

Project Name: Monocytes\_cocomo\_single

# NormalyzerDE (ver 1.14.0 )

Report created on: 2023-02-09

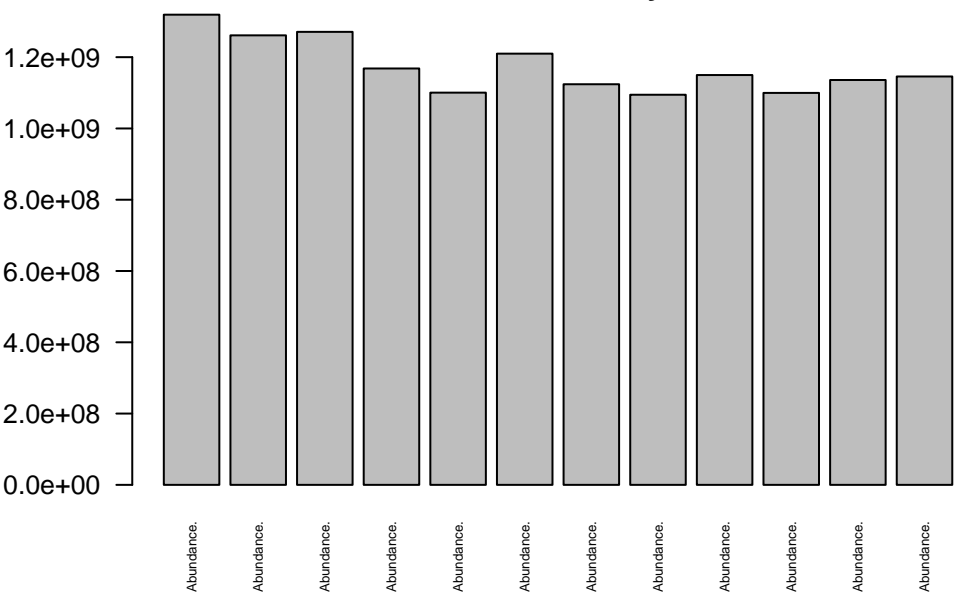
Citation: NormalyzerDE: Online Tool for Improved Normalization of Omics Expression Data and High-Sensitivity Differential Expression Analysis  
Journal of Proteome Research (2018), 10.1021/acs.jproteome.8b00523

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

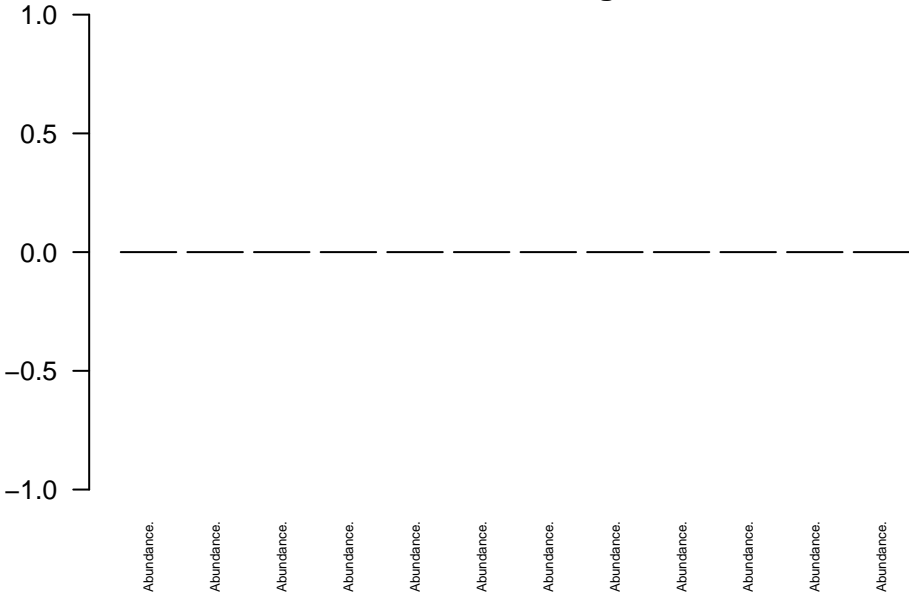
Group nbr.	Design group	Nbr. samples in cond.
1	1A	1
2	1B	1
3	2A	1
4	2B	1
5	3A	1
6	3B	1
7	4A	1
8	4B	1
9	5A	1
10	5B	1
11	6A	1
12	6B	1

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

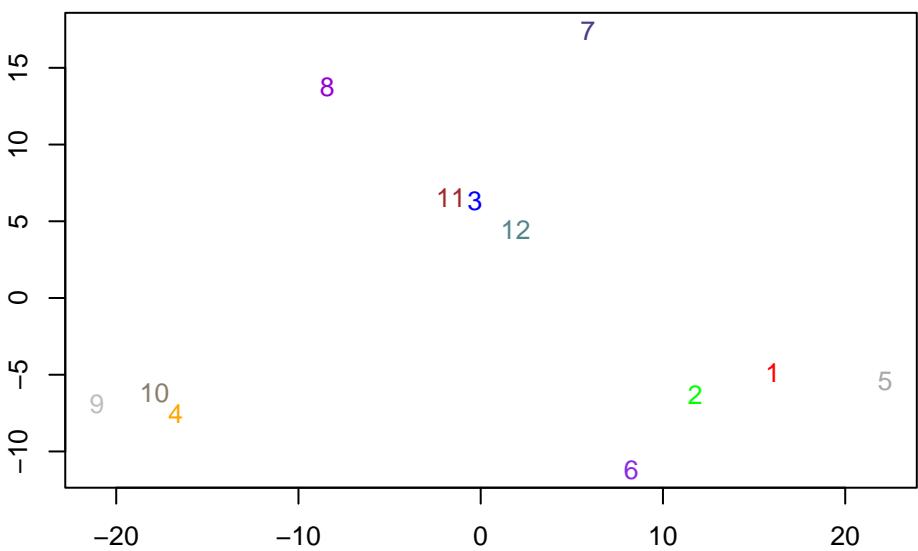
Total intensity

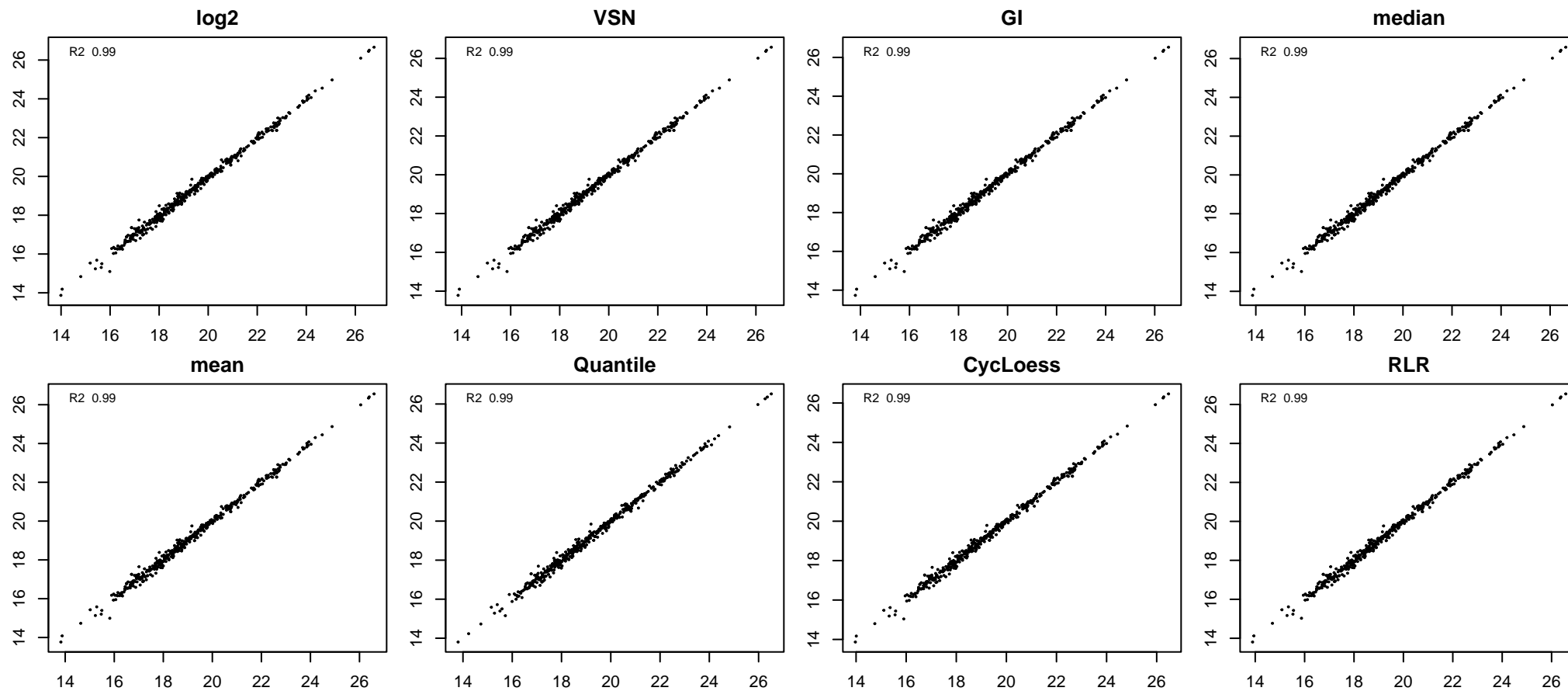


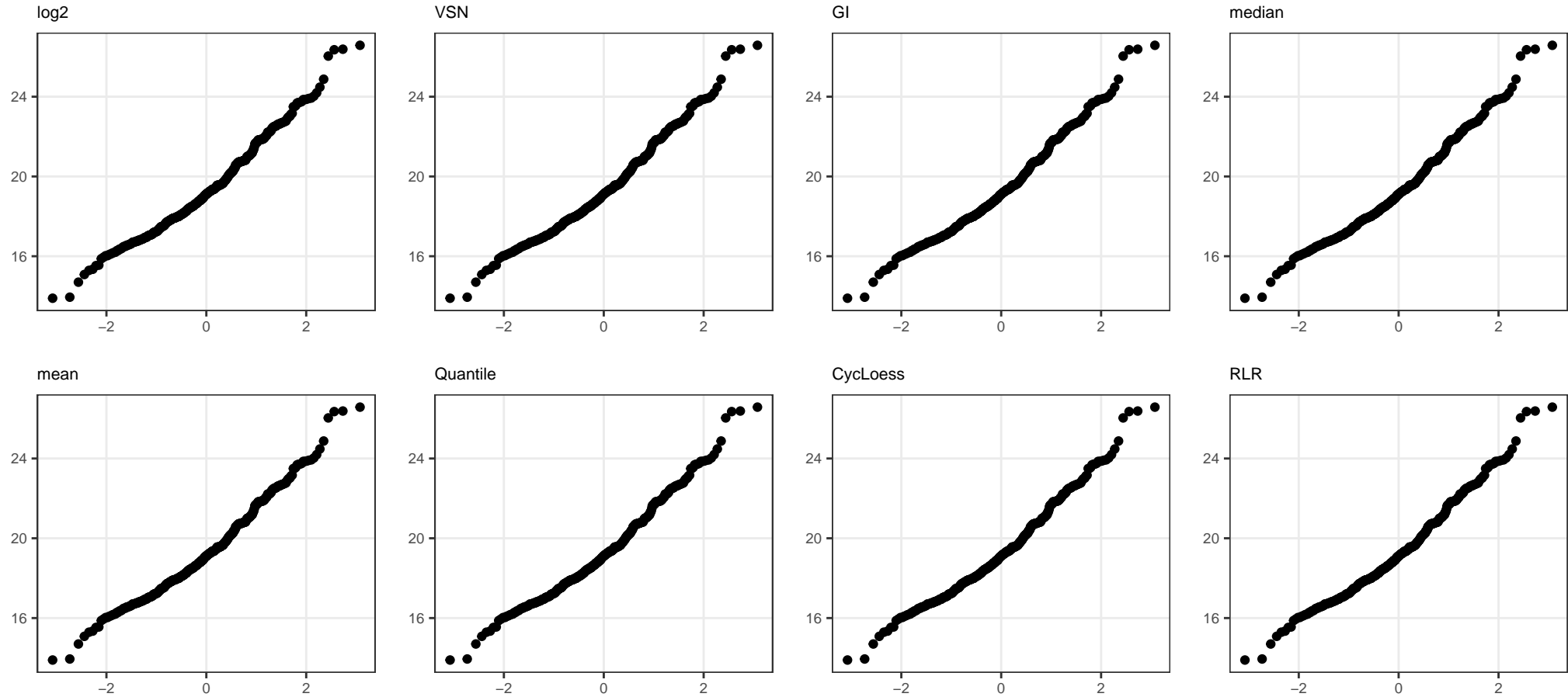
Total missing

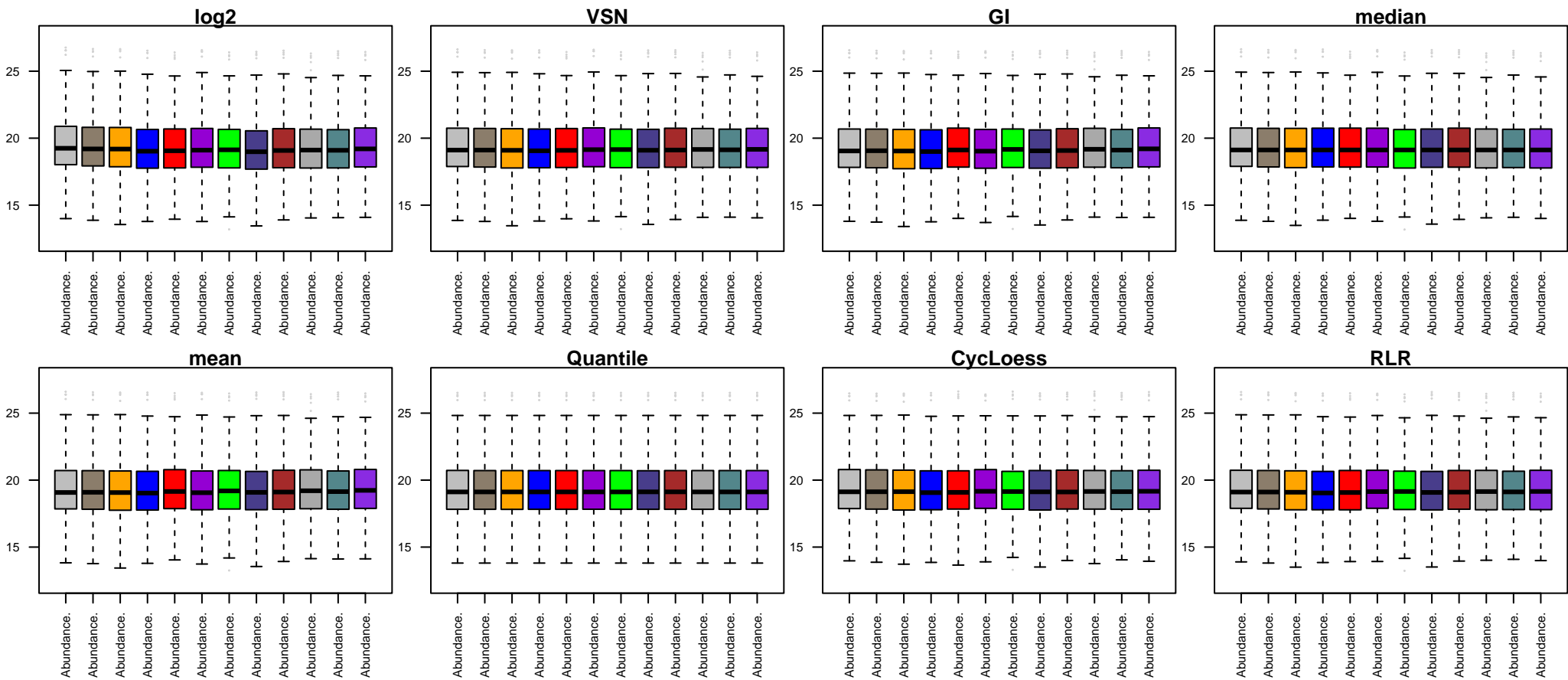


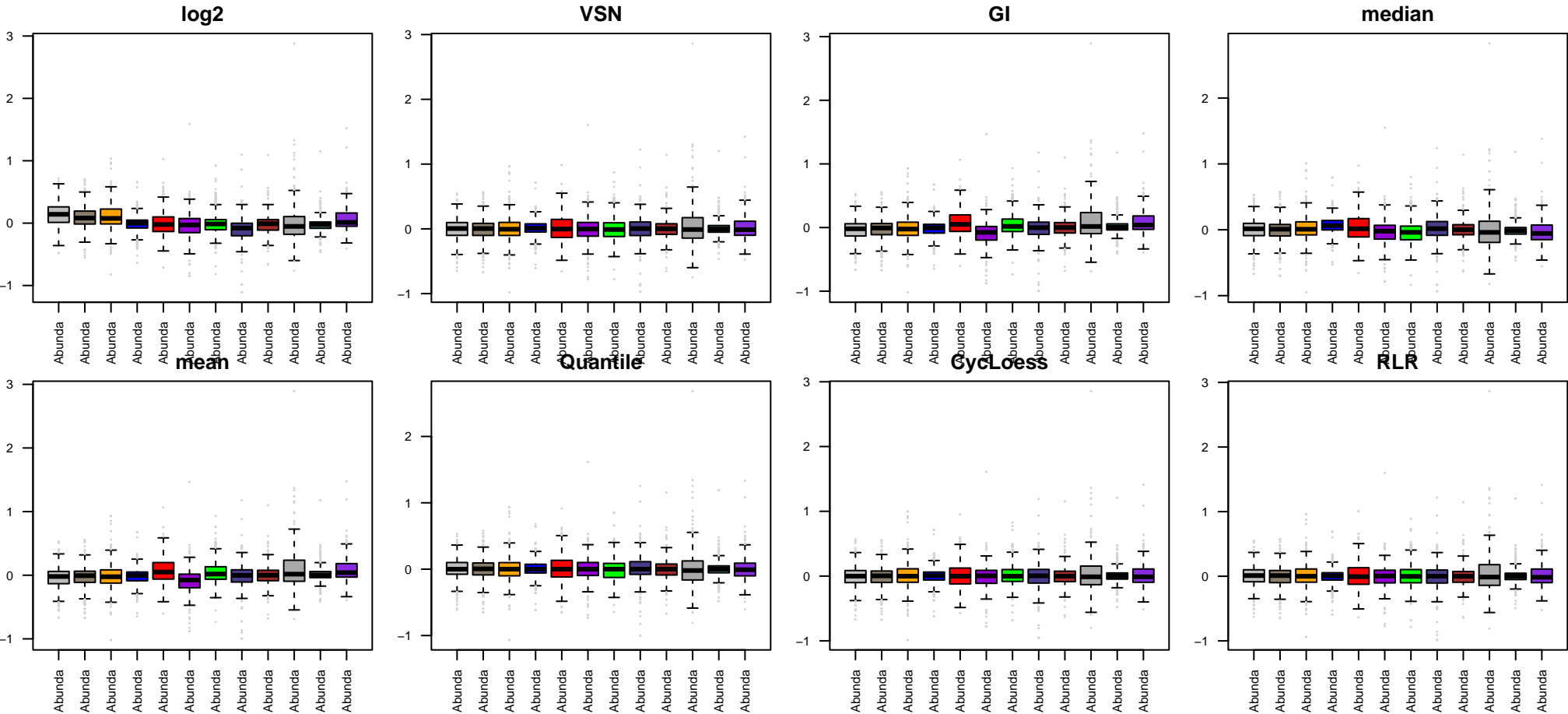
Log2-MDS plot



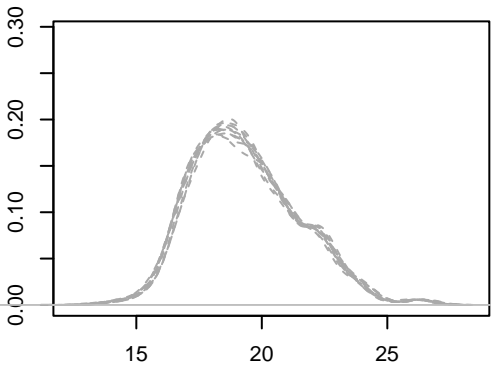




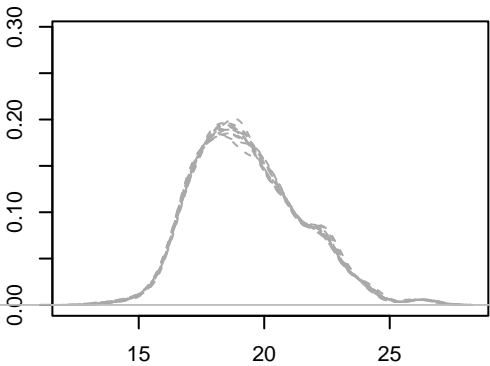




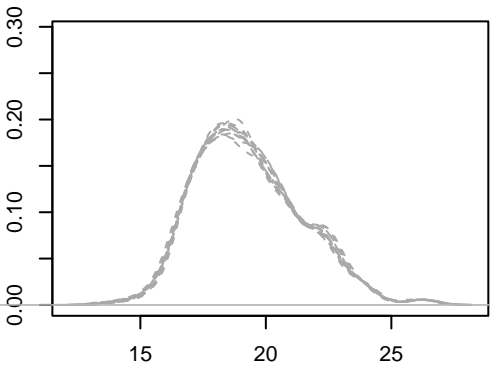
log2



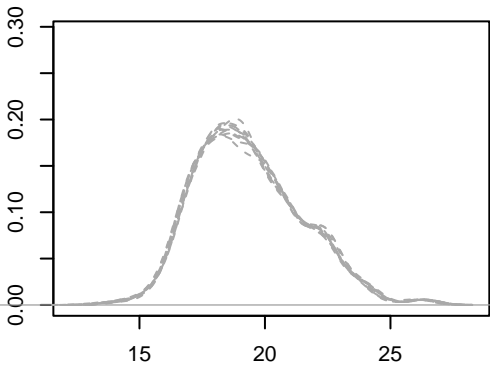
VSN



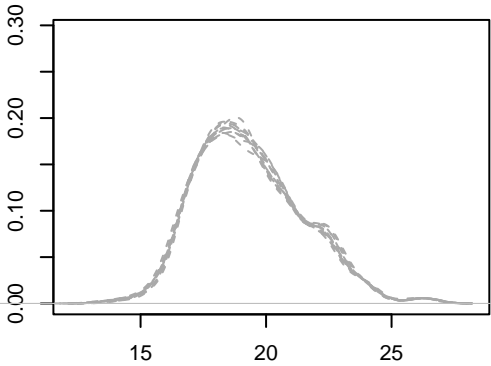
GI



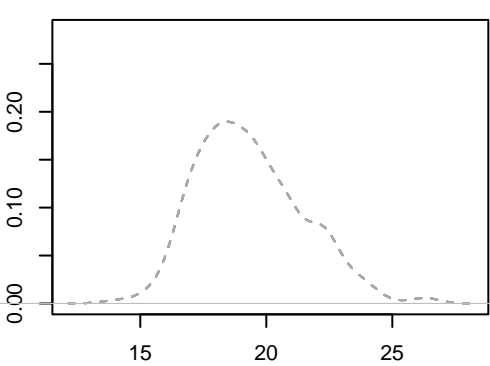
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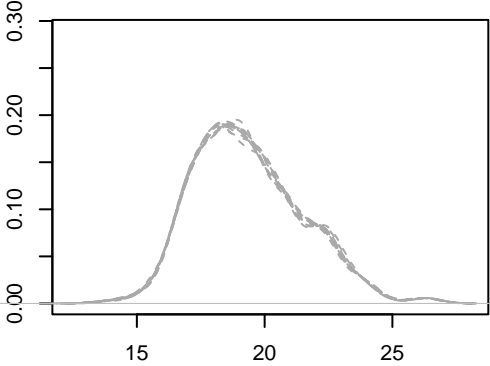
mean



Quantile



CycLoess



RLR

