

Project Name: organoids_prot_frac

NormalizerDE (ver 1.14.0)

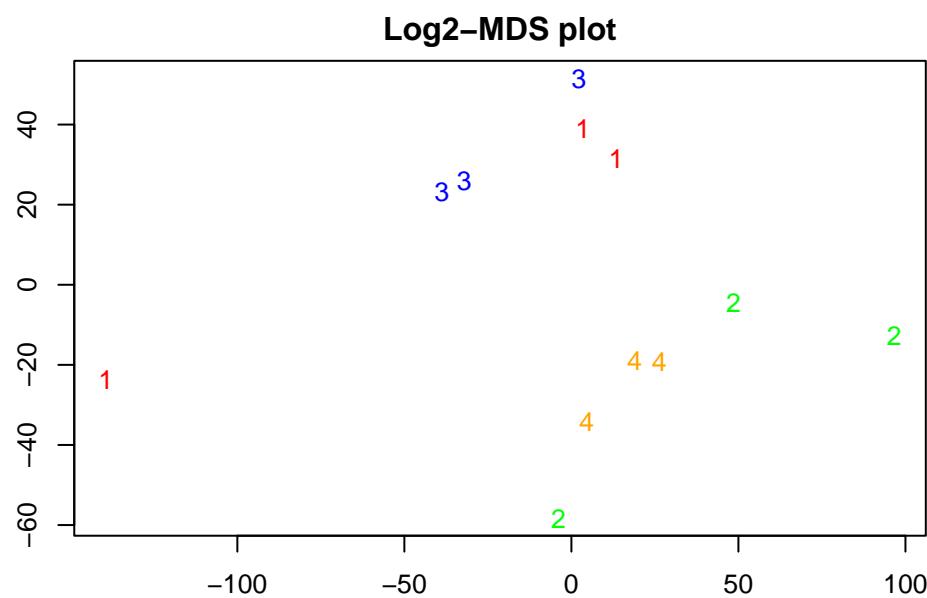
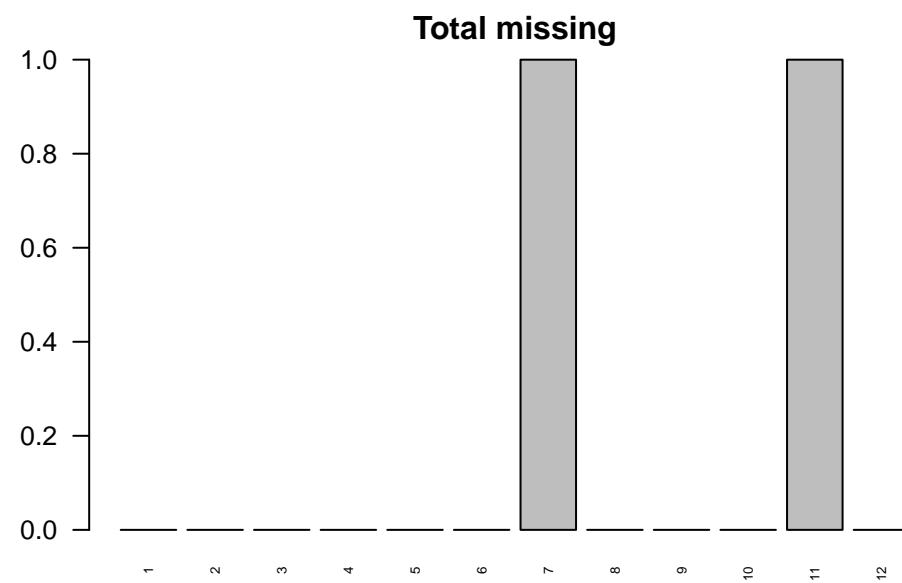
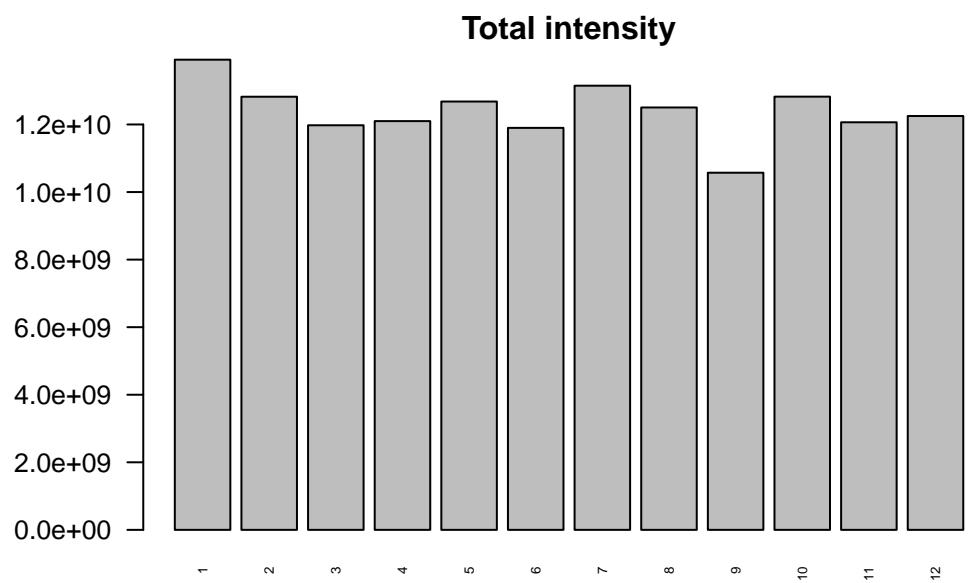
Report created on: 2023-05-27

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

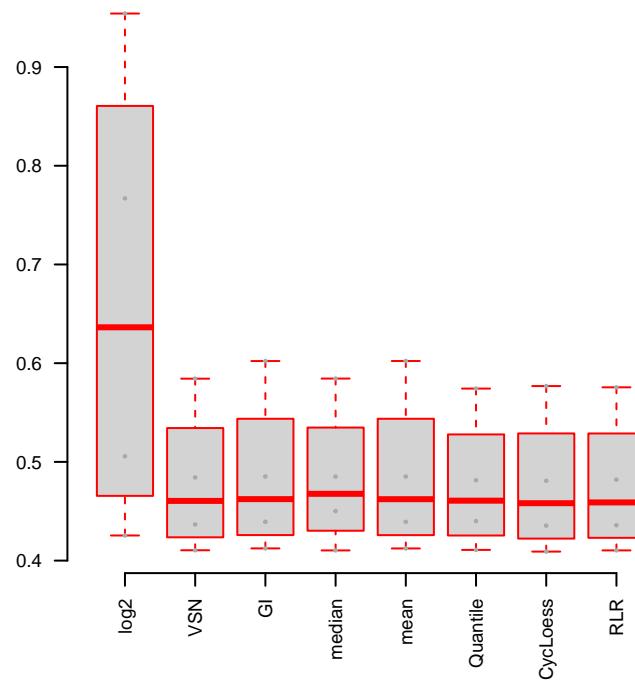
Documentation for analyzing this report can be found at <http://quantitativeproteomics.org/normalizer/help.php>

Group nbr.	Design group	Nbr. samples in cond.
1	At.risk	3
2	Control	3
3	HC.like	3
4	SPD	3

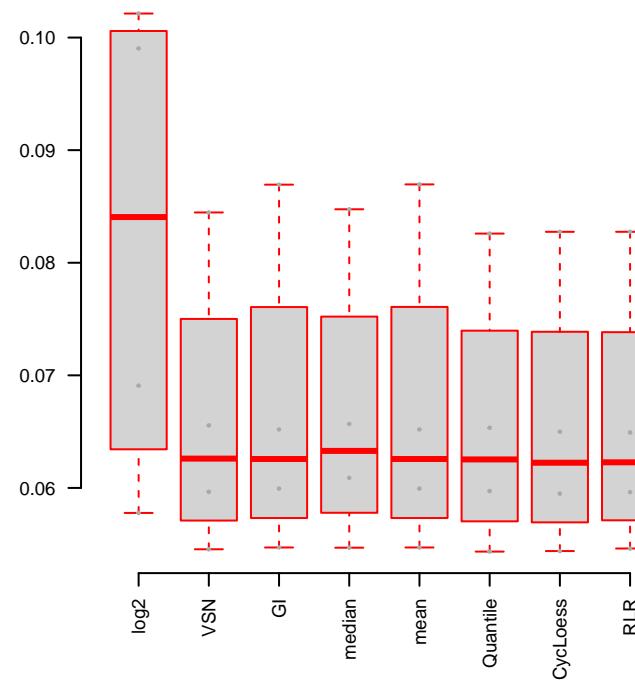
Please note that the grouping only impacts evaluation measures and visuals seen in the report, they do not impact the performed normalizations.



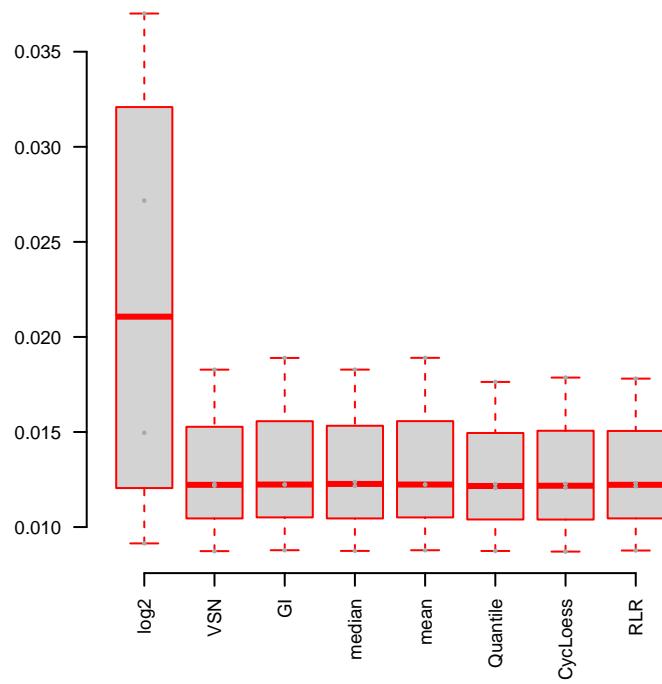
PCV – Intragroup



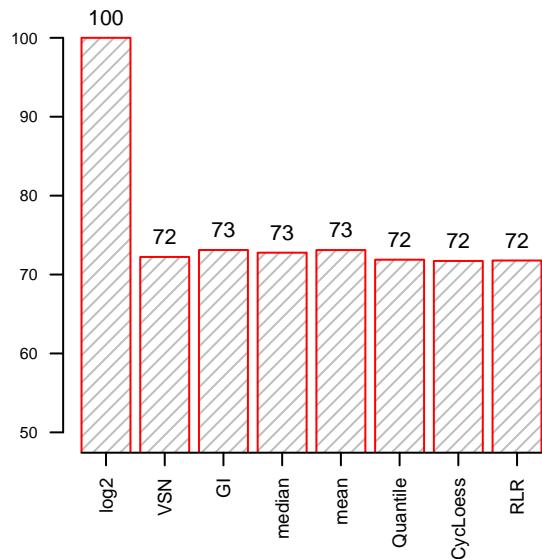
PMAD – Intragroup



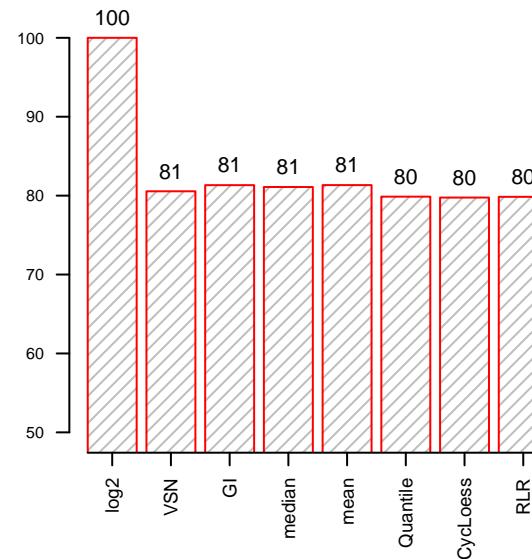
PEV – Intragroup



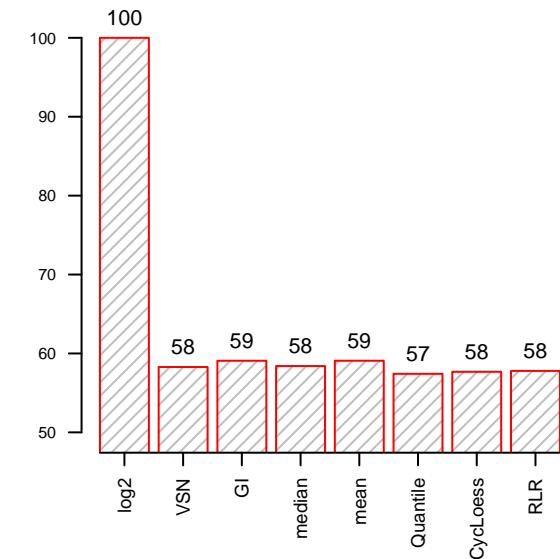
PCV compared to log2



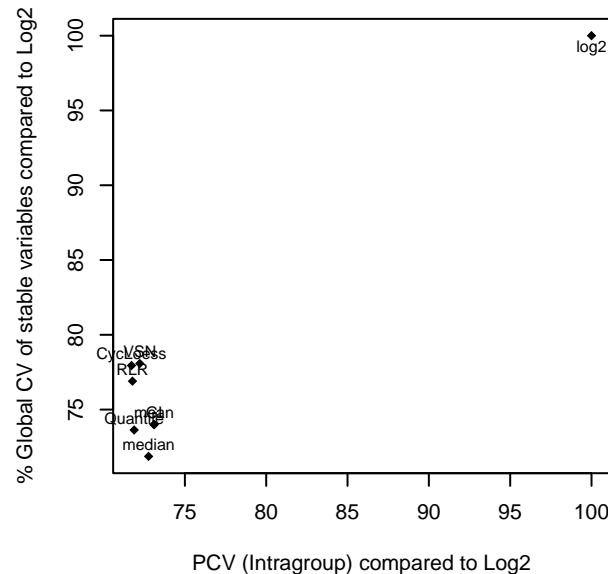
PMAD compared to log2

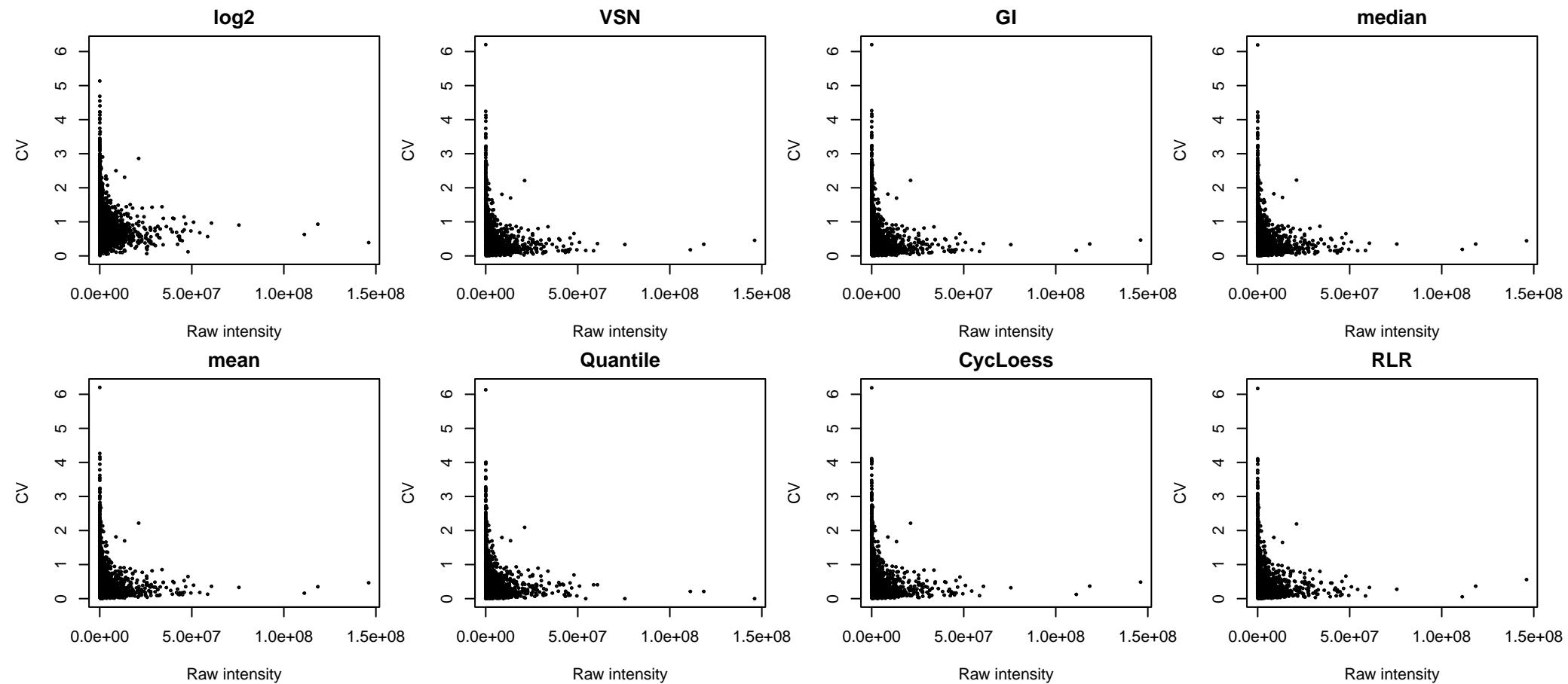


%PEV – compared to log2



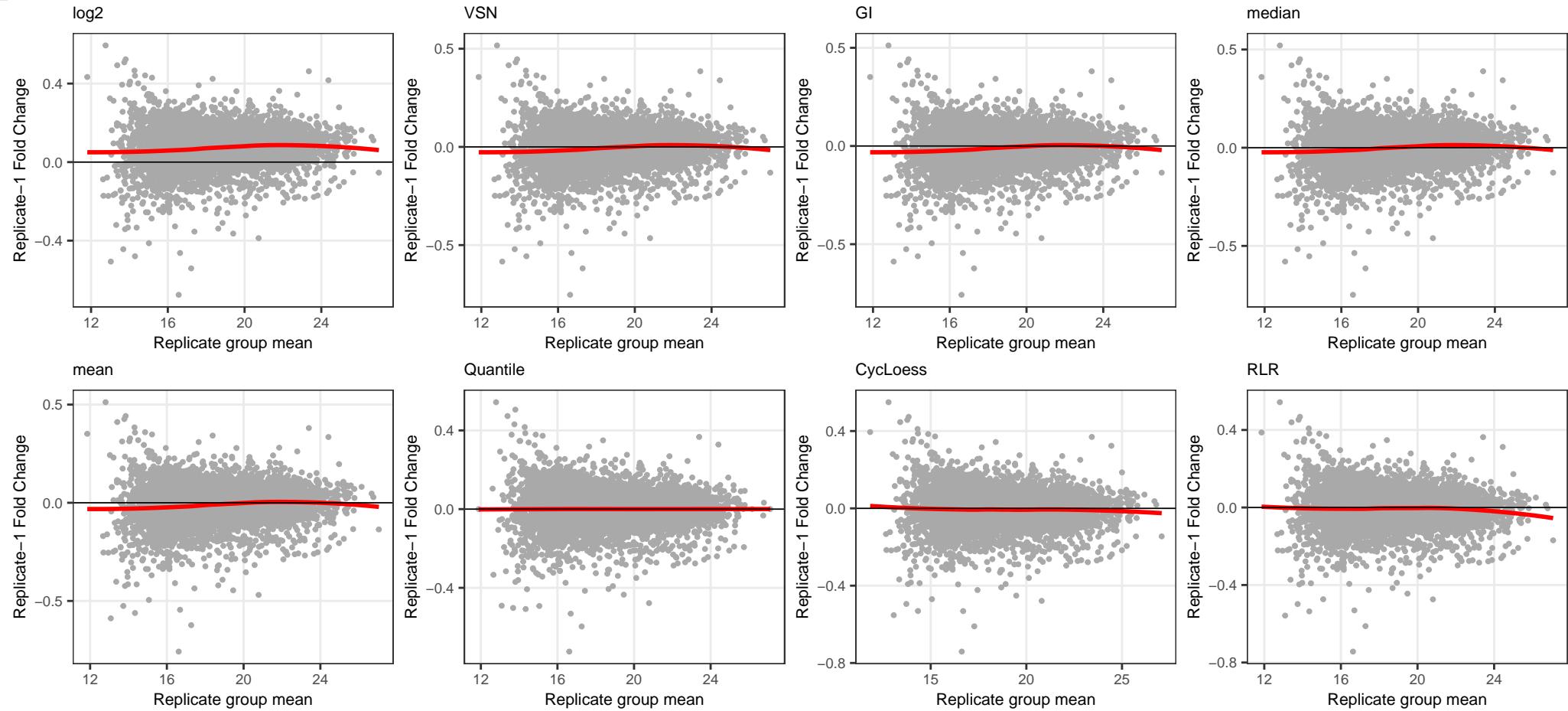
Stable variables plot

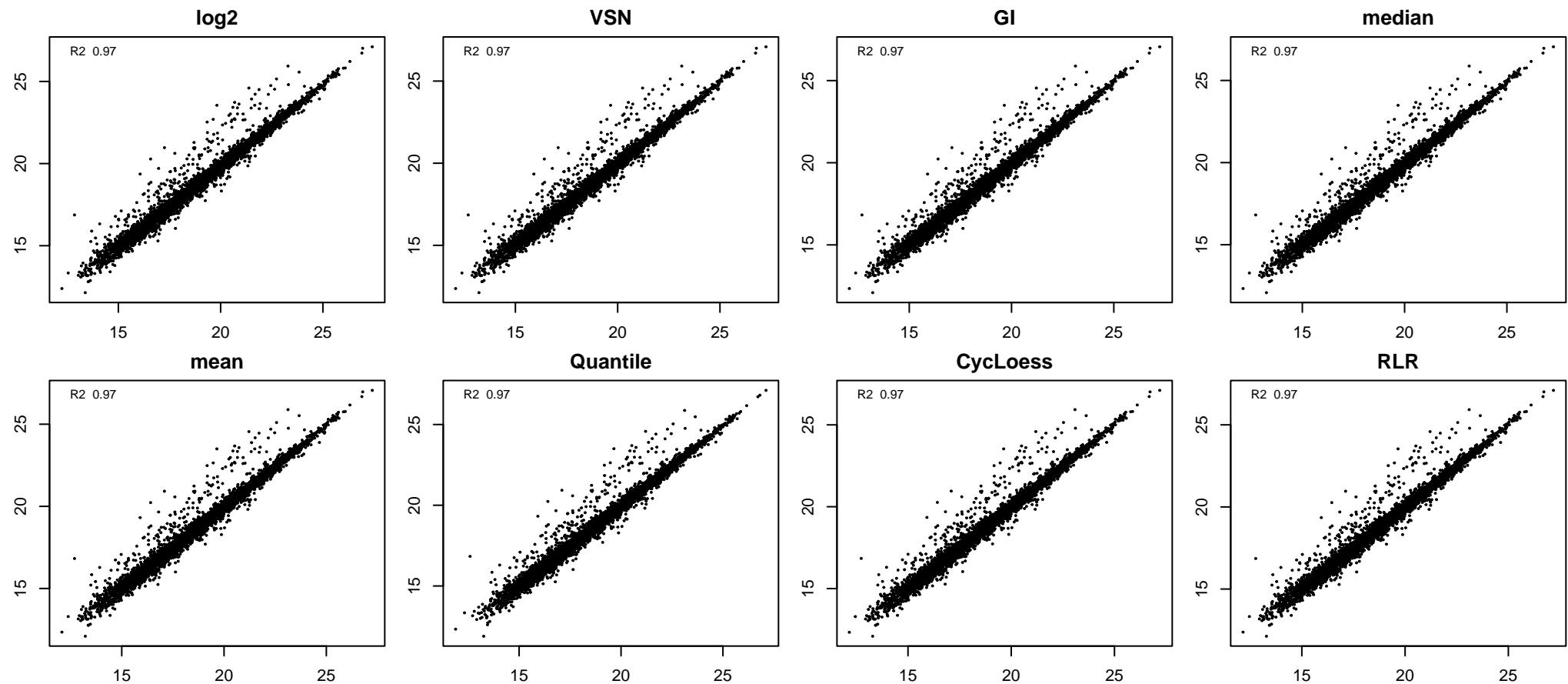




MA plots

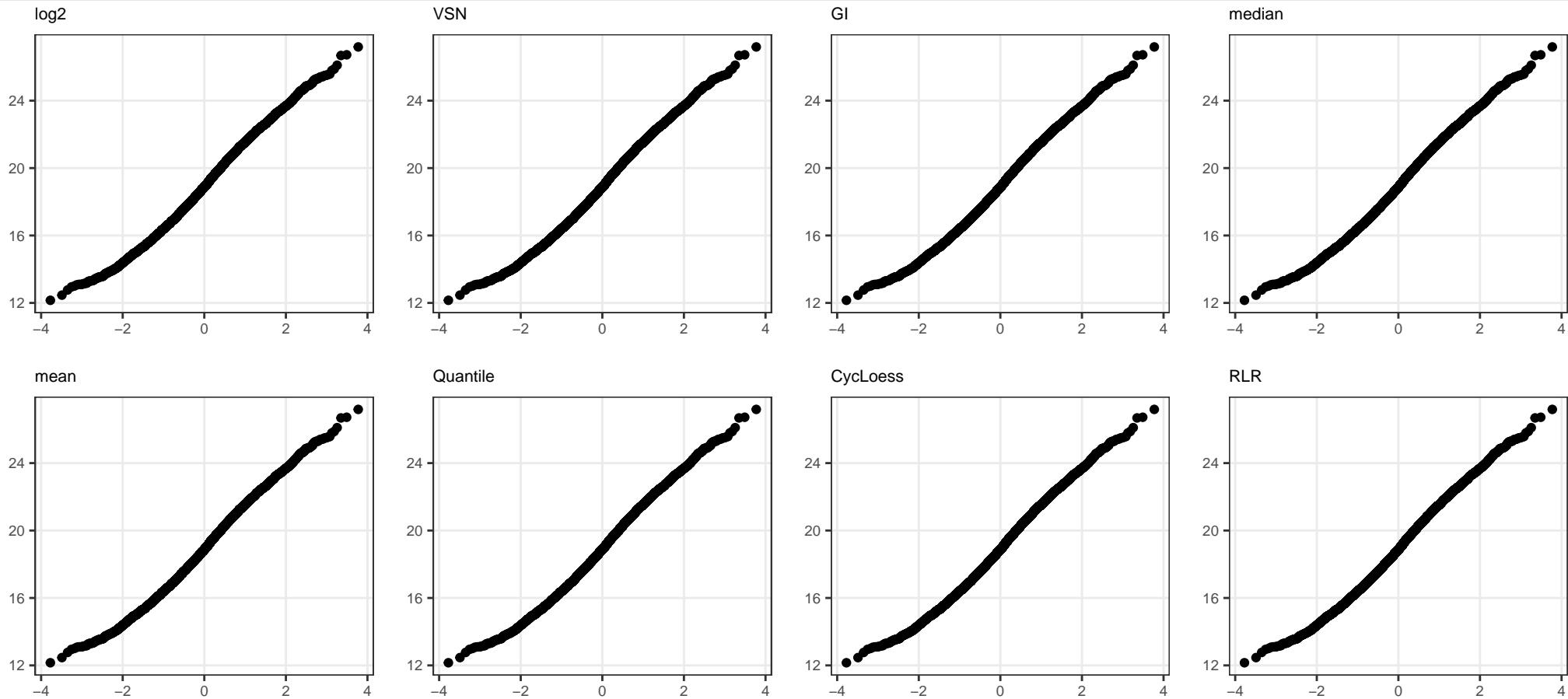
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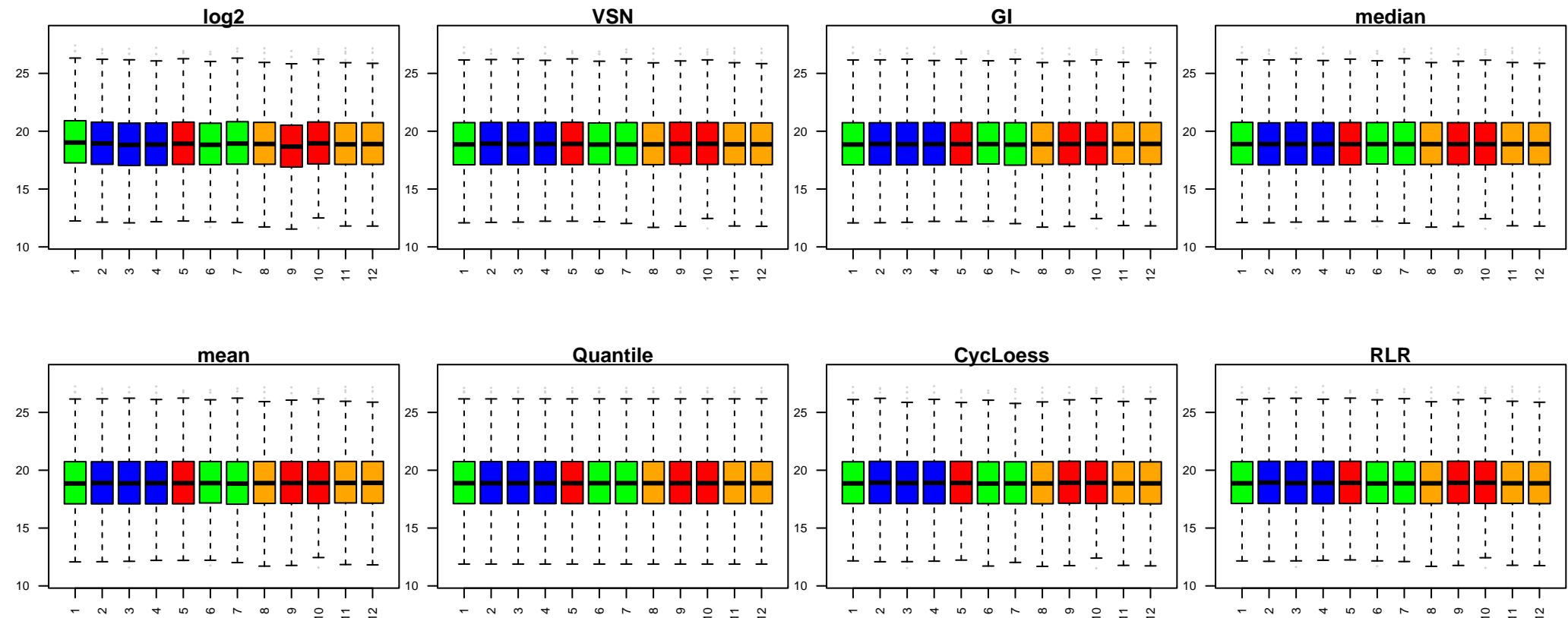


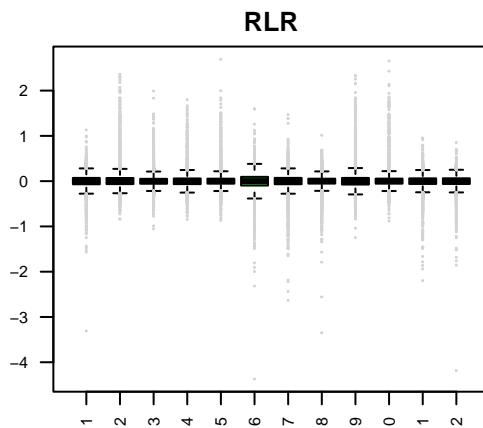
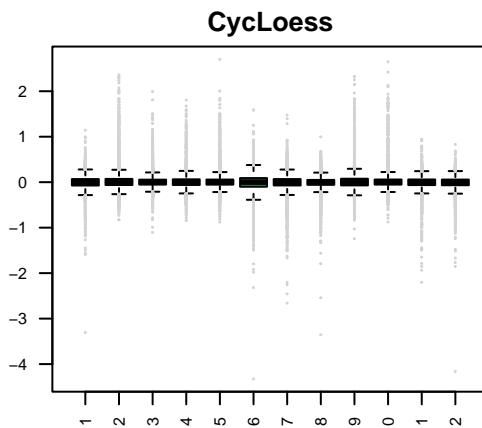
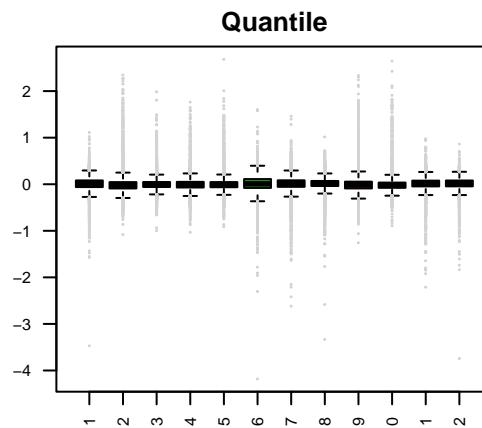
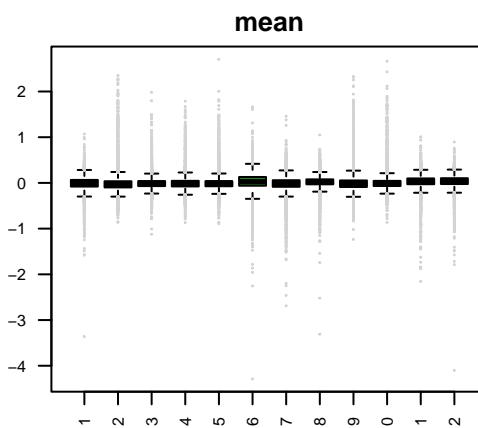
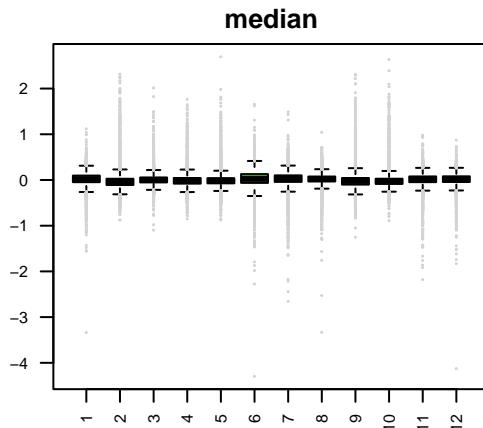
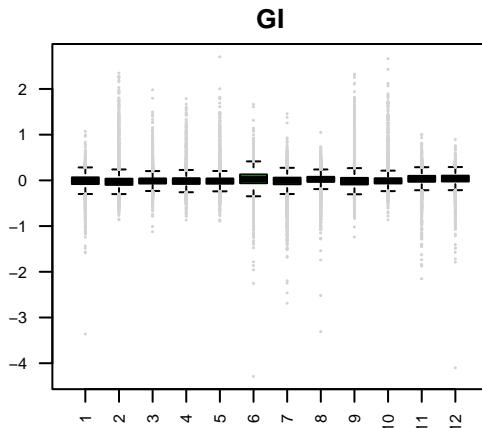
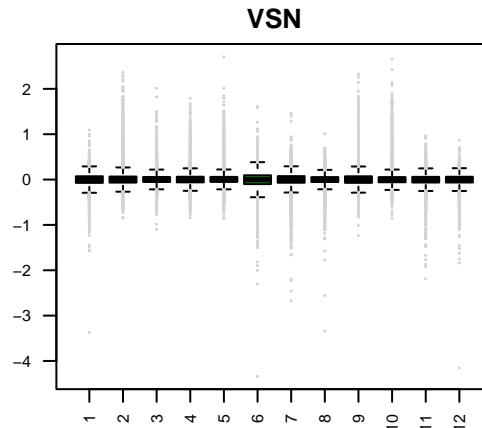
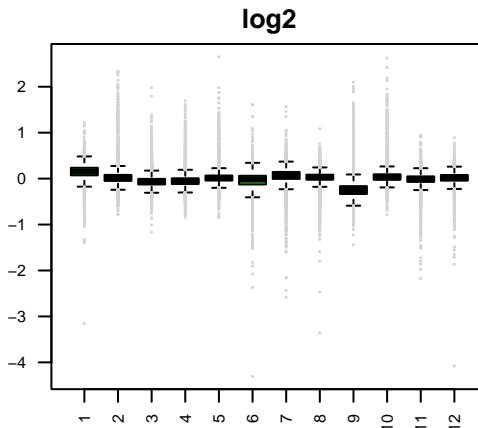


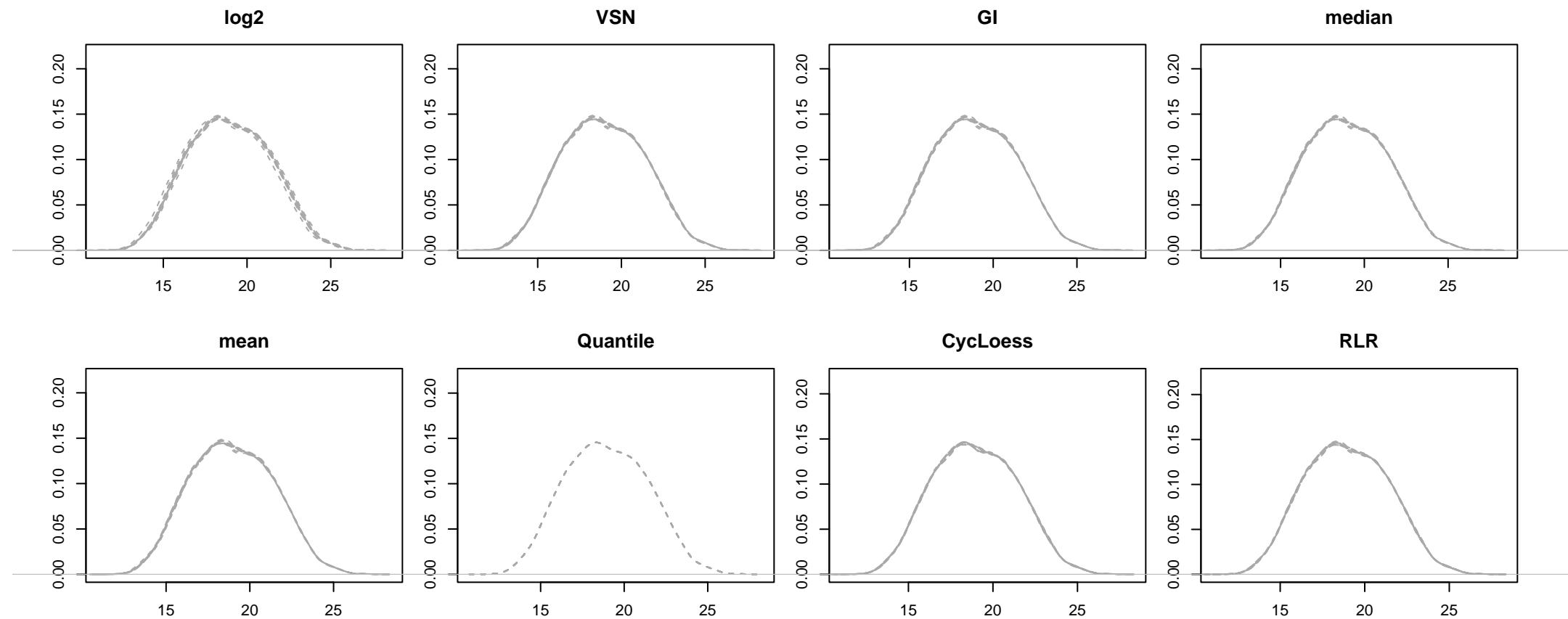
Q-Q plots

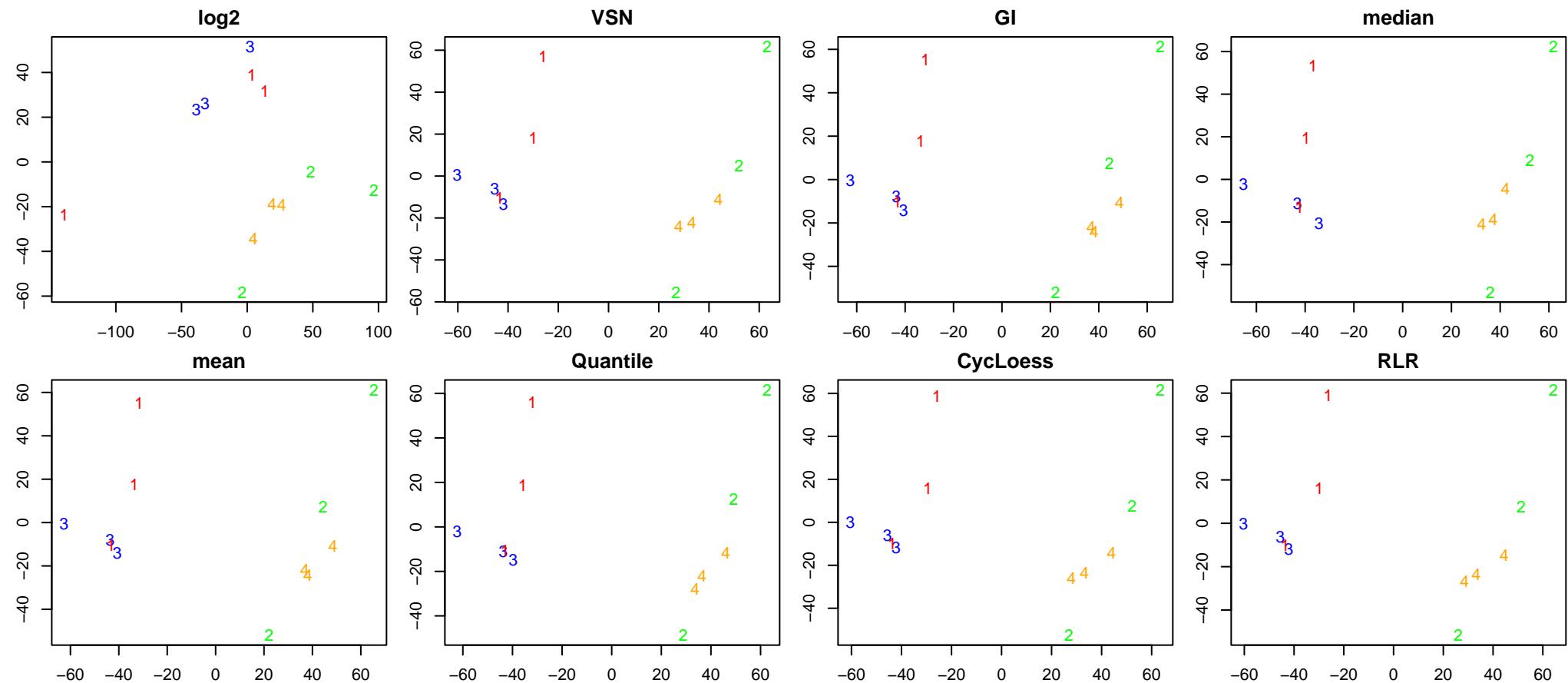
NormalizerDE Report

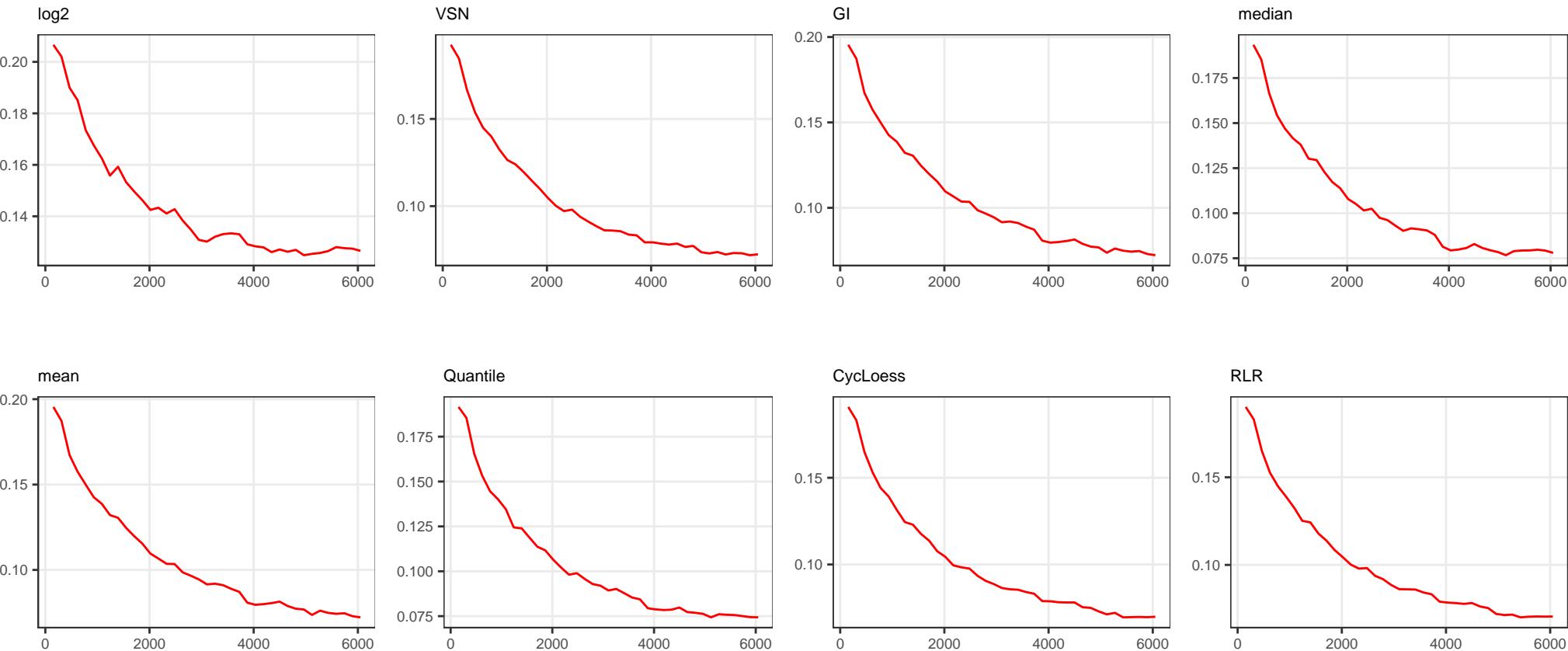




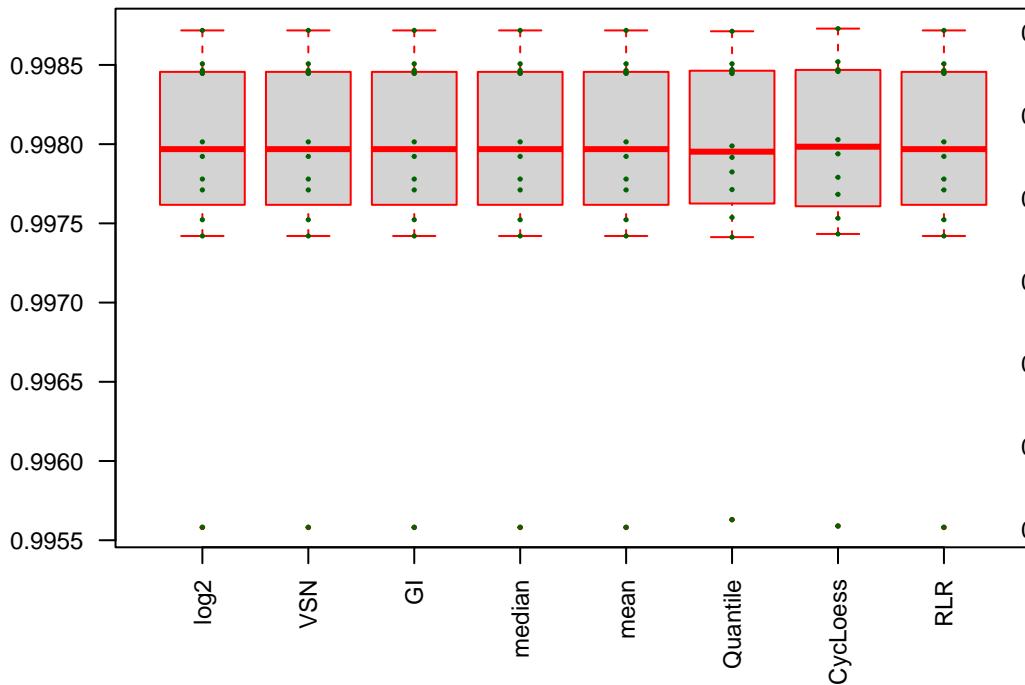








Pearson correlation – Intragroup



Spearman correlation – Intragroup

