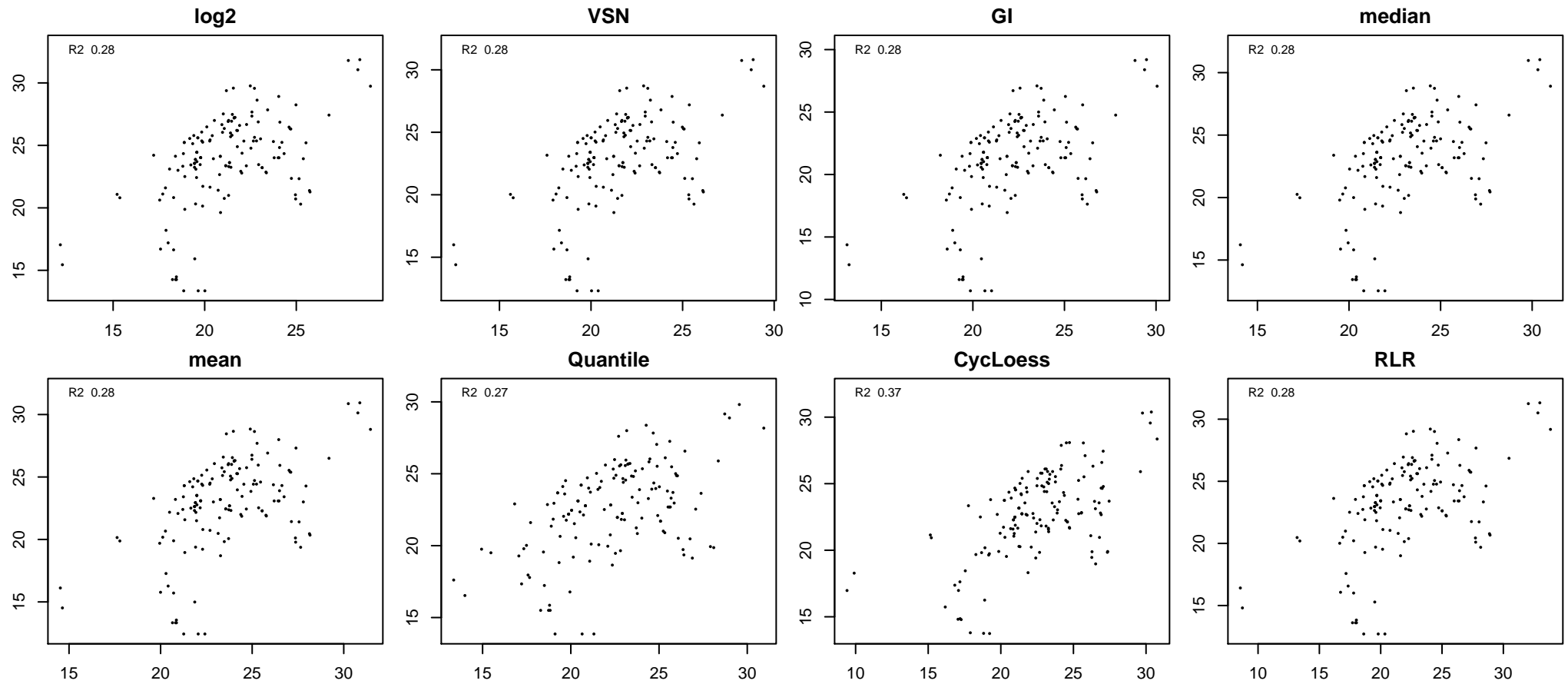


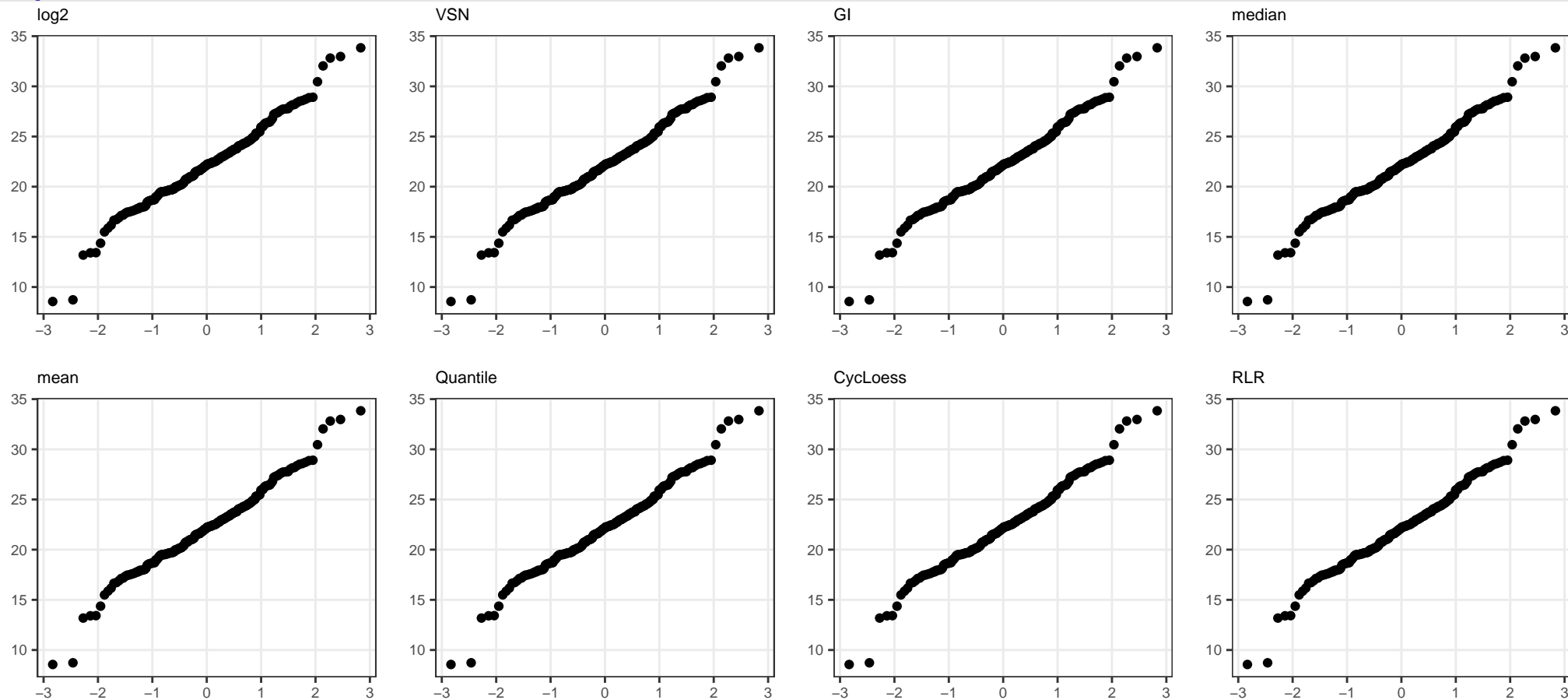
Stable variables plot

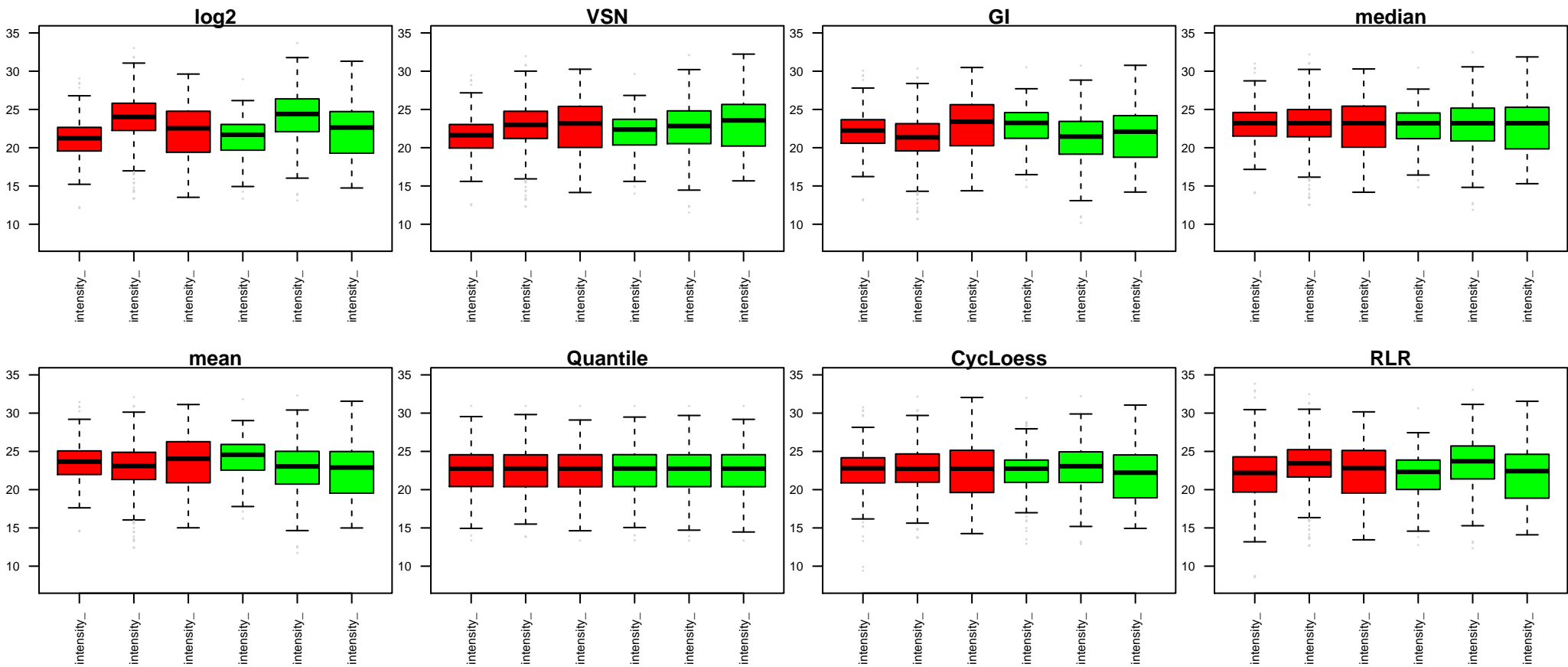


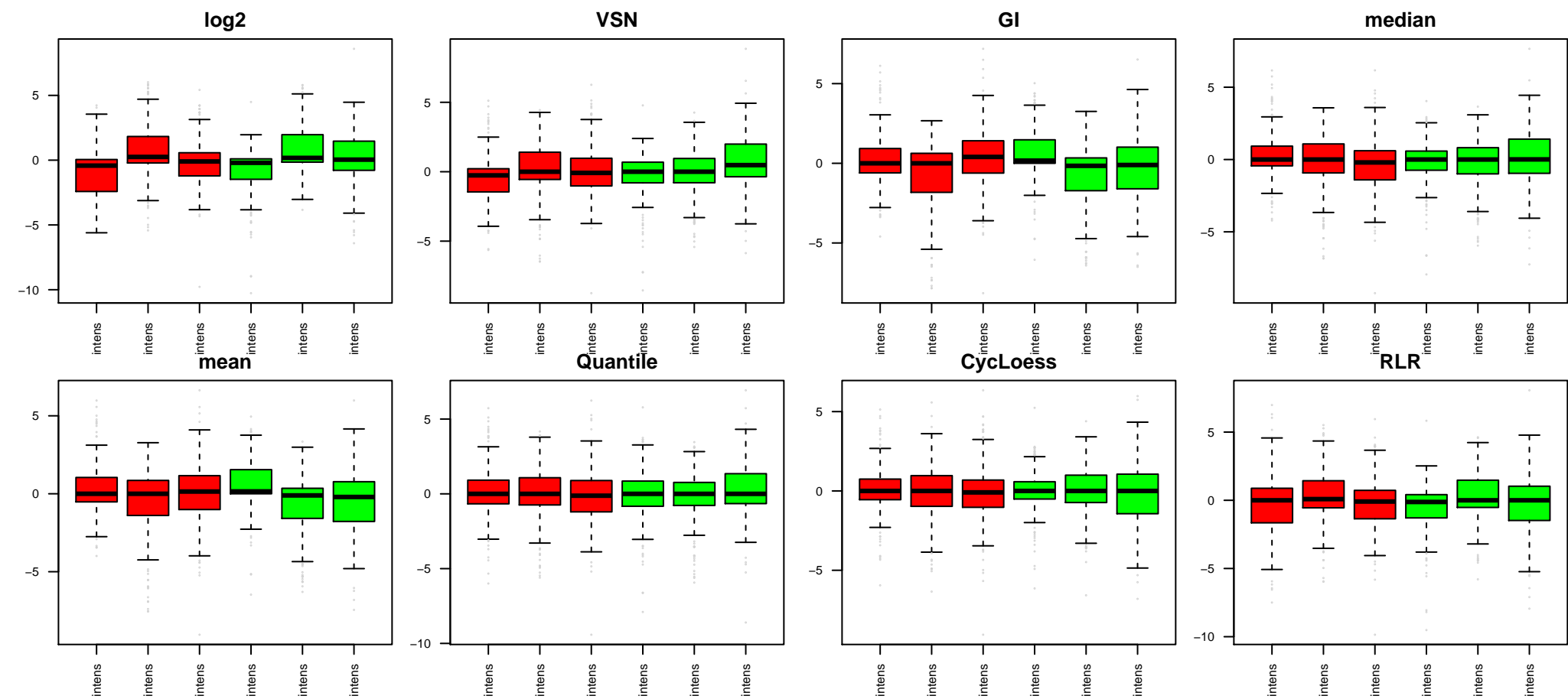




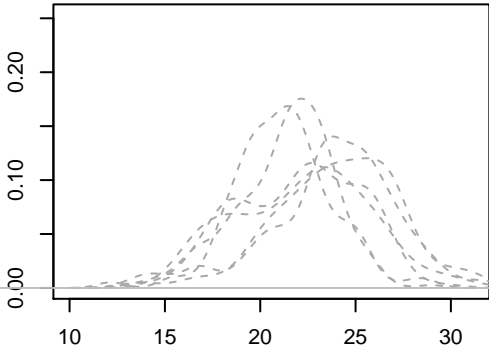




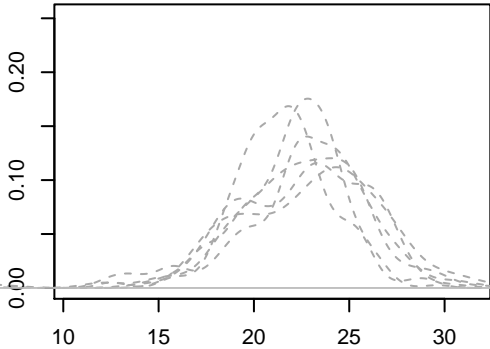




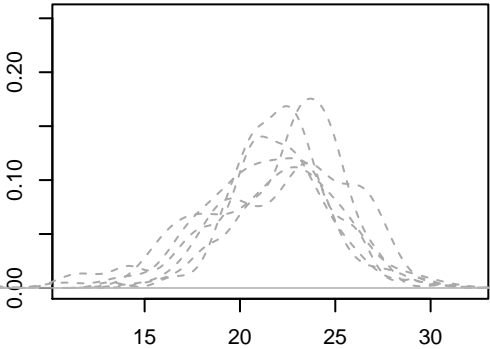
log2



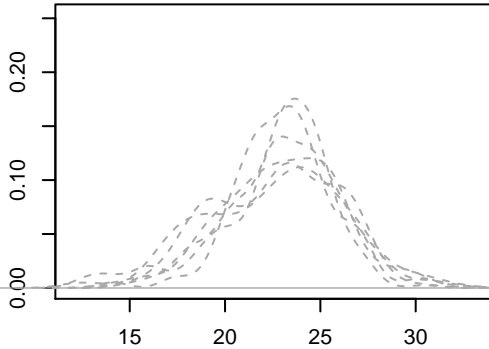
VSN



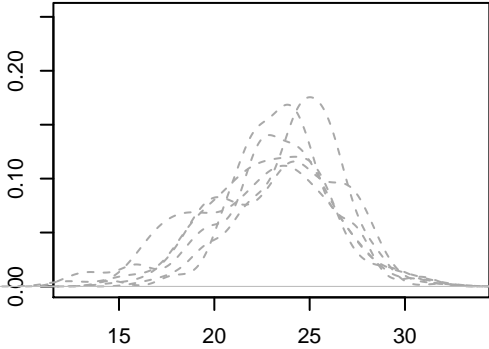
GI



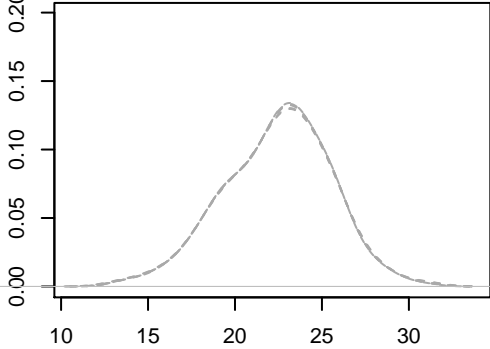
median



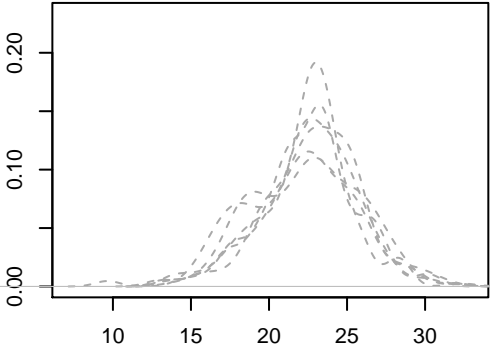
mean



Quantile



CycLoess



RLR

