

Project Name: NormalyzerPeptides

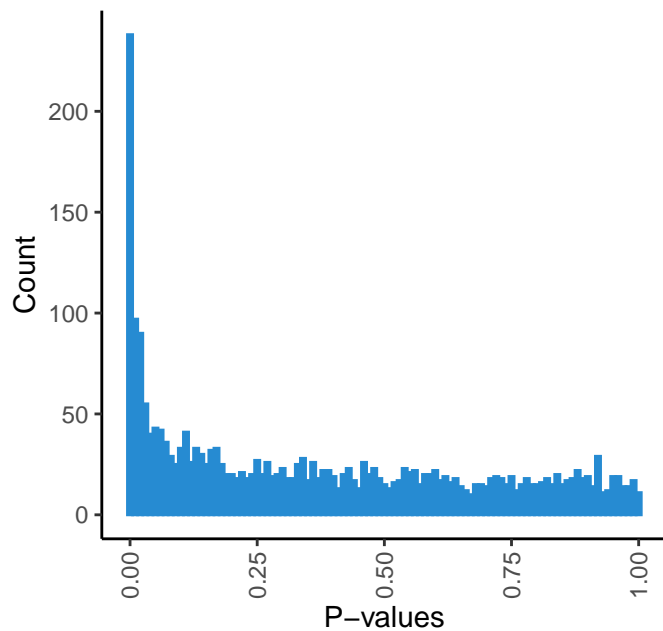
NormalyzerDE (ver 1.14.0)

Report created on: 2022-06-28

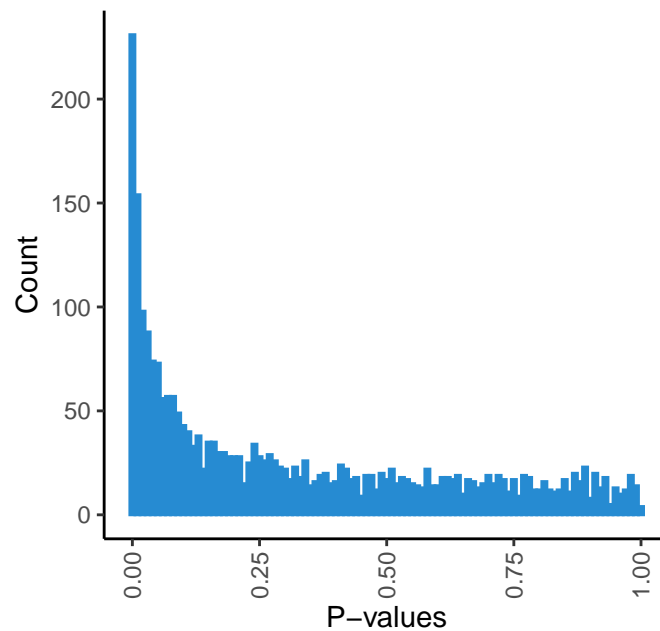
Citation: NormalyzerDE: Online Tool for Improved Normalization of Omics Expression Data and High-Sensitivity Differential Expression Analysis
Journal of Proteome Research (2018), 10.1021/acs.jproteome.8b00523

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

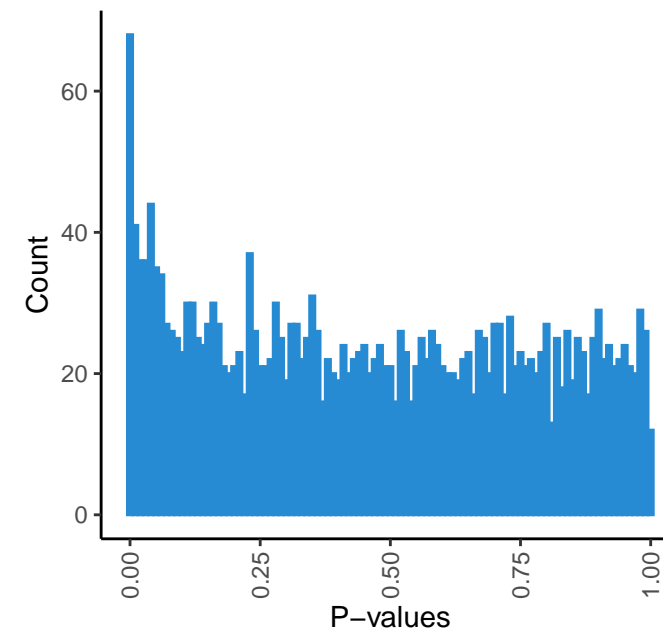
Contrast: 125.amol–12500.amol



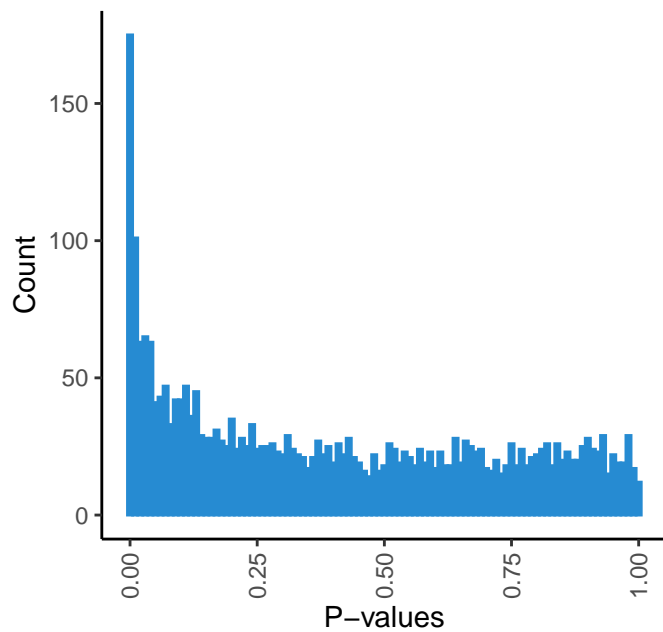
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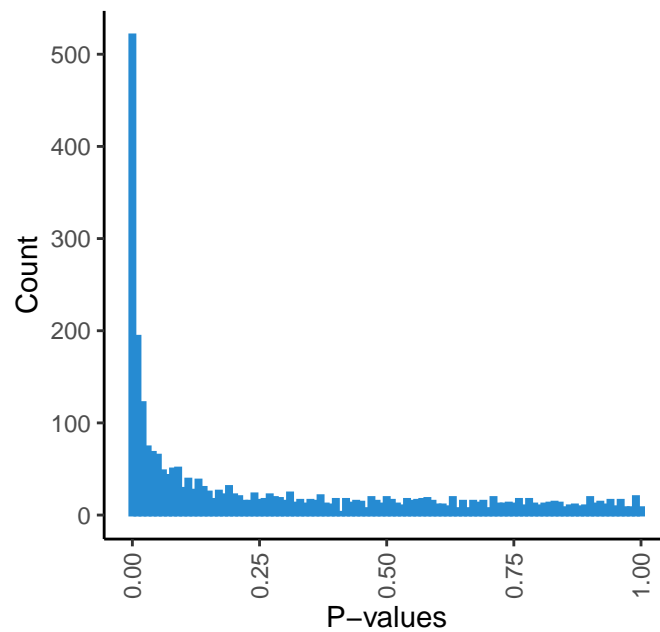
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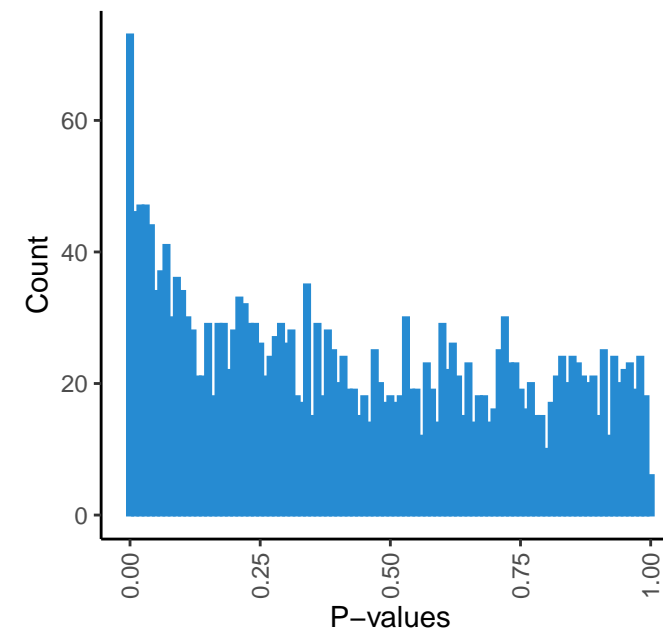
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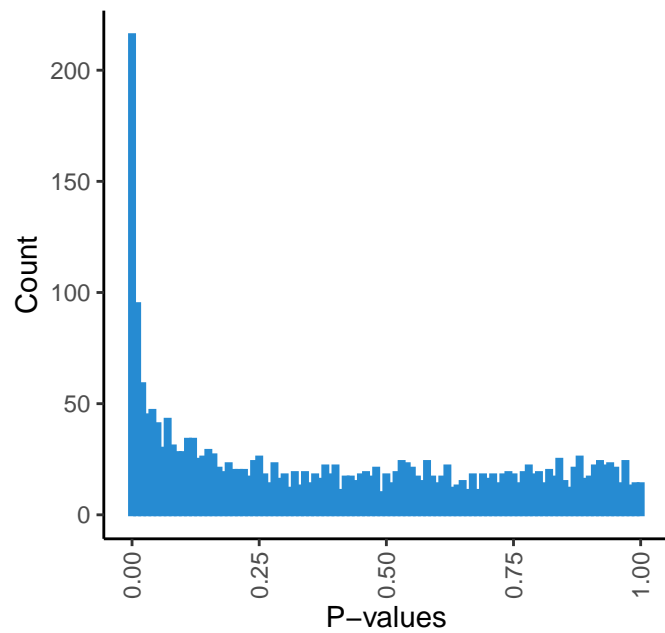
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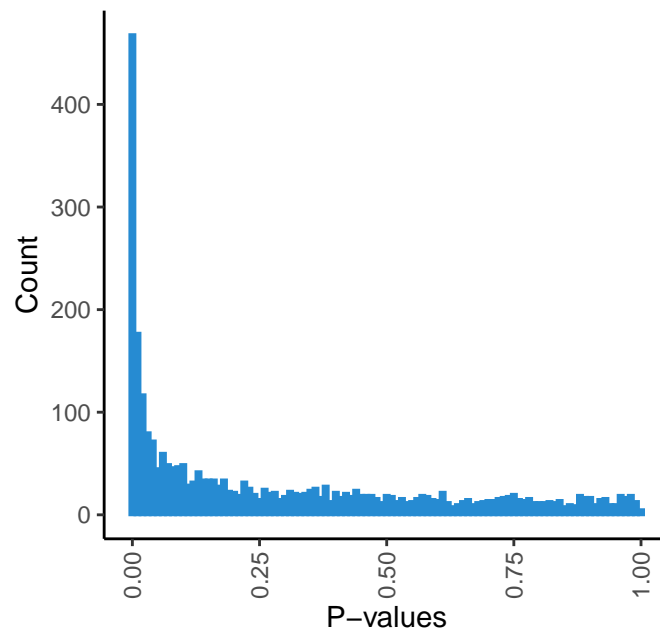
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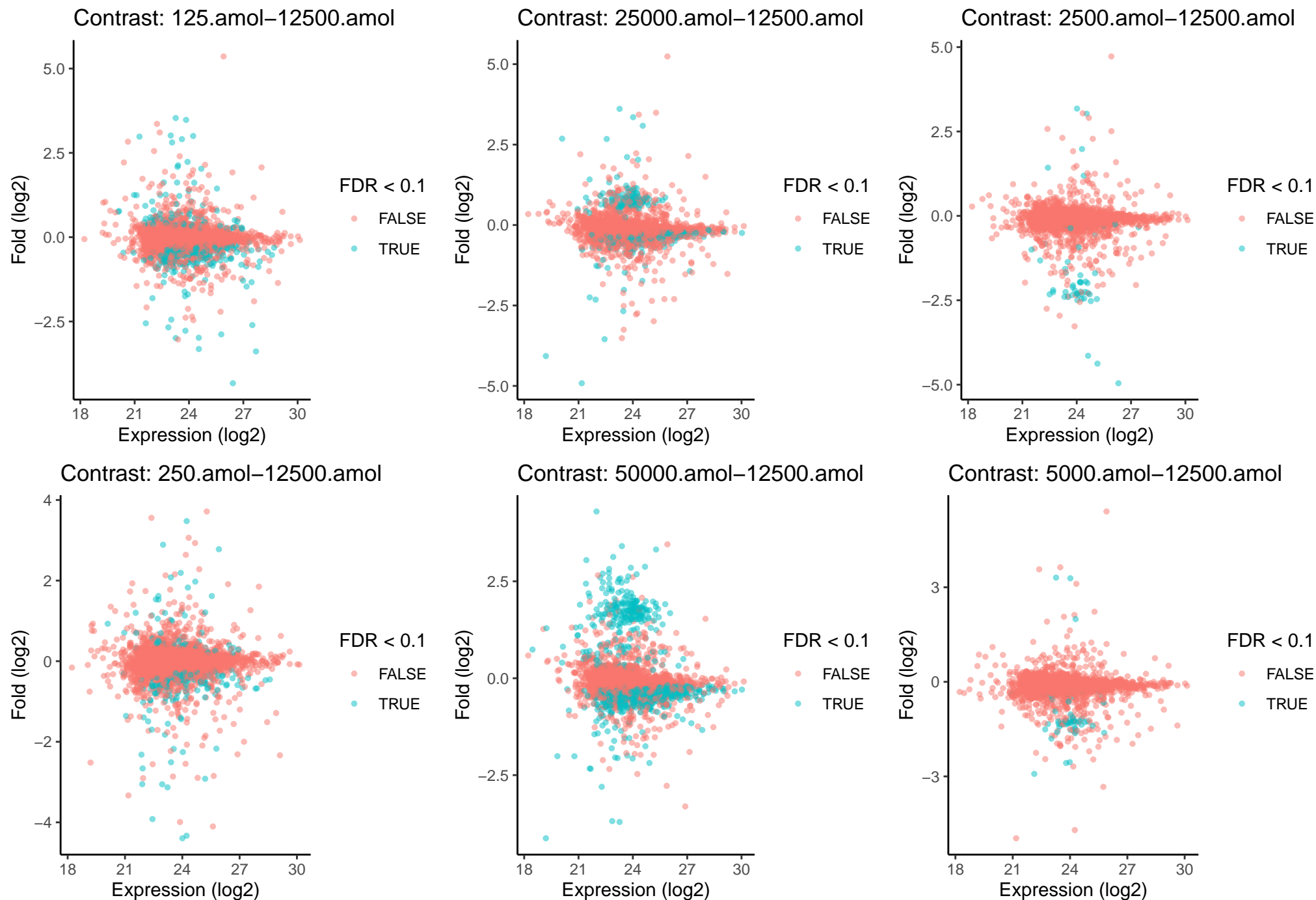


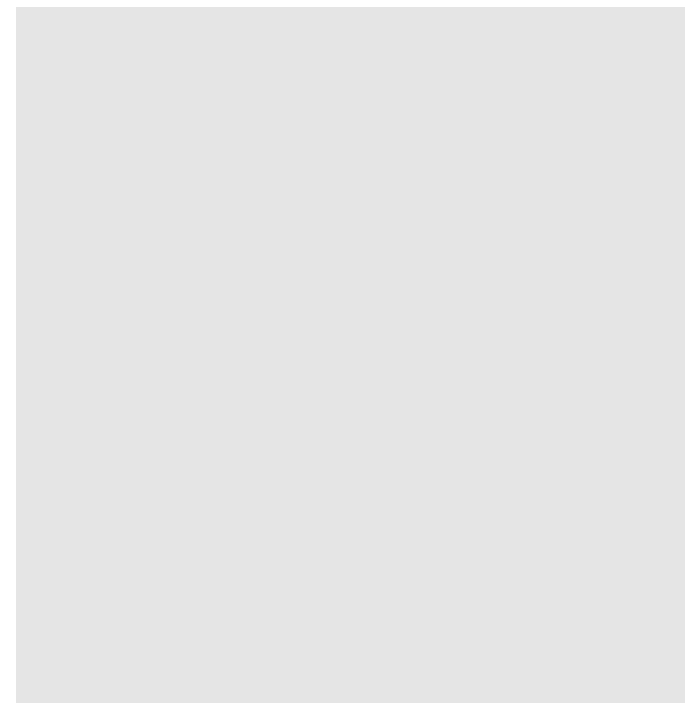
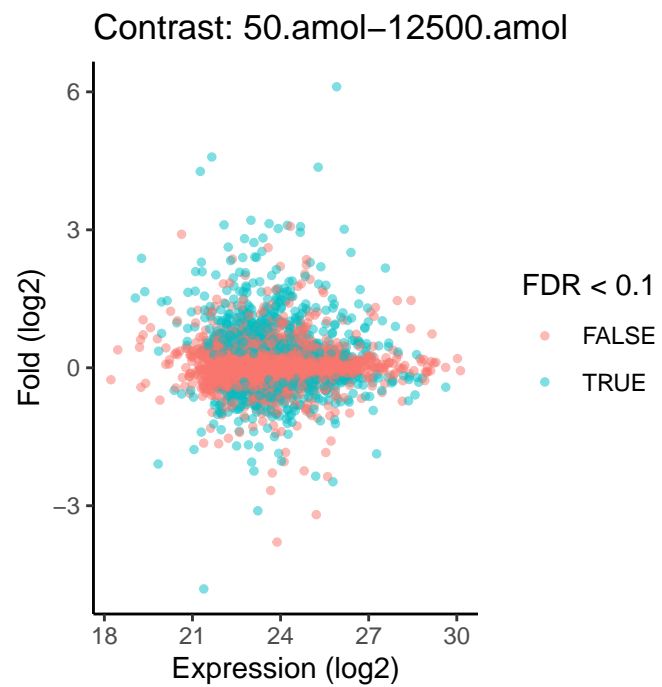
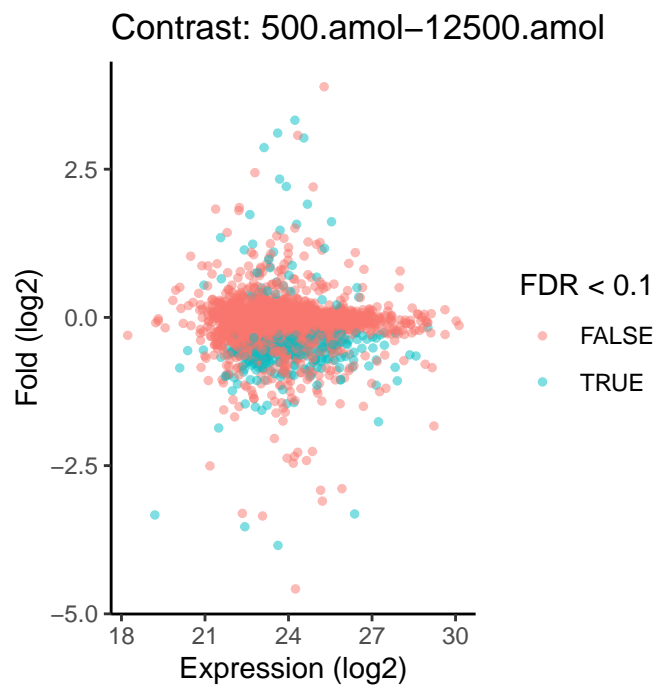
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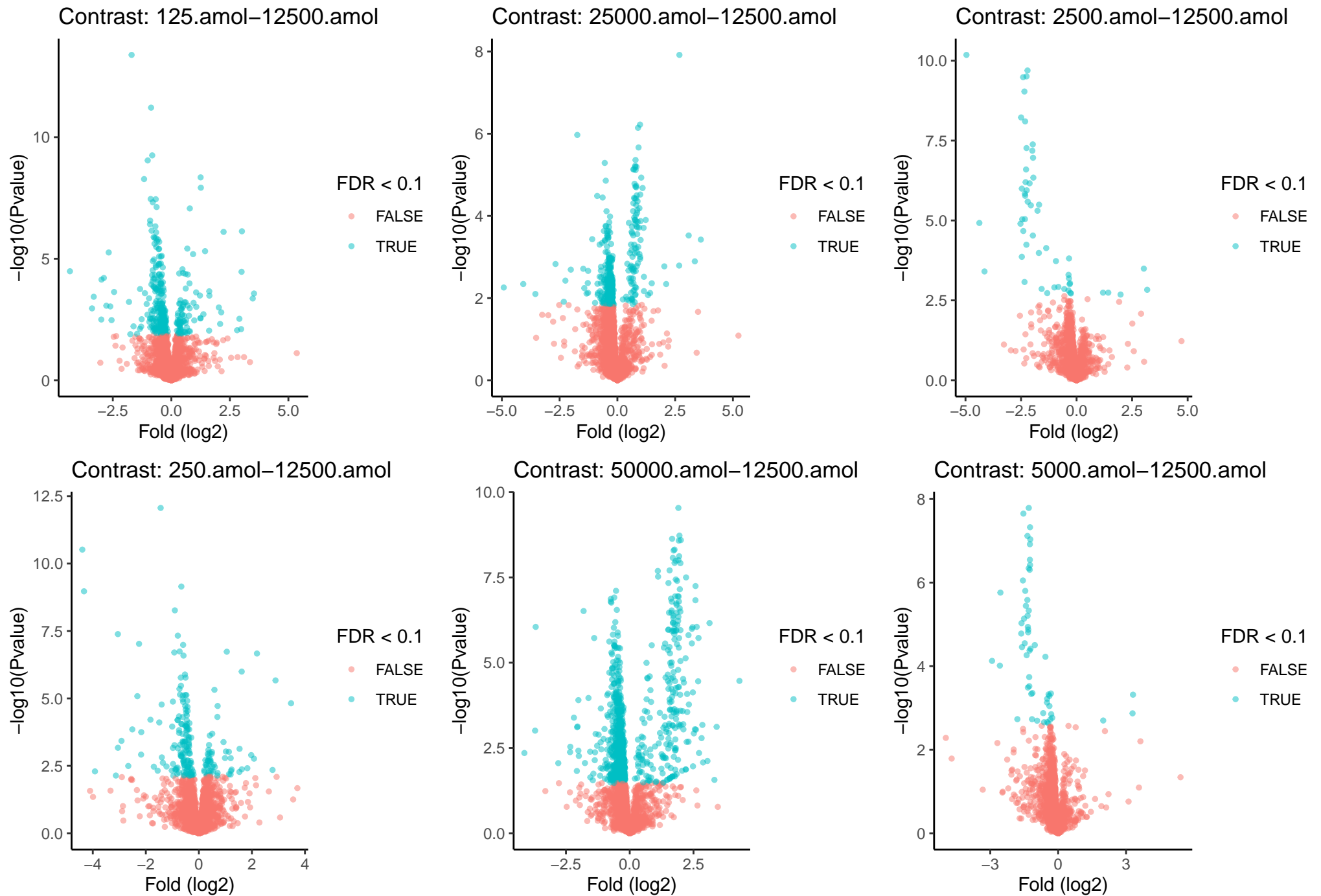


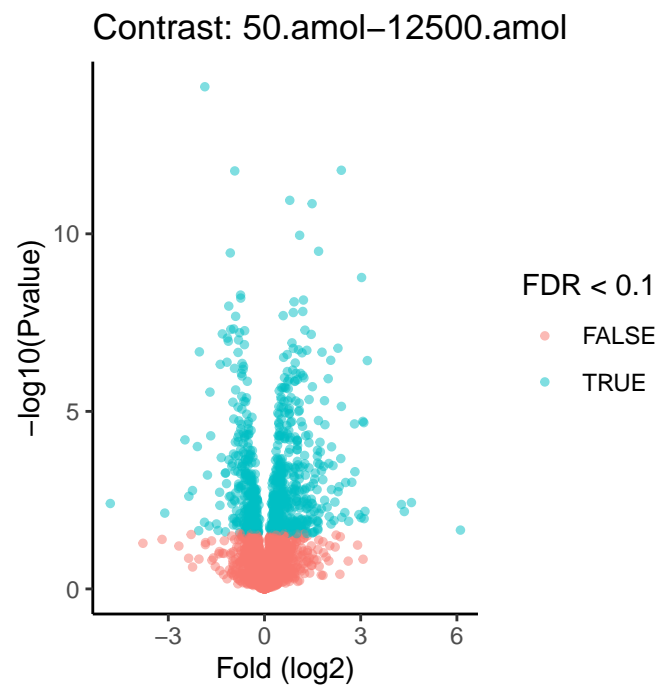
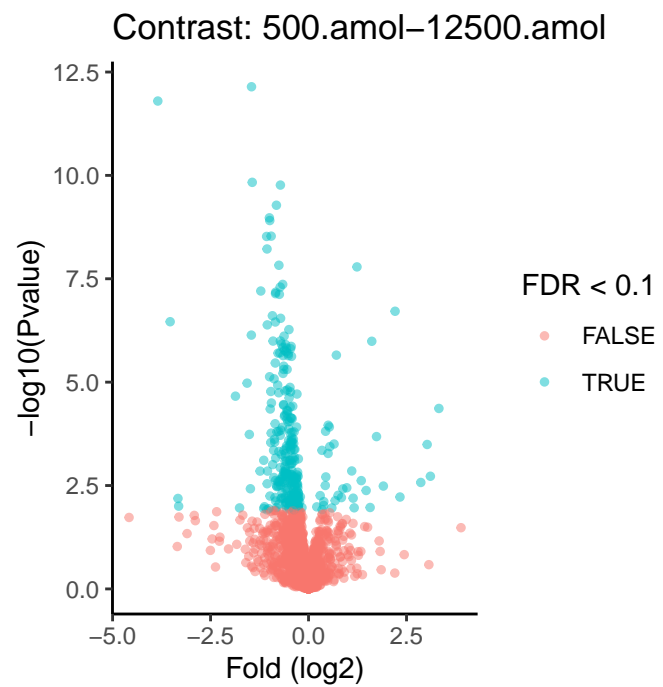
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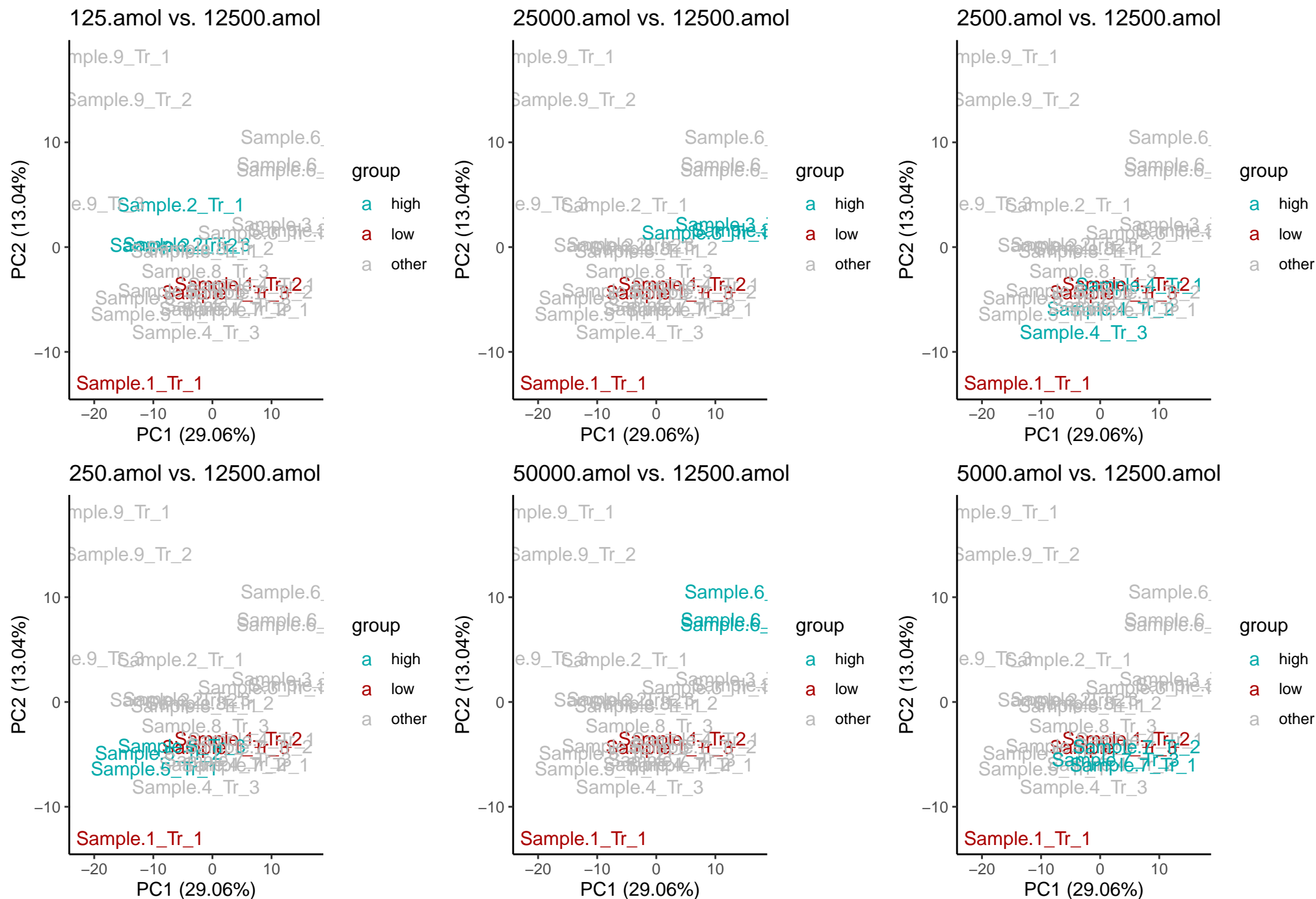


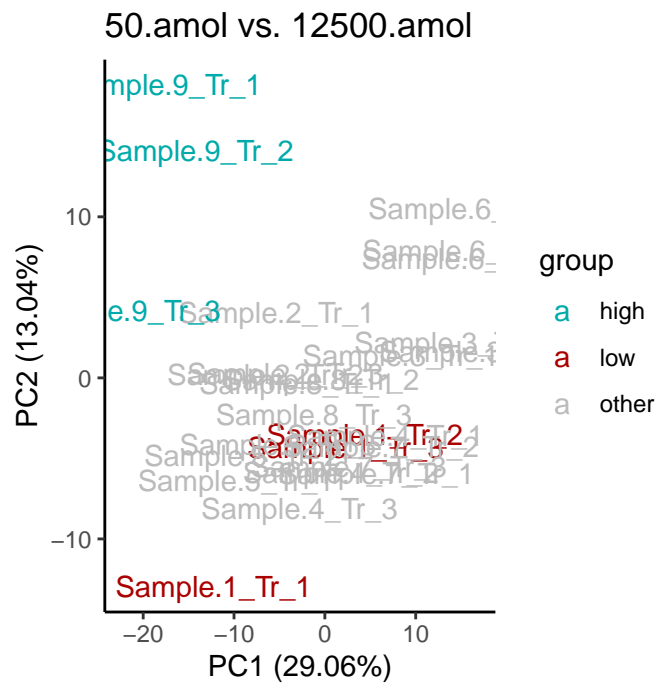
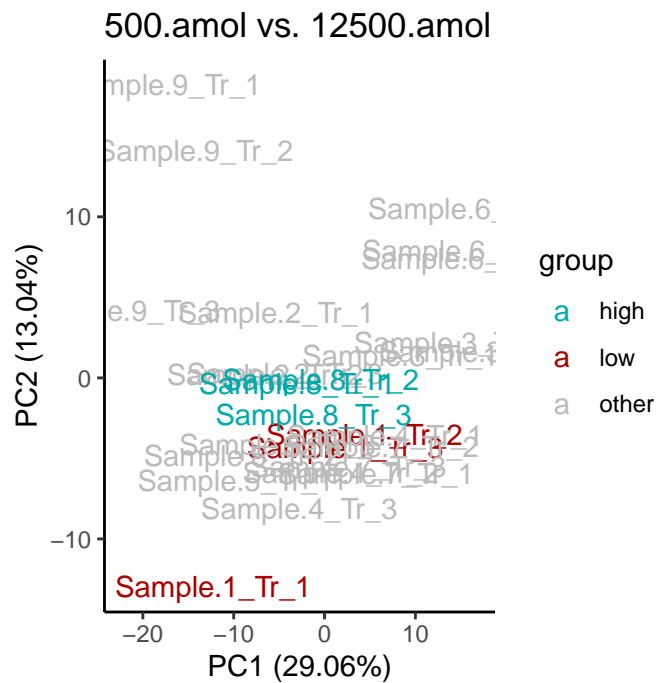


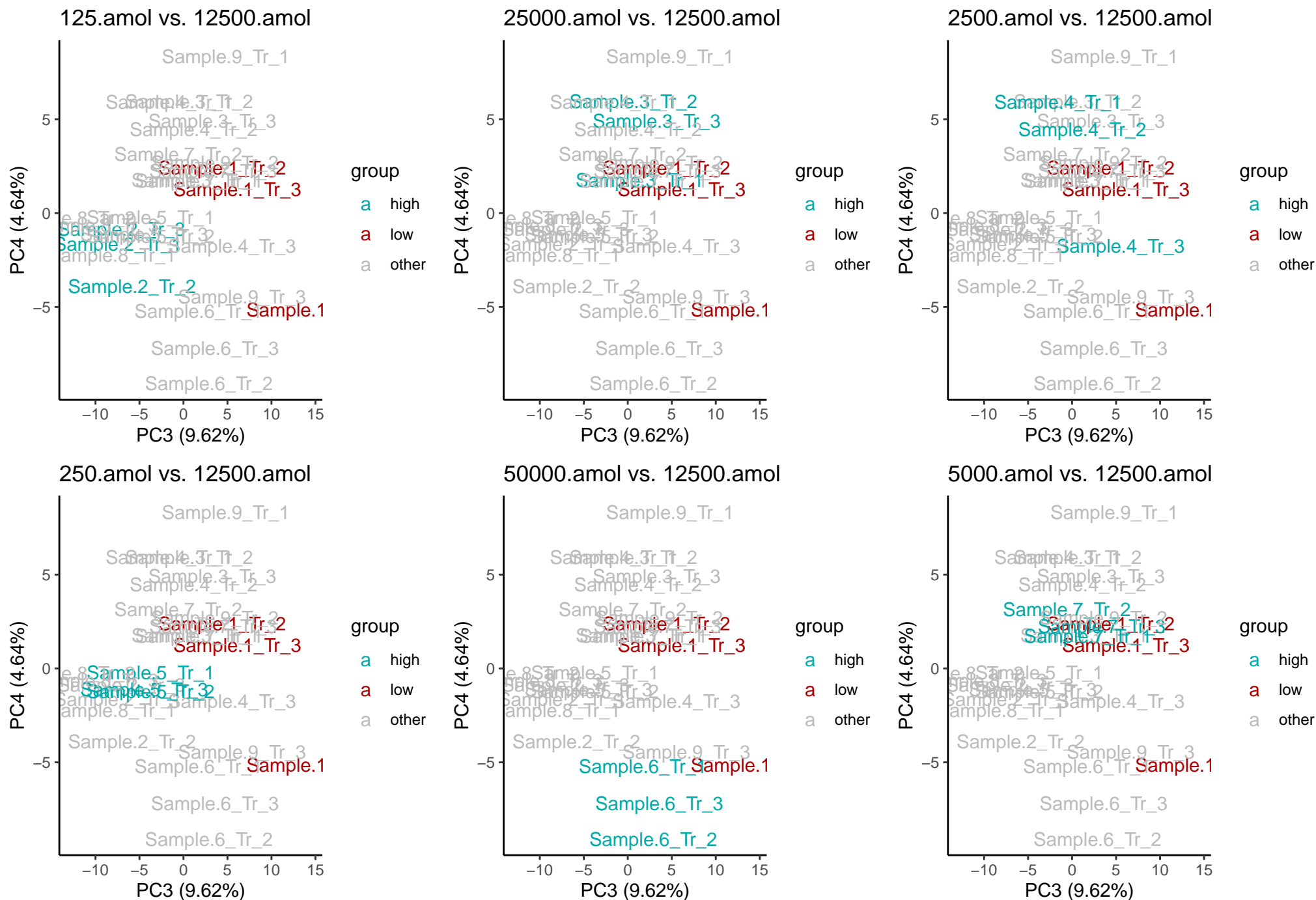


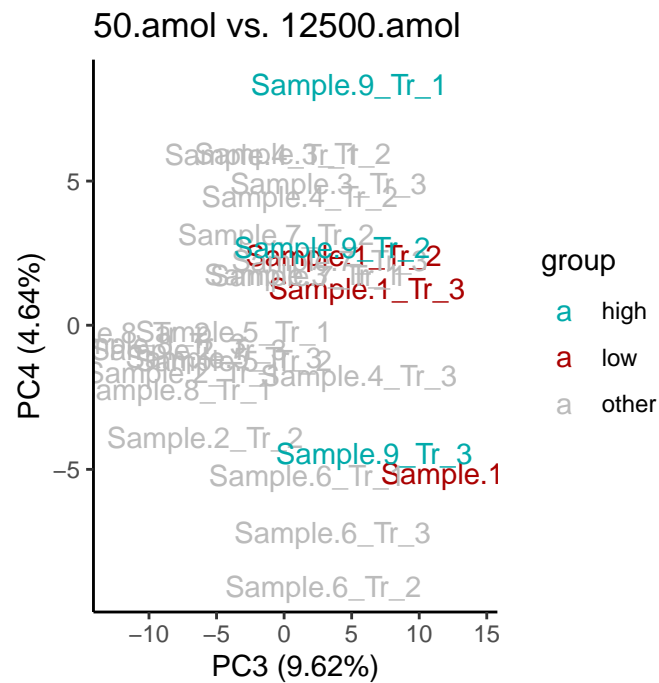
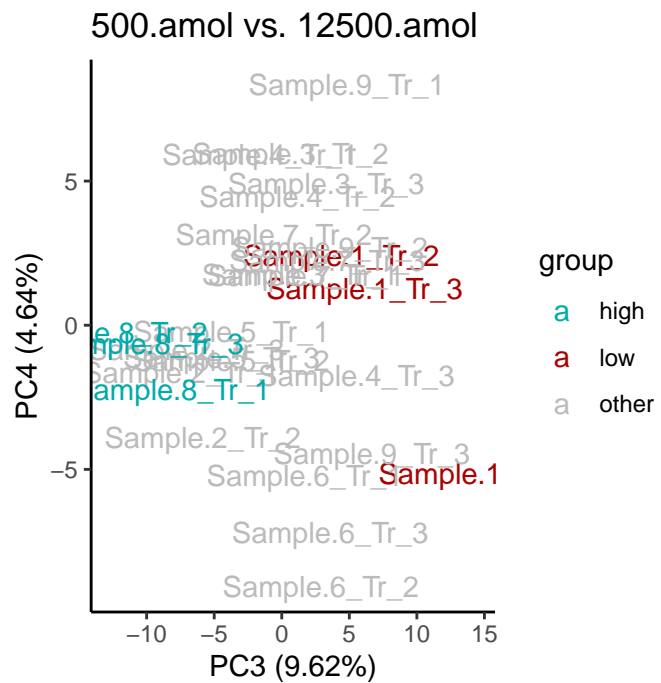




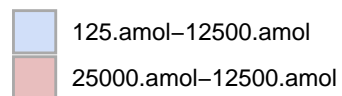
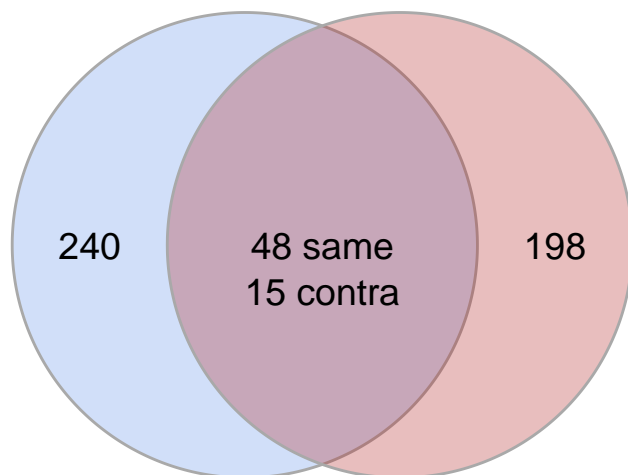




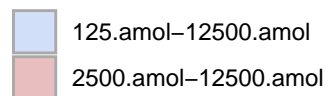
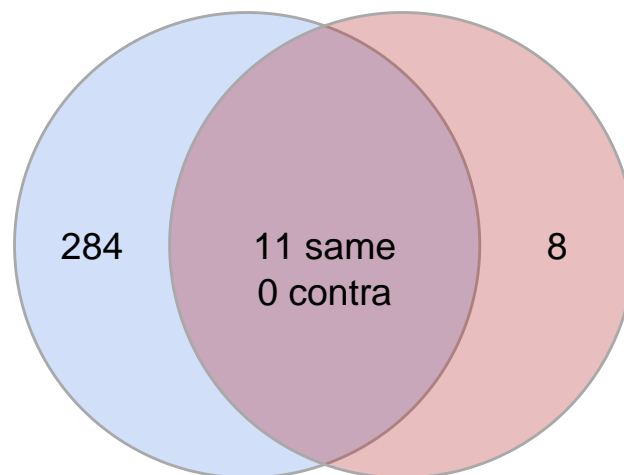




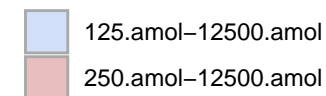
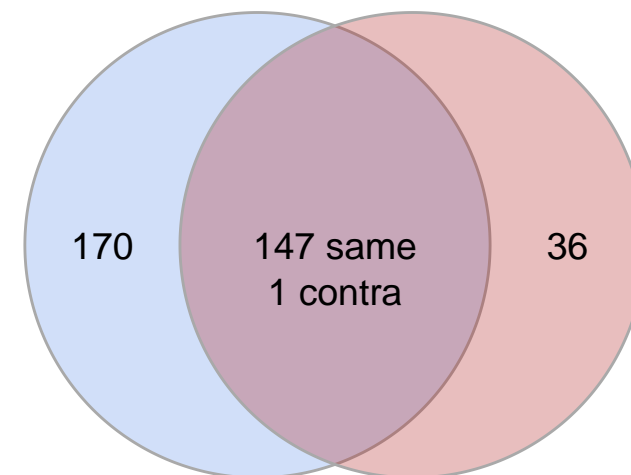
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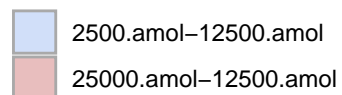
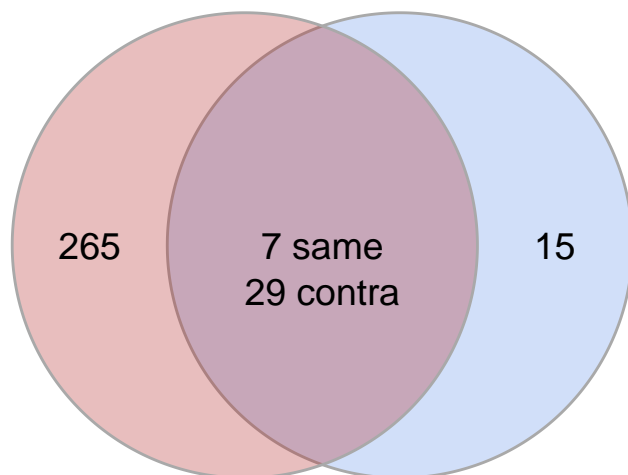
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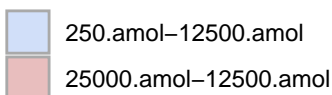
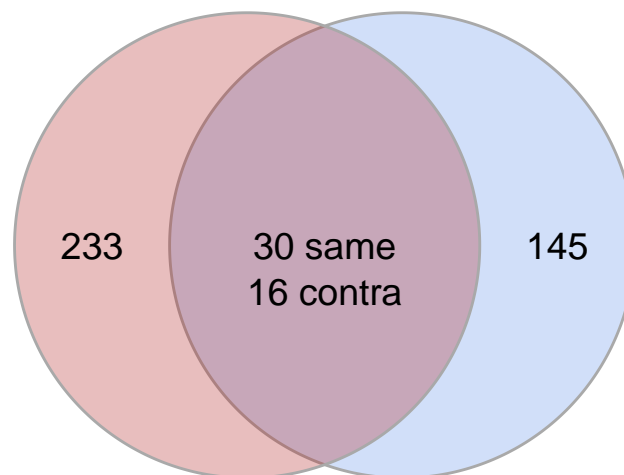
fdr < 0.1, |log2 fold| >= 0



fdr < 0.1, |log2 fold| >= 0



fdr < 0.1, |log2 fold| >= 0



fdr < 0.1, |log2 fold| >= 0

