

Project Name: AT_Humedad_norm

NormalizerDE (ver 1.20.0)

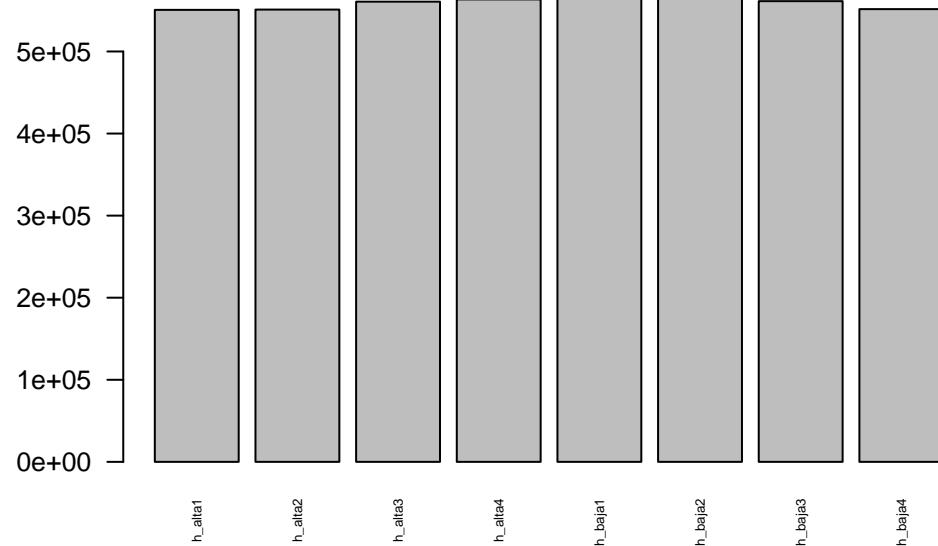
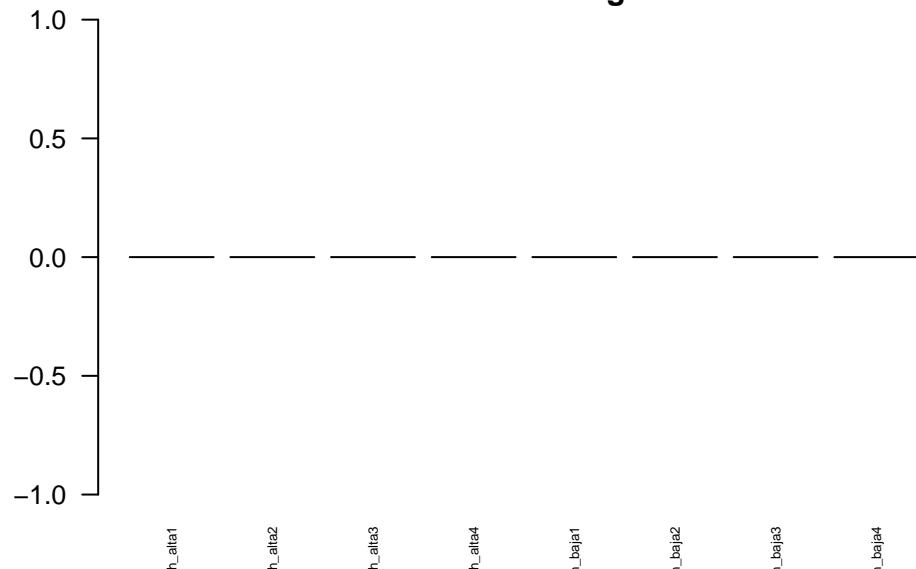
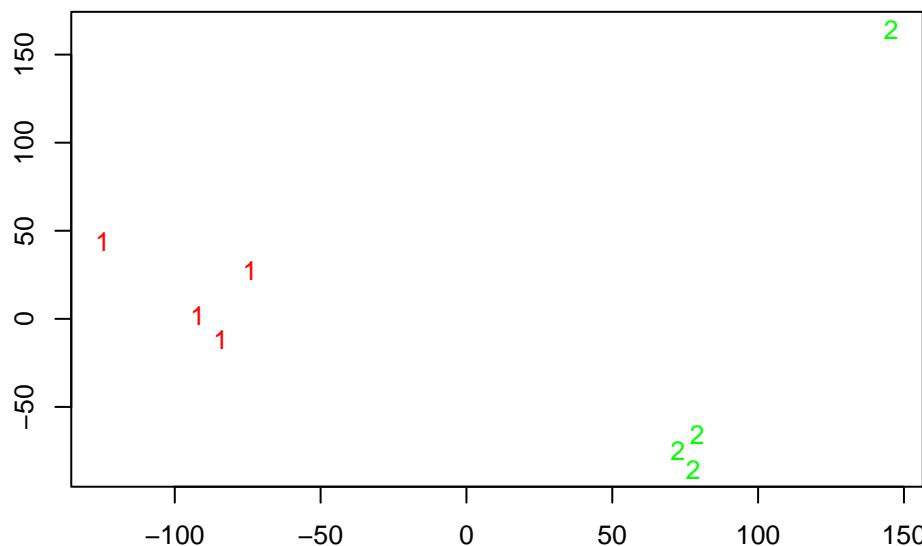
Report created on: 2024–01–24

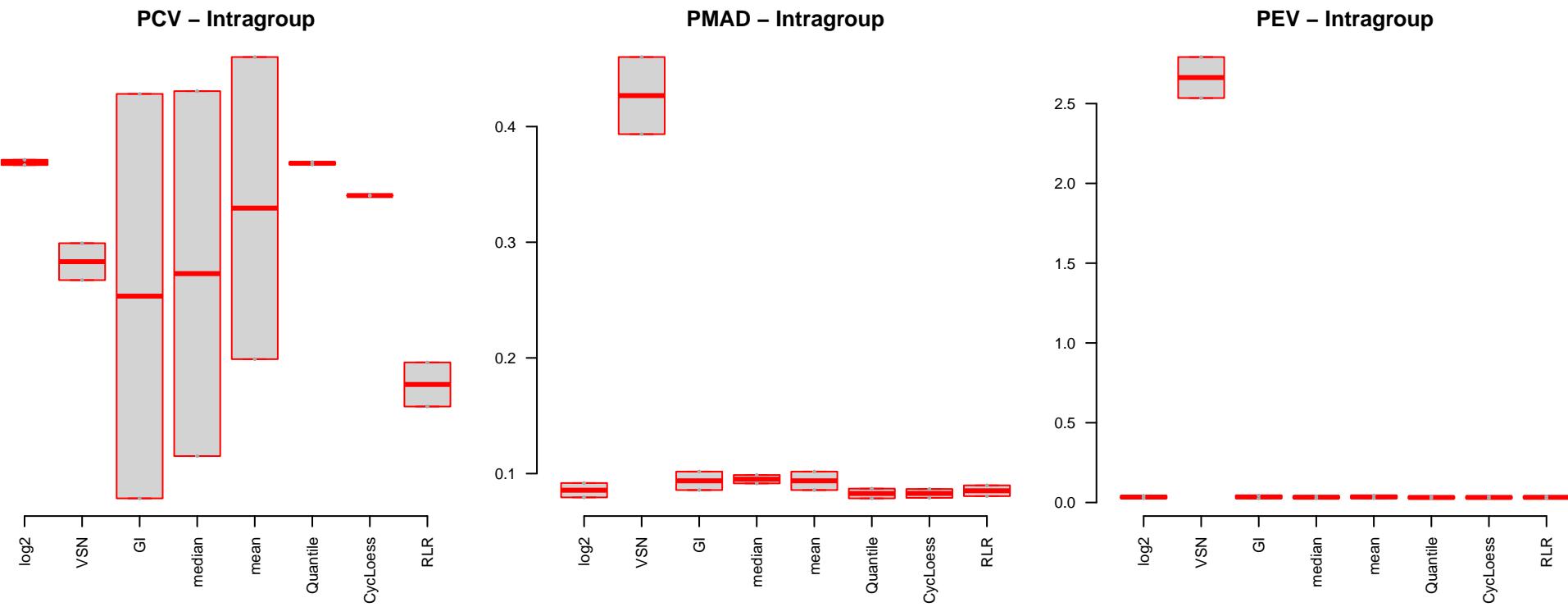
Citation: NormalizerDE: Online Tool for Improved Normalization of Omics Expression Data and High-Sensitivity Differential Expression Analysis
Journal of Proteome Research (2018), 10.1021/acs.jproteome.8b00523

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

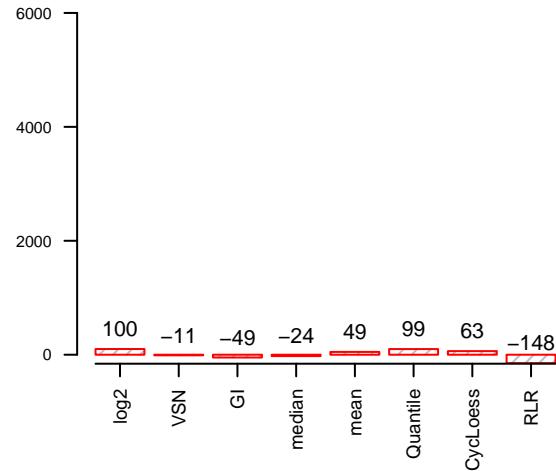
Group nbr.	Design group	Nbr. samples in cond.
1	h_alta	4
2	h_baja	4

Please note that the grouping only impacts evaluation measures and visuals seen in the report, they do not impact the performed normalizations.

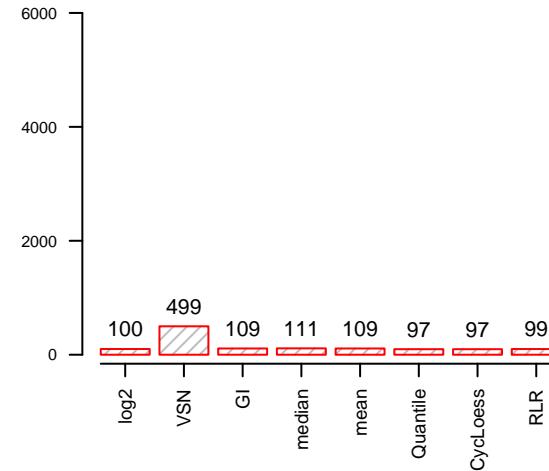
Total intensity**Total missing****Log2-MDS plot**



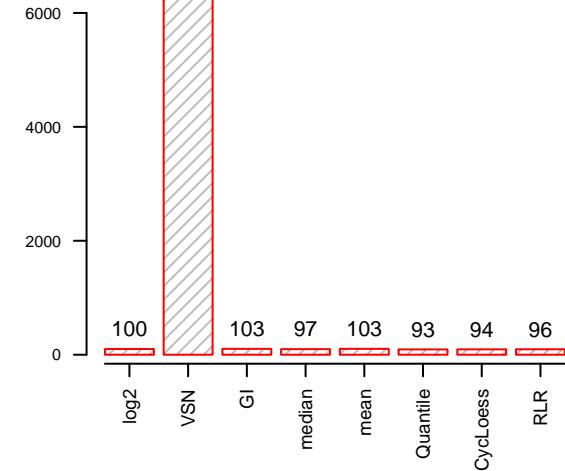
PCV compared to log2



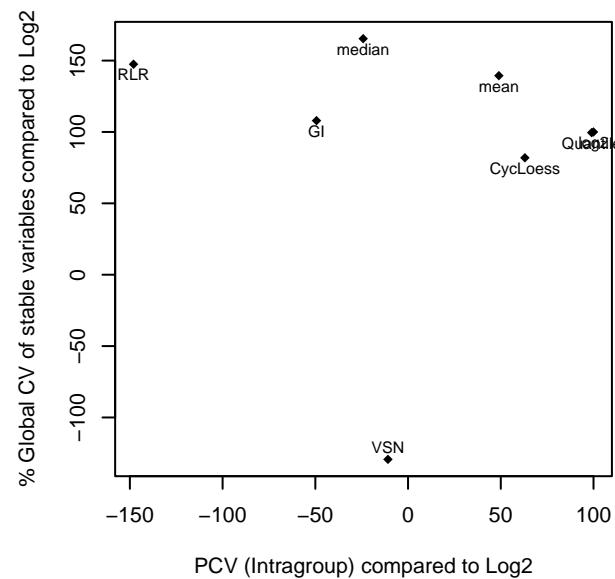
PMAD compared to log2

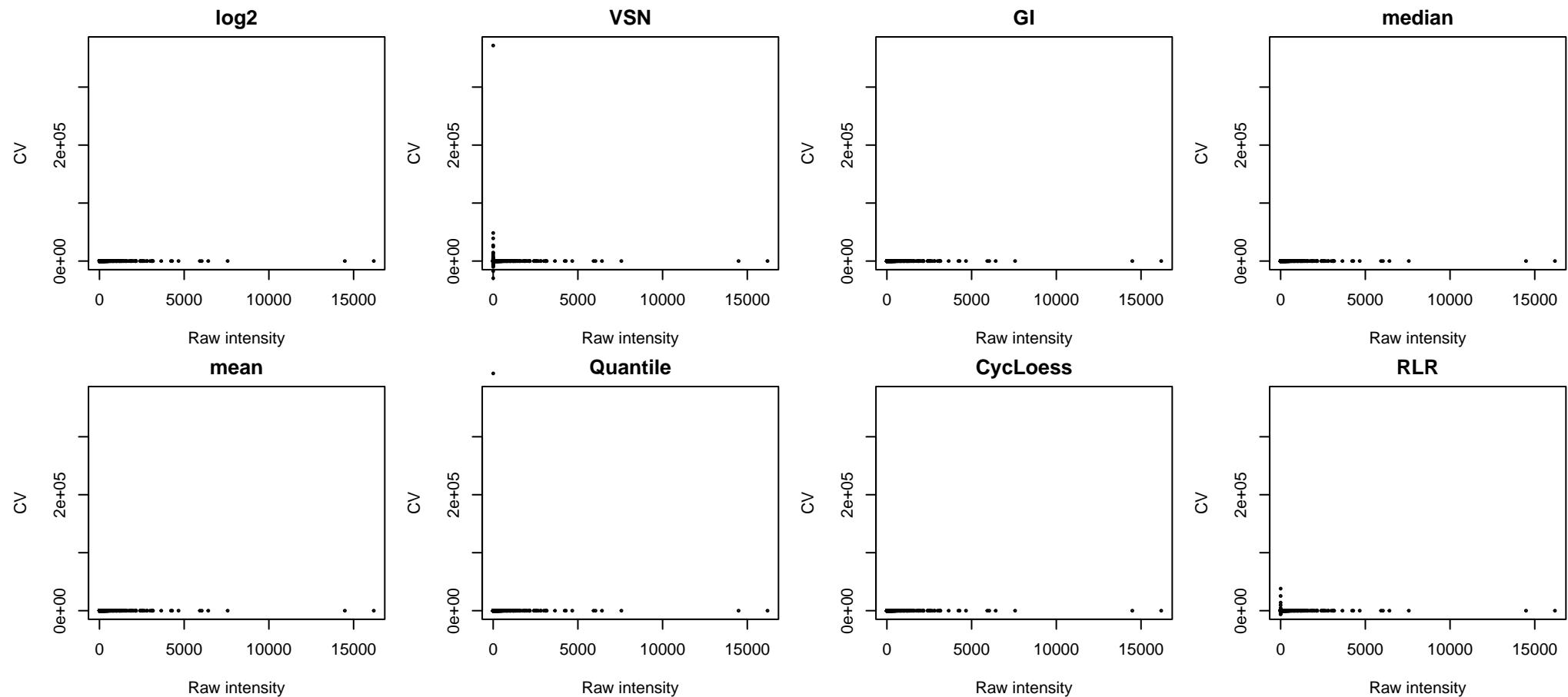


%PEV – compared to log2



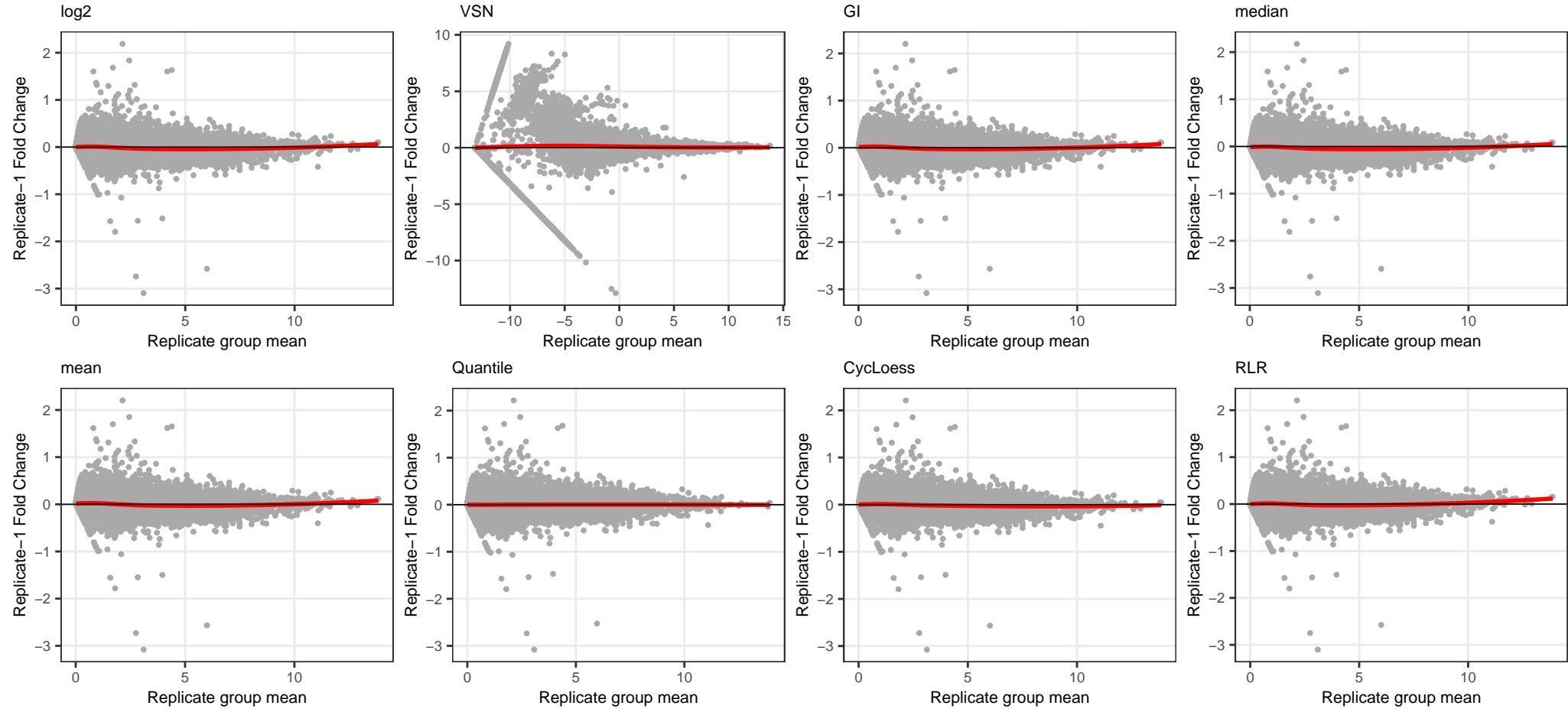
Stable variables plot

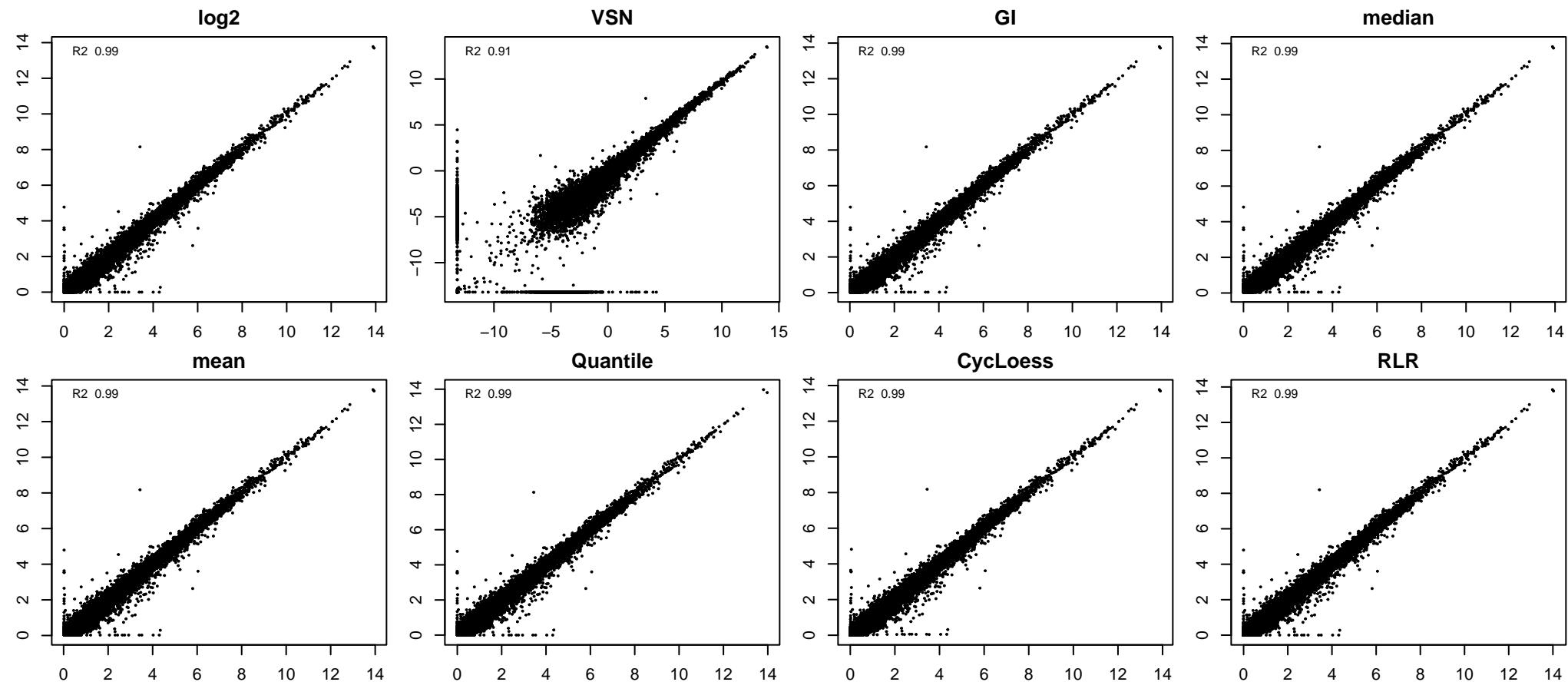




MA plots

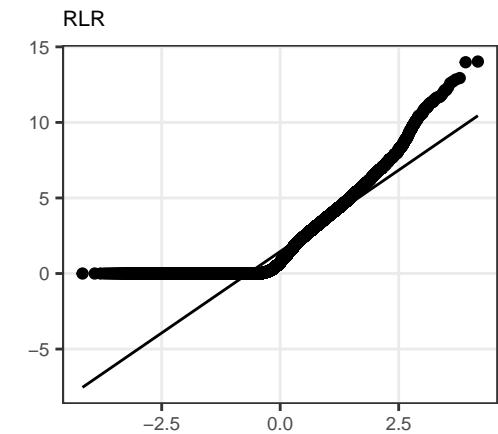
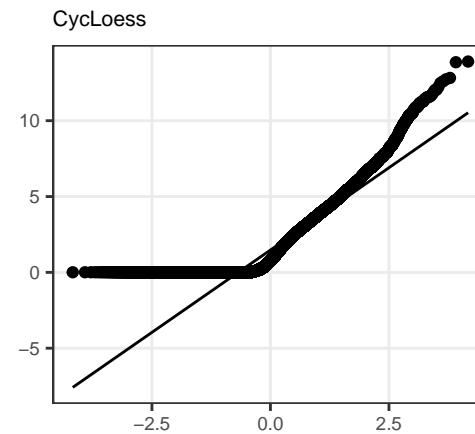
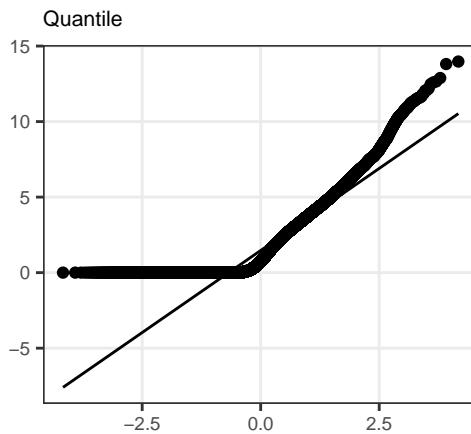
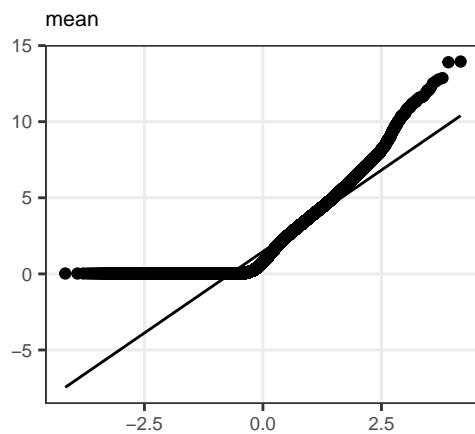
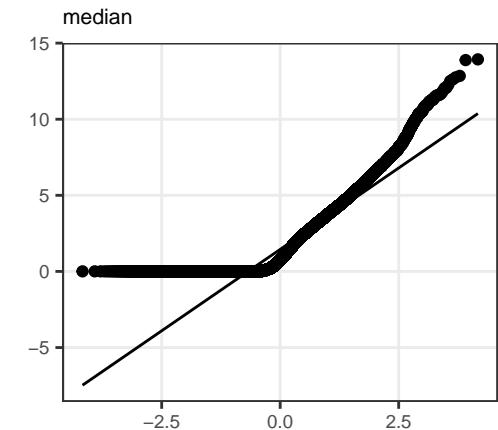
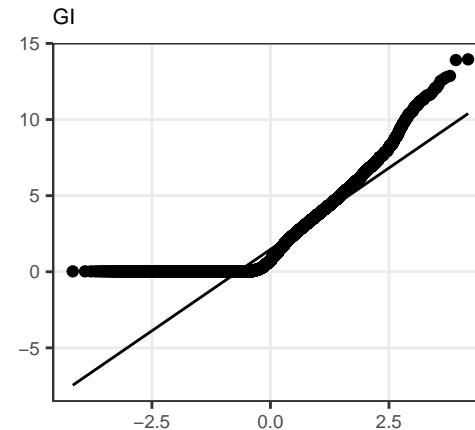
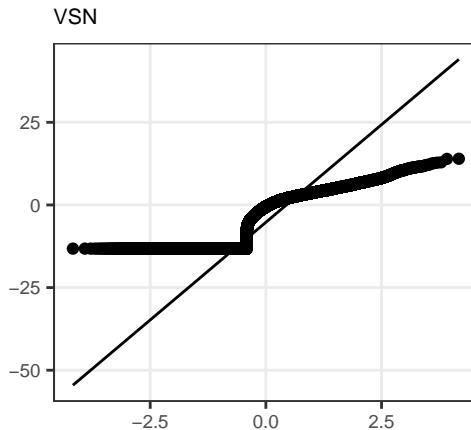
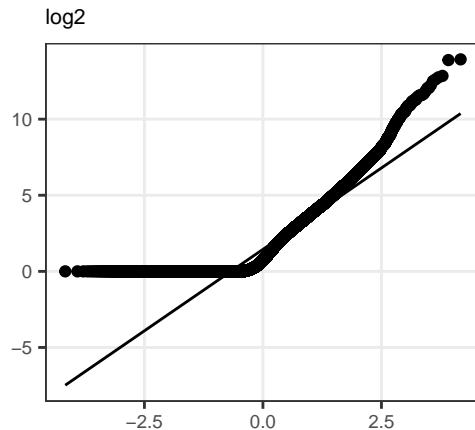
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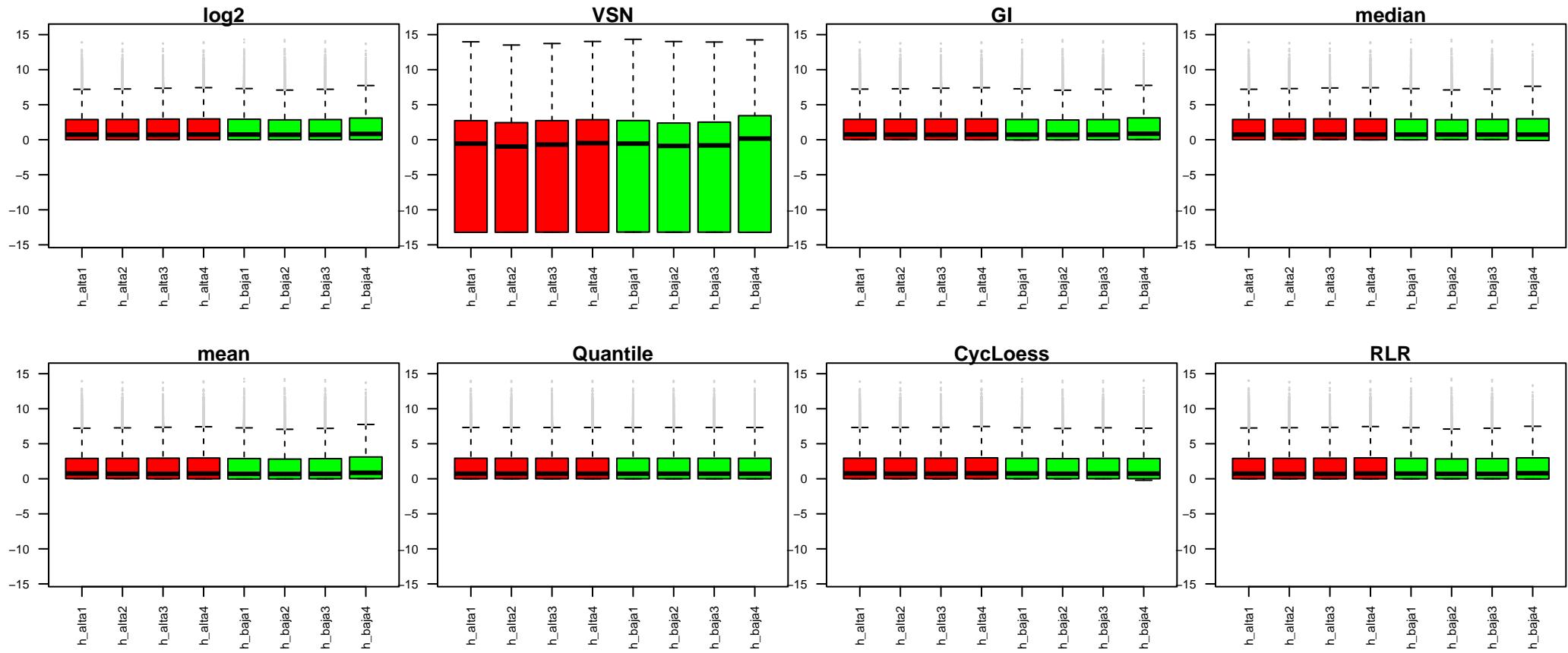




Q-Q plots

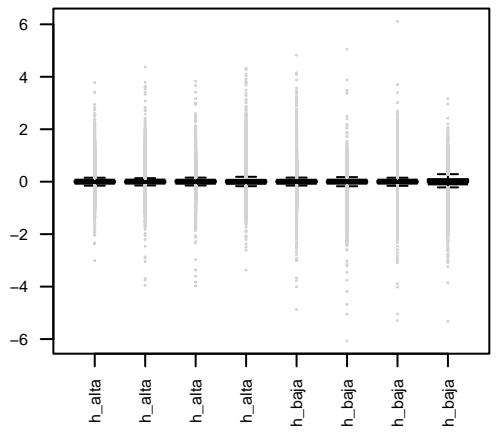
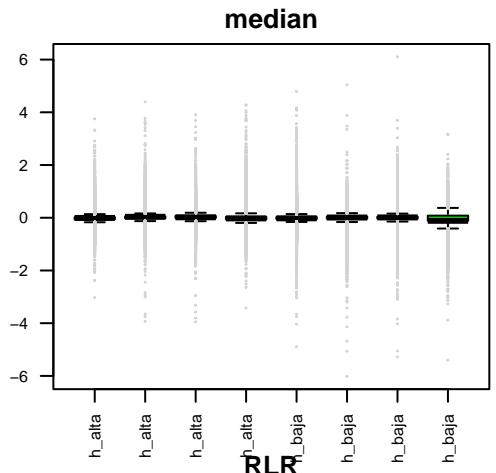
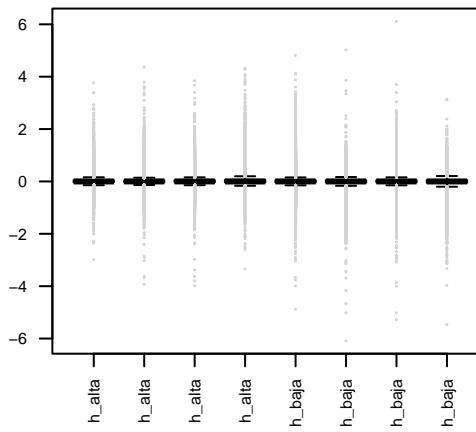
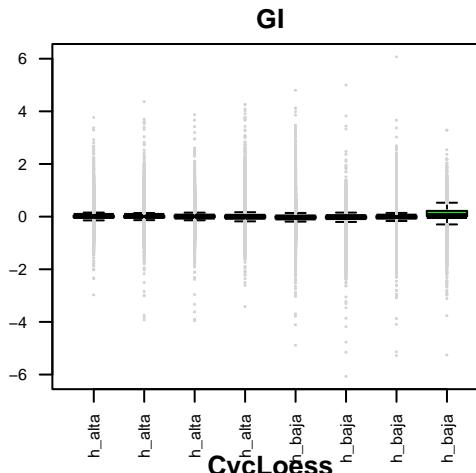
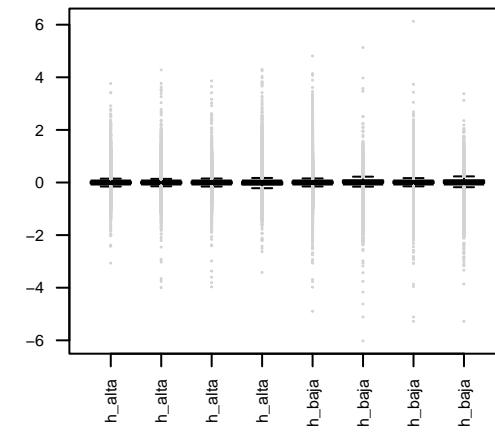
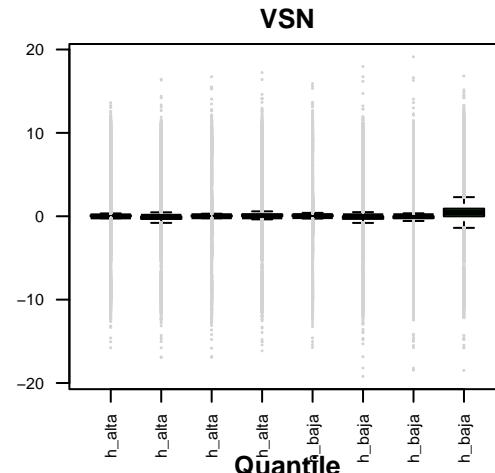
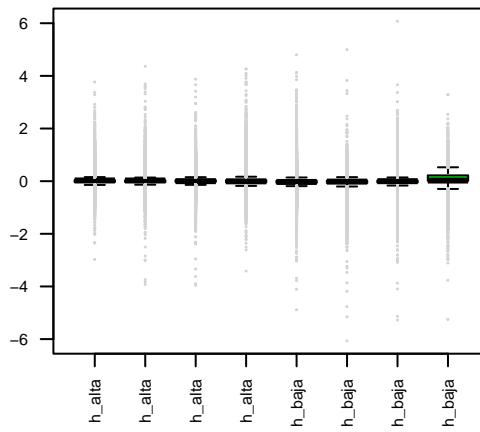
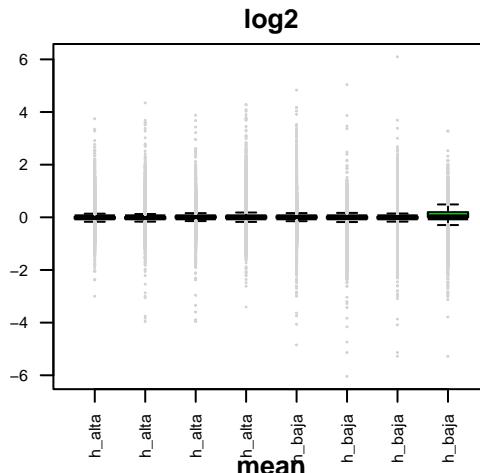
NormalizerDE Report

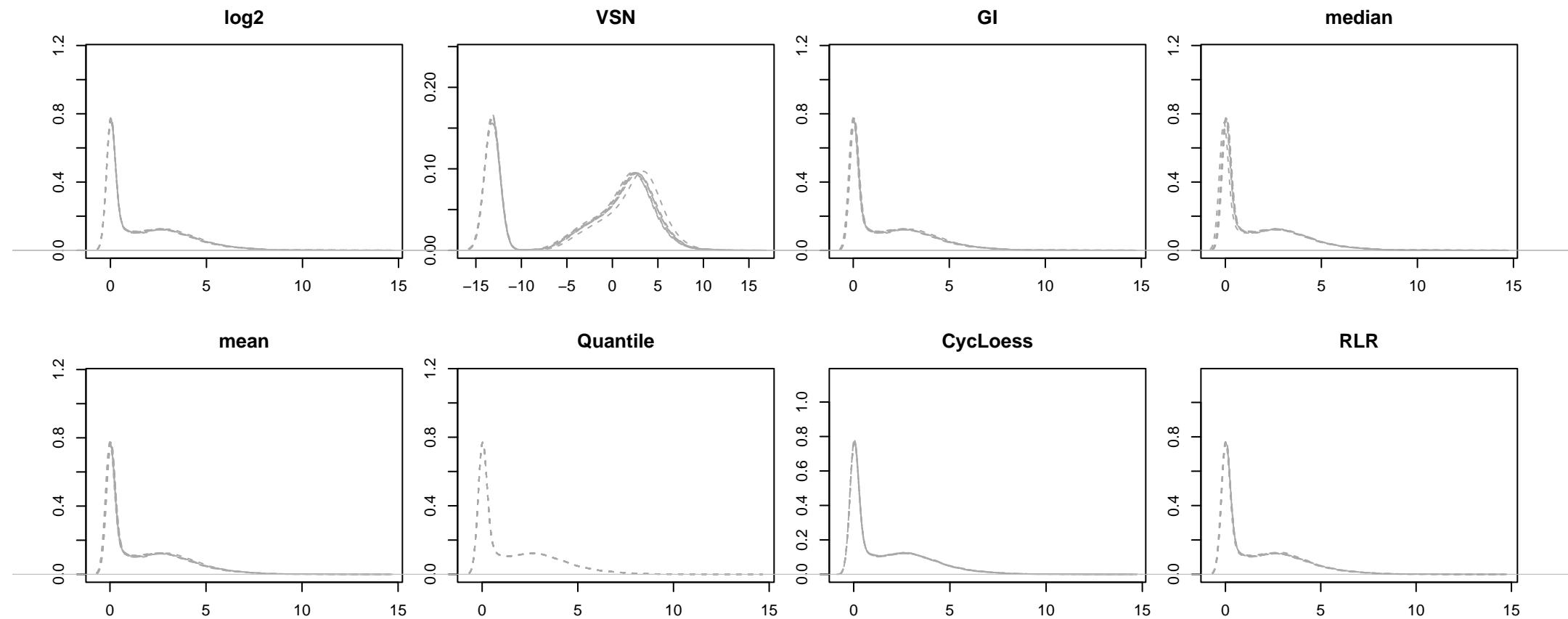


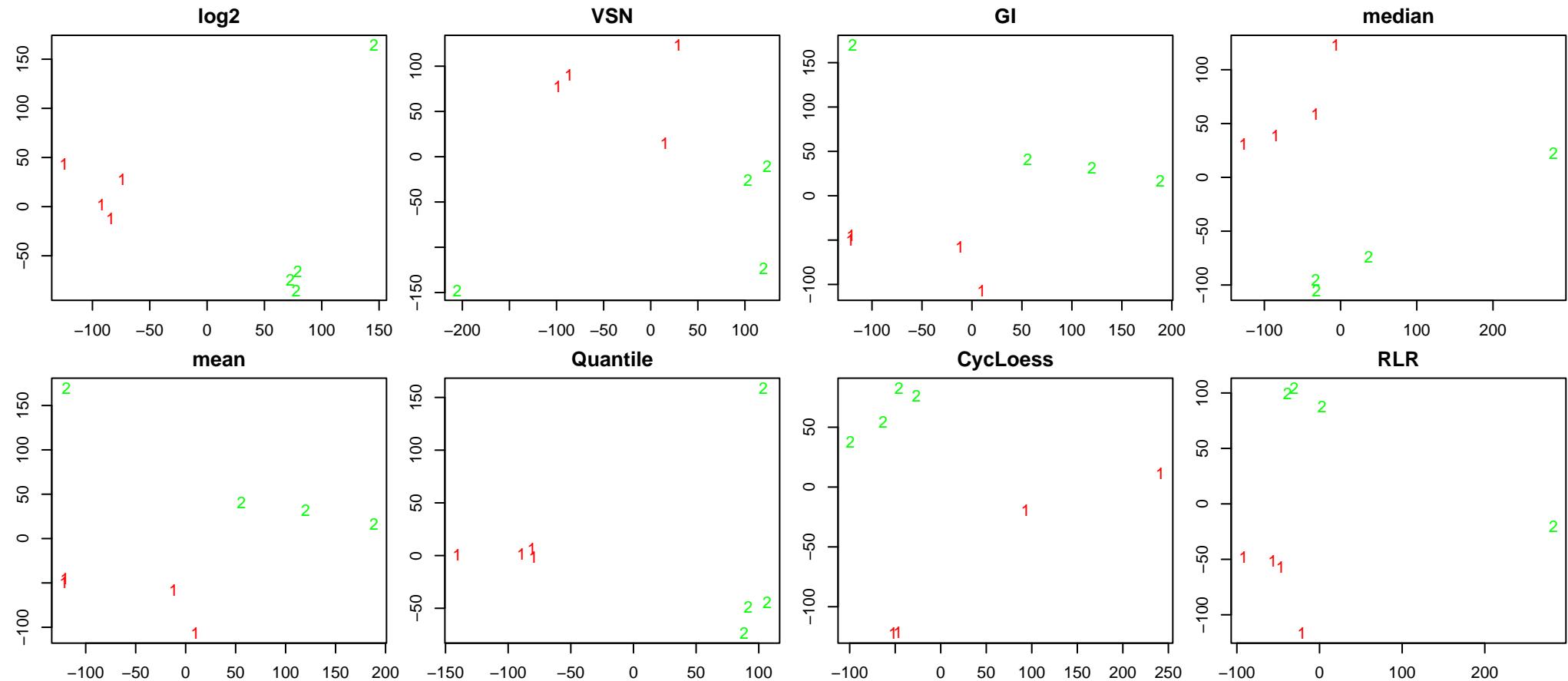


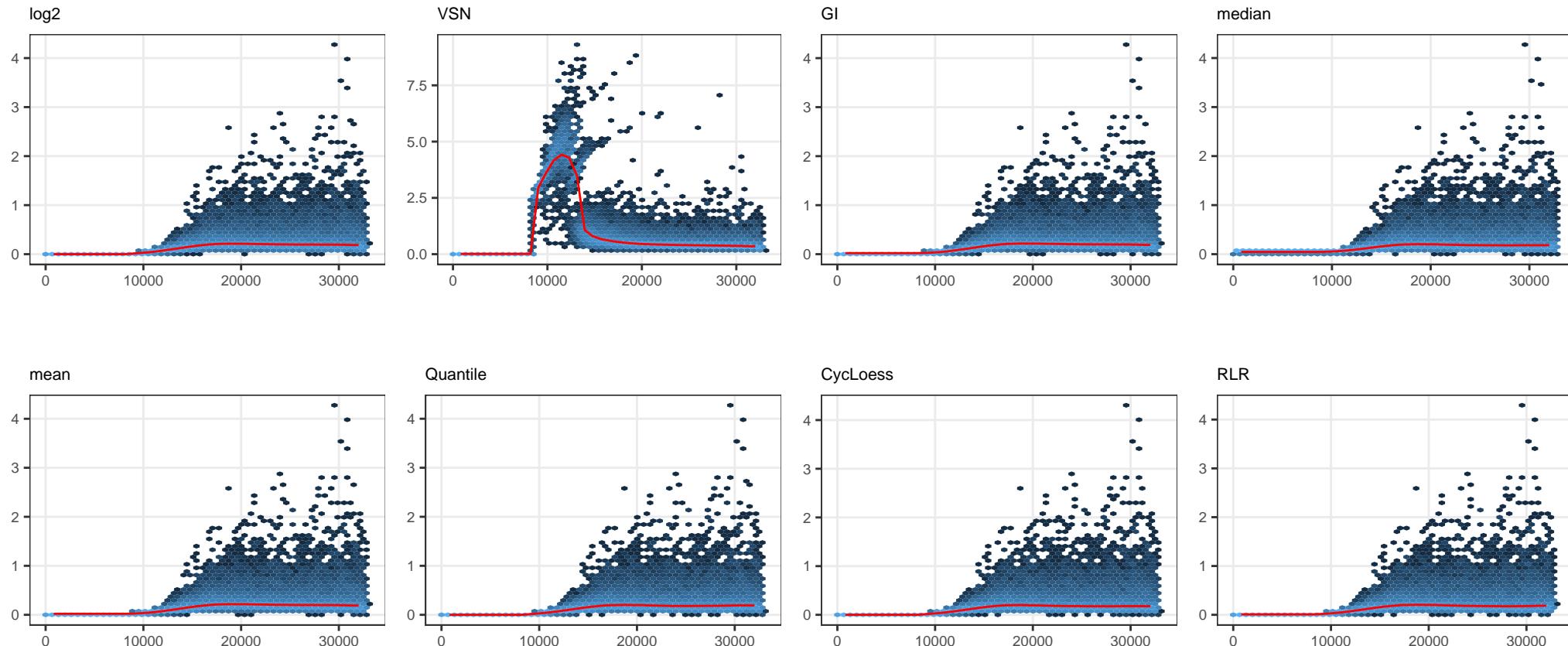
Relative Log Expression (RLE) plots

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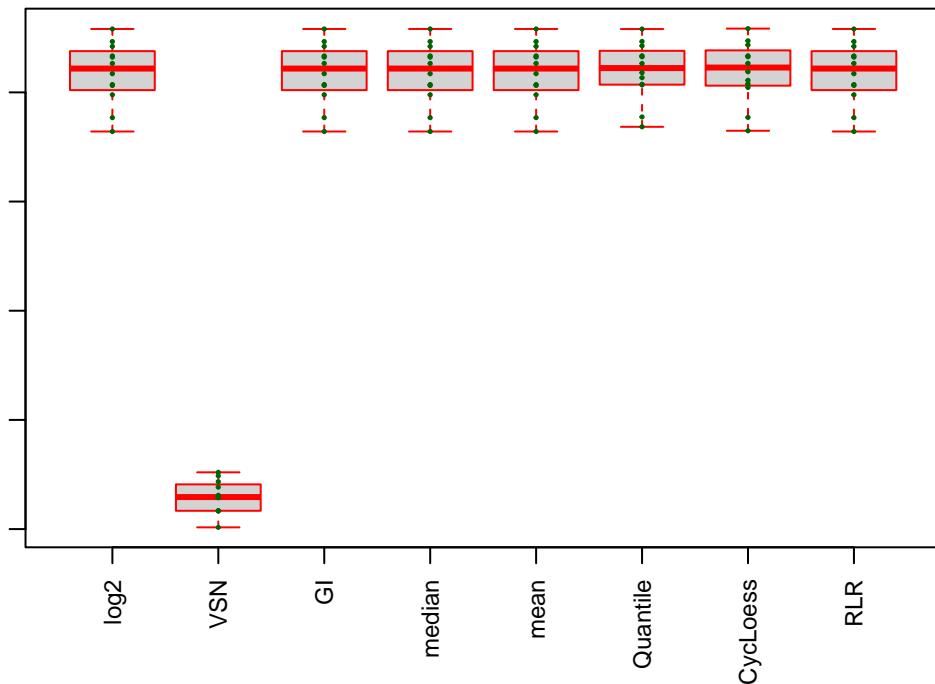




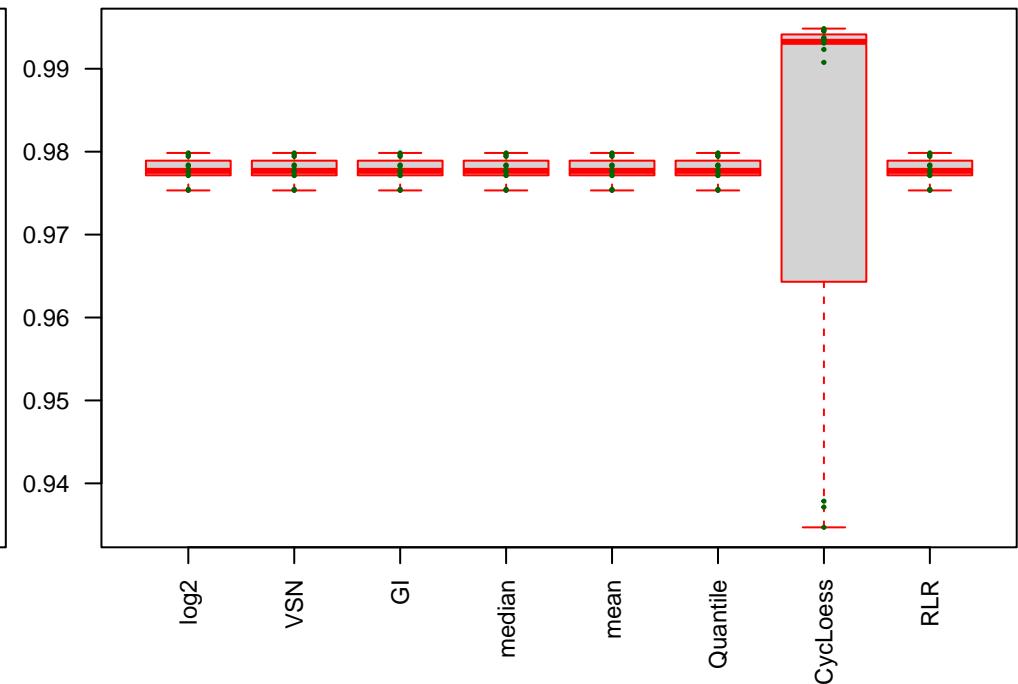


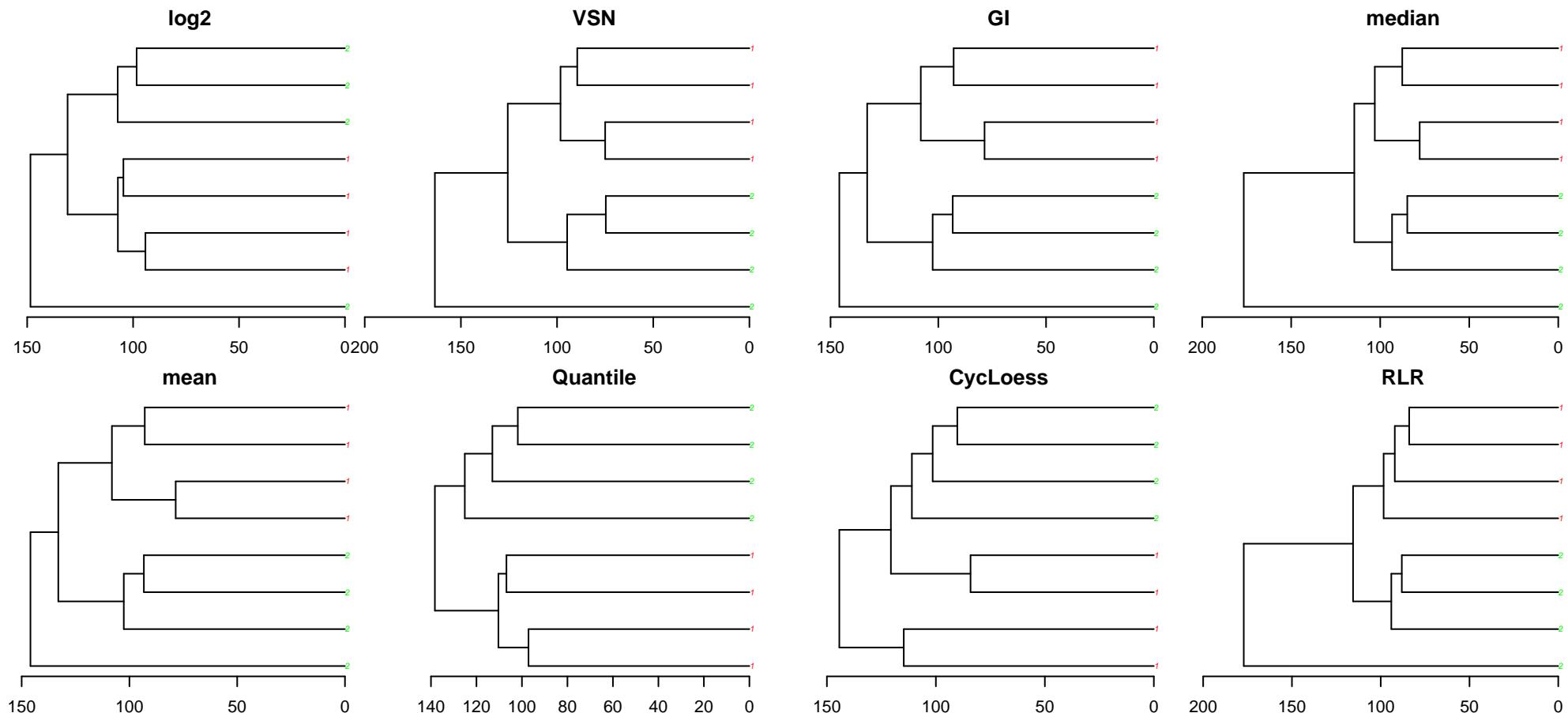


Pearson correlation – Intragroup



Spearman correlation – Intragroup





HistPlots

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