Exhaustive and Reliable Data Analysis for Untargeted Metabolomics

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From Raw Data to Compound List

Use high quality expert curated data for your downstream statistical analysis!



Retention Time (min)

Example Analysis

• The Association of Biomolecular Resource Facilities (ABRF) Metabolomics Research Group (MRG) 2016 Study

https://abrf.org/research-group/metabolomics-research-group-mrg https://abrf.org/sites/default/files/2016_mrg_study_outline_060716.pdf

- Urine samples from mice exposed to 5 Gray of external X-ray (n = 5) and those exposed to sham irradiation control group (n = 5)
- Each sample analyzed twice in both positive and negative polarity modes (UPLC-Q-TOF)
- Pooled sample analyzed in MS^{*E*} mode in both positive and negative polarity
- Authors provided raw data as well as peak lists from automated analysis
- The assignment: "...report on the top 50 features that show statistically significant differences in relative abundances between the two study groups..."

Compound Interpretation



Compounds with Significant Differences

None of the peaks below were found in peak lists provided by the authors of the study.



Thank You!

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Additional References

Untargeted Metabolomics Suffers from Incomplete Data Analysis

https://doi.org/10.1101/143818

• Unexpected Metabolites

http://pubs.acs.org/doi/abs/10.1021/ac1020112 http://www.mdpi.com/1660-3397/11/10/3617/htm

• Unexpected Metabolic Capabilities

http://dx.doi.org/10.1039/c1mb05196b

Metabolic Interactions

http://www.nature.com/articles/ncomms9289

• Interpretation of Mass Spectra

http://pubs.acs.org/doi/abs/10.1021/ac402180c

• Metabolite Turnover

http://pubs.acs.org/doi/abs/10.1021/acschembio.6b00890

• Enzyme Discovery

http://pubs.acs.org/doi/abs/10.1021/pr0600576 http://pubs.acs.org/doi/abs/10.1021/cb300477w

• Biomarker Discovery

http://www.jbc.org/content/281/24/16768.full