Project Name: con\_sp.hsa.canonical.isoforms\_junc.exons\_peptides

## Normalyzer (ver 1.4.0)

Report created on: 2020-05-07

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

**Total missing** 

1h\_vs\_ud|I

l|pn\_sv\_d9

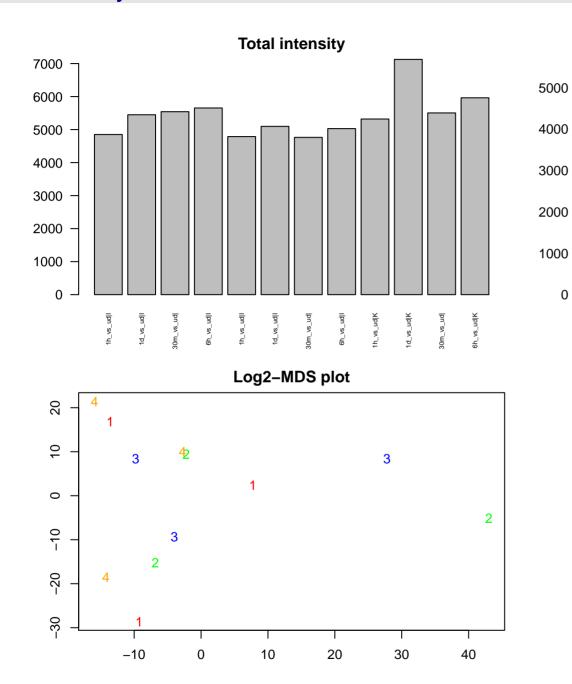
1h\_vs\_ud|I

1d\_vs\_ud||

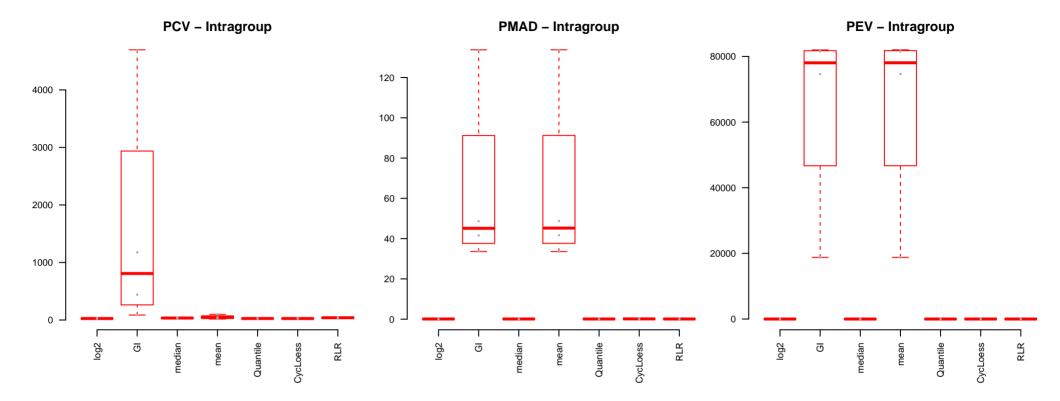
30m\_vs\_ud

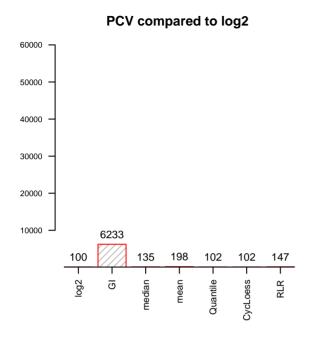
6h\_vs\_ud||

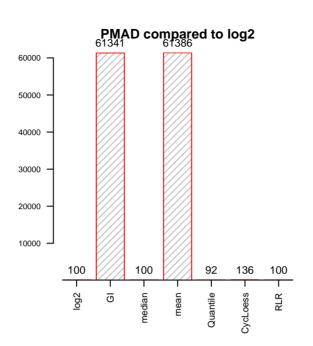
1h\_vs\_ud|K

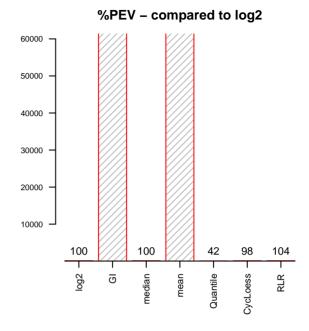


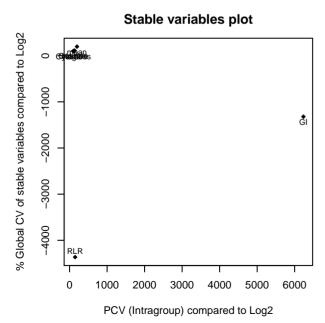


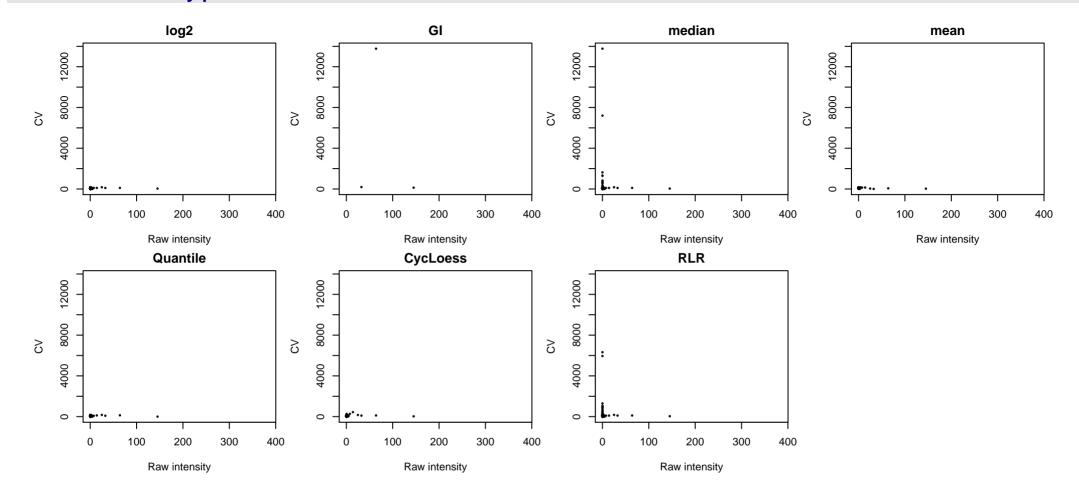


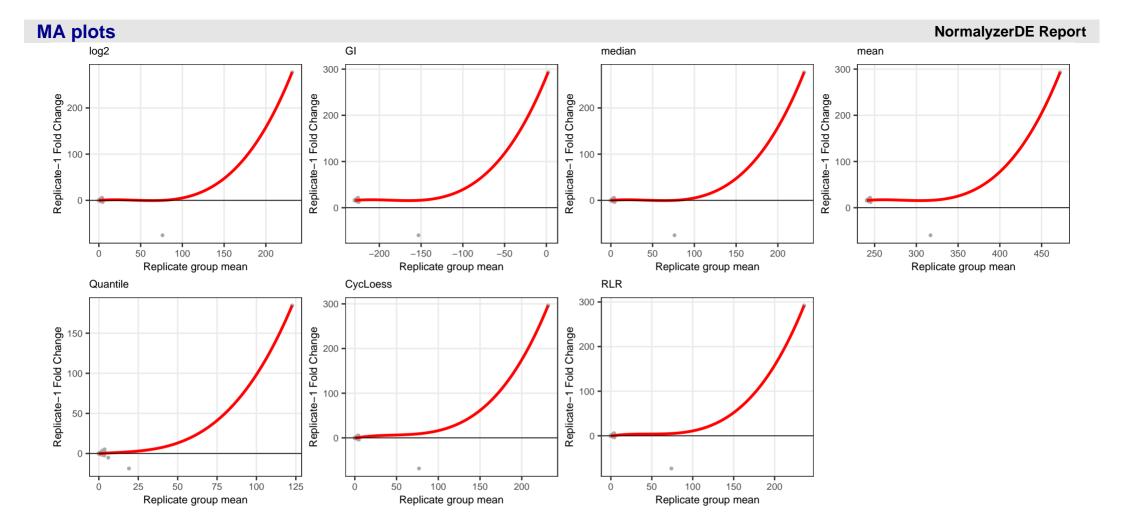




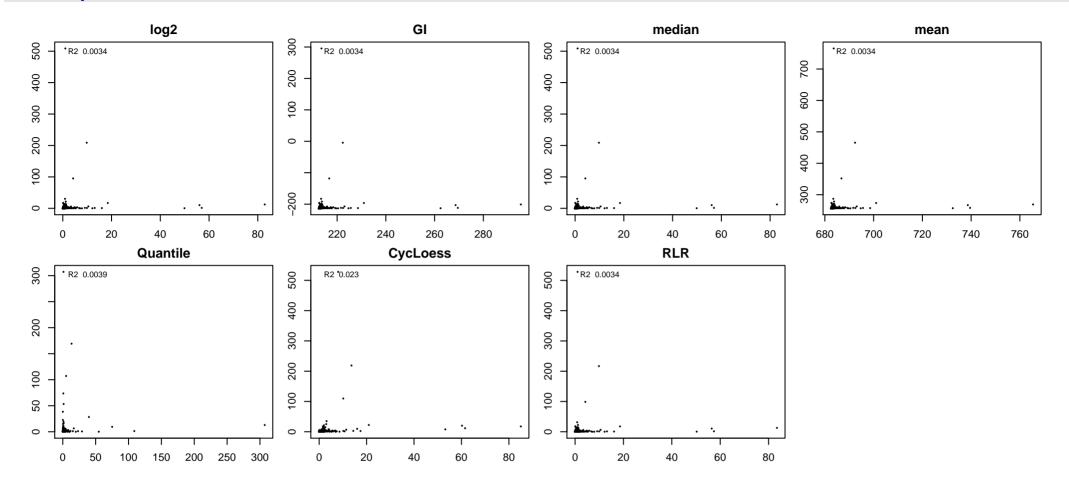


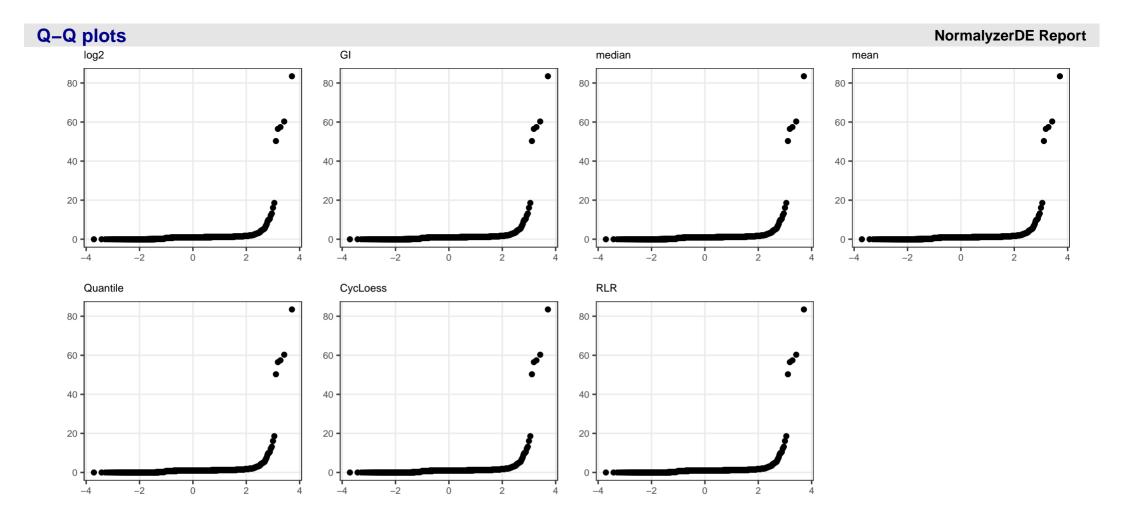




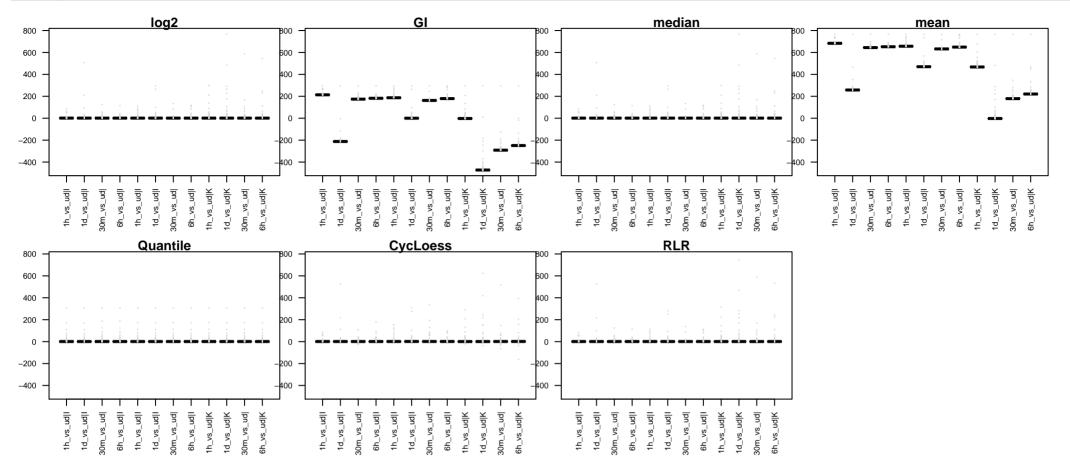


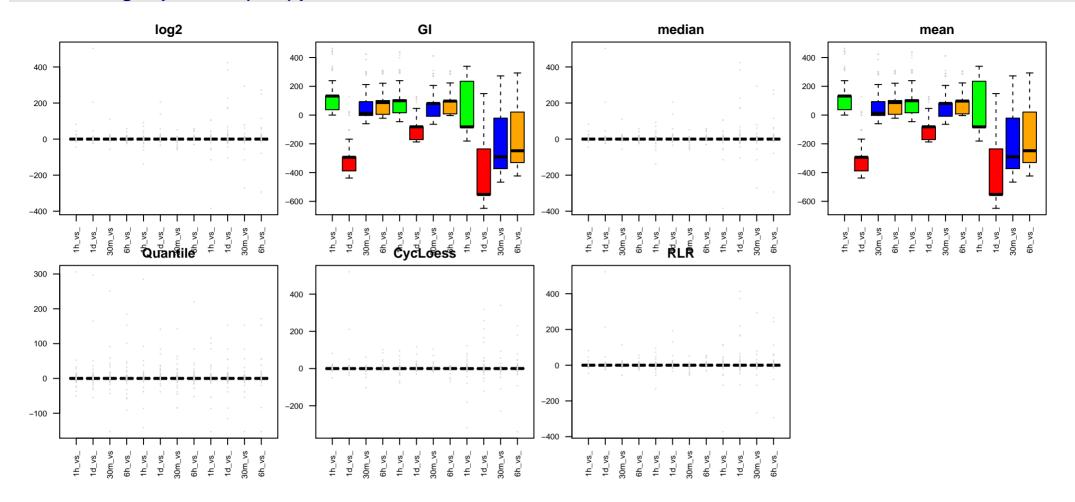
**Scatterplots** 

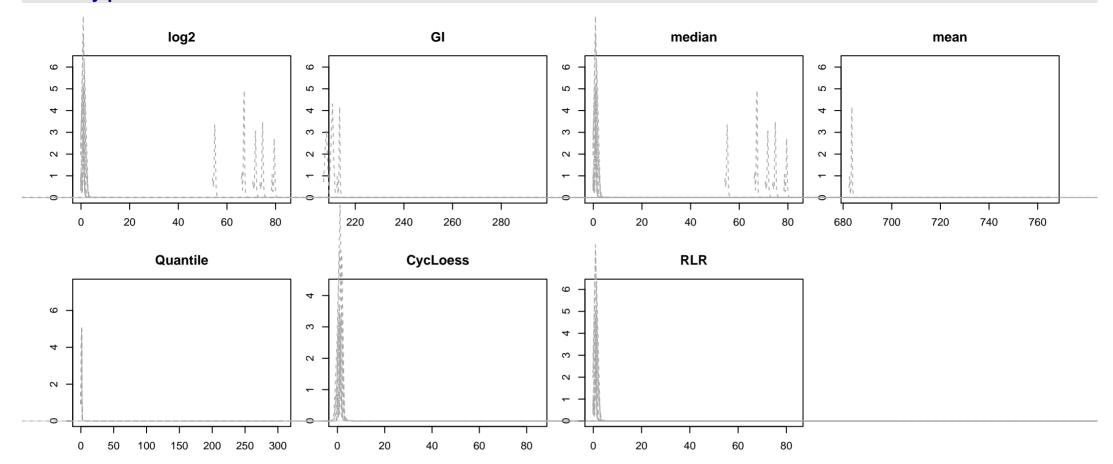


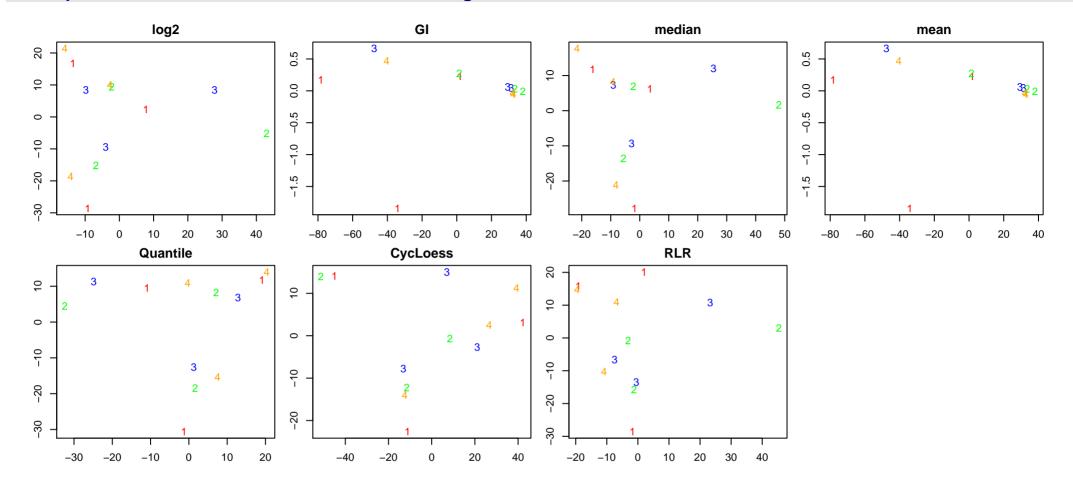


Boxplots Normalyzer Report

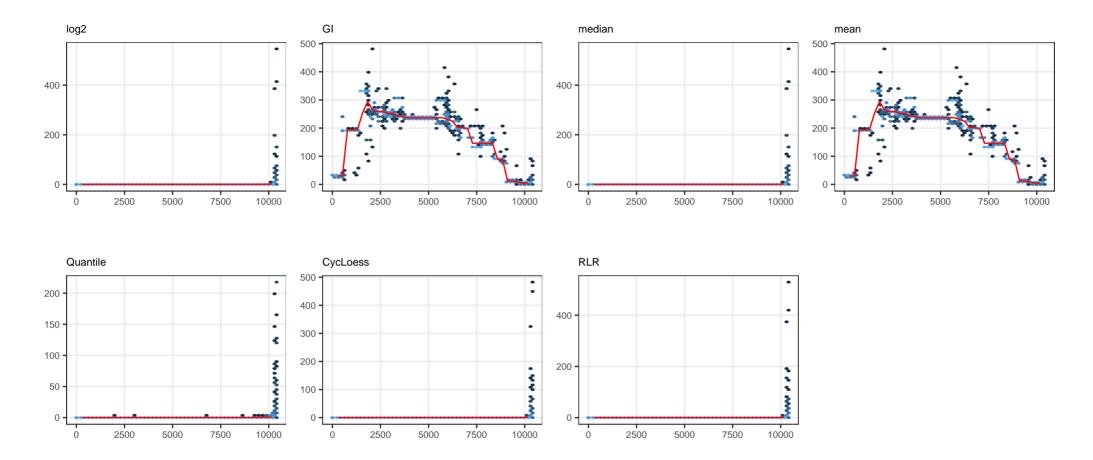








**MeanSDplots** 



**Correlation plots** 

