

Project Name: 2024-04-16\_msp-ms\_normalize\_output

# NormalizerDE (ver 1.19.7 )

Report created on: 2024-04-16

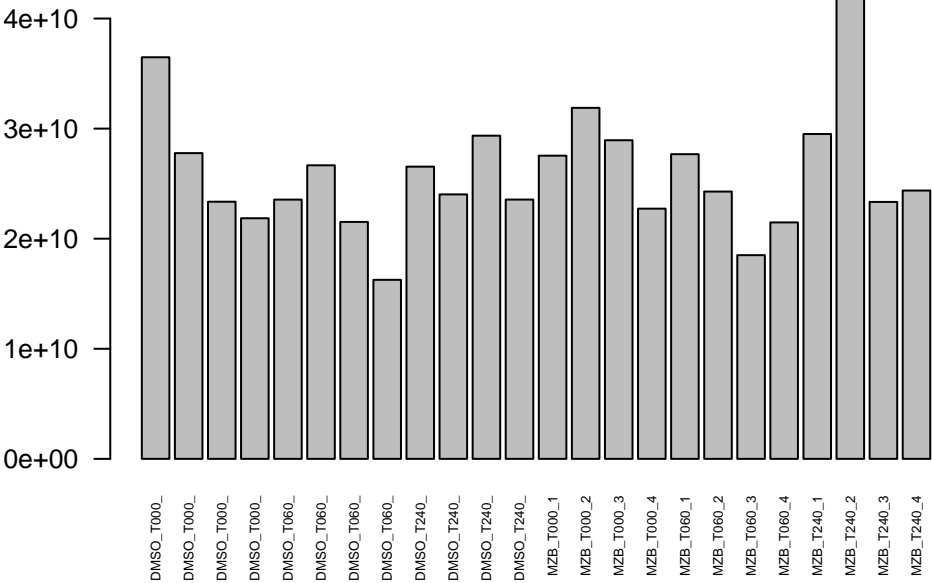
Citation: NormalizerDE: Online Tool for Improved Normalization of Omics Expression Data and High-Sensitivity Differential Expression Analysis  
Journal of Proteome Research (2018), 10.1021/acs.jproteome.8b00523

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

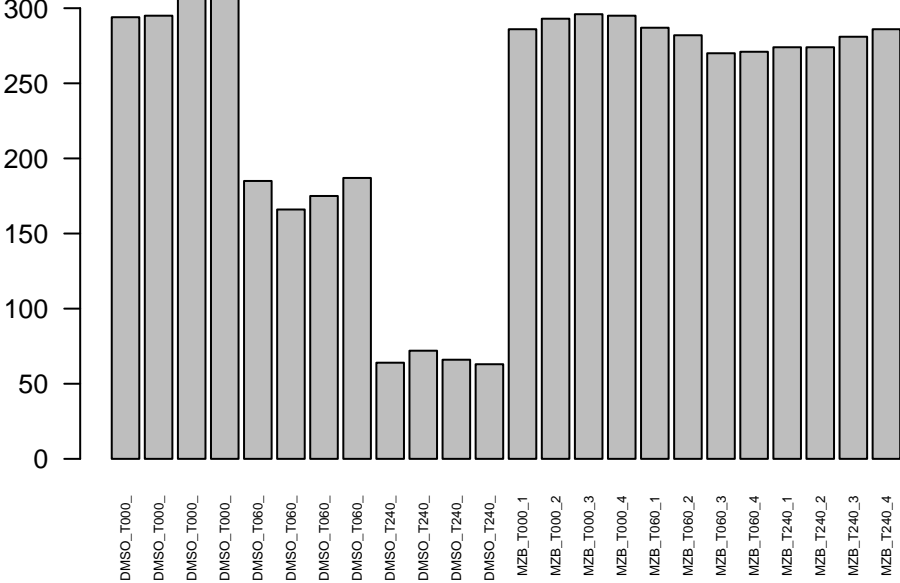
Group nbr.	Design group	Nbr. samples in cond.
1	DMSO_T0	4
2	DMSO_T240	4
3	DMSO_T60	4
4	MZB_T0	4
5	MZB_T240	4
6	MZB_T60	4

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

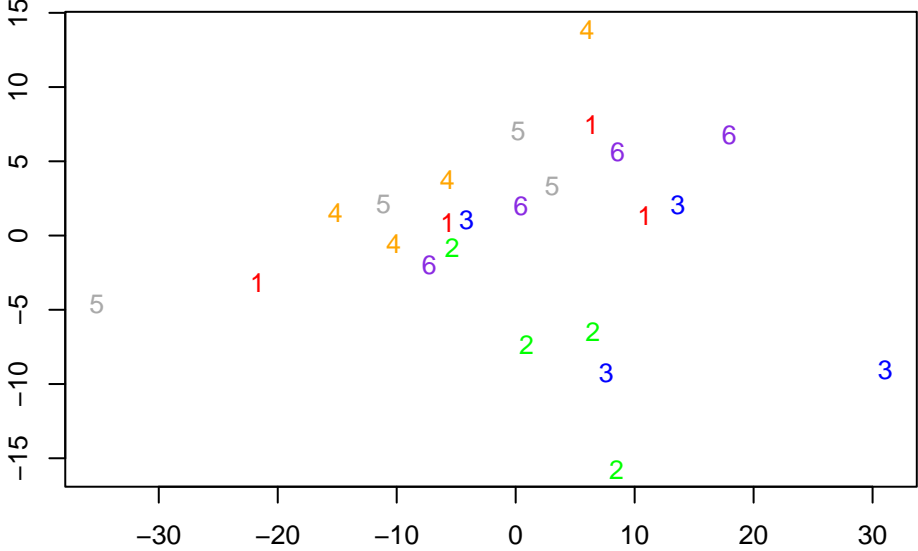
Total intensity

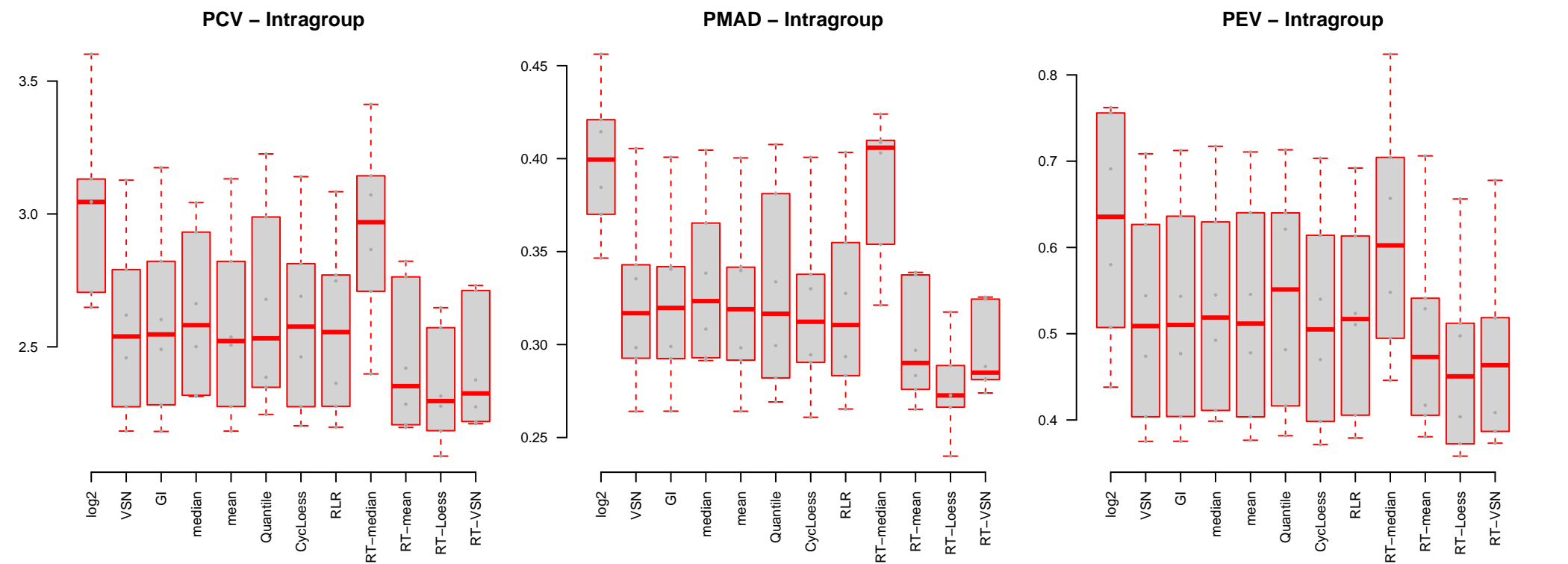


Total missing

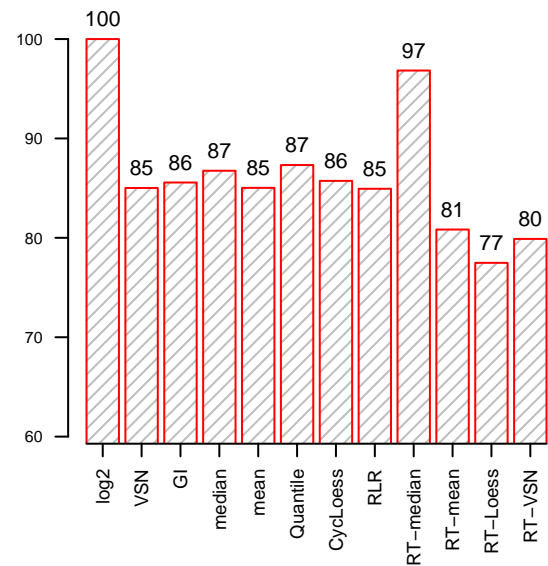


Log2-MDS plot

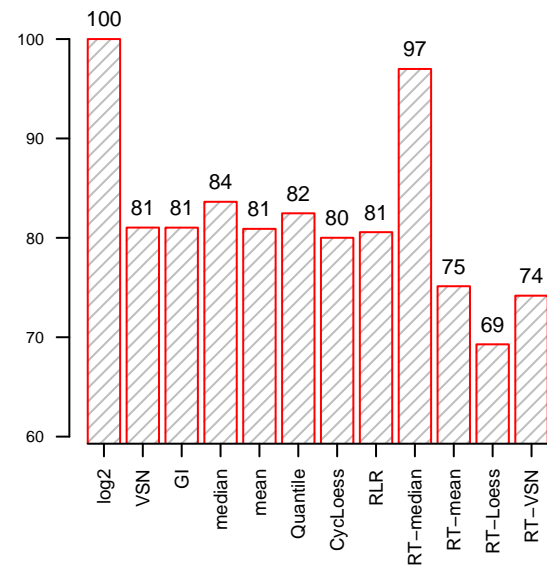




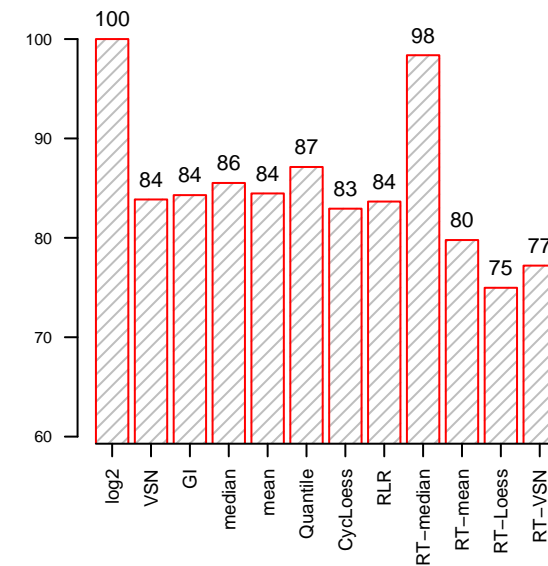
PCV compared to log2



PMAD compared to log2



%PEV – compared to log2



Stable variables plot

