

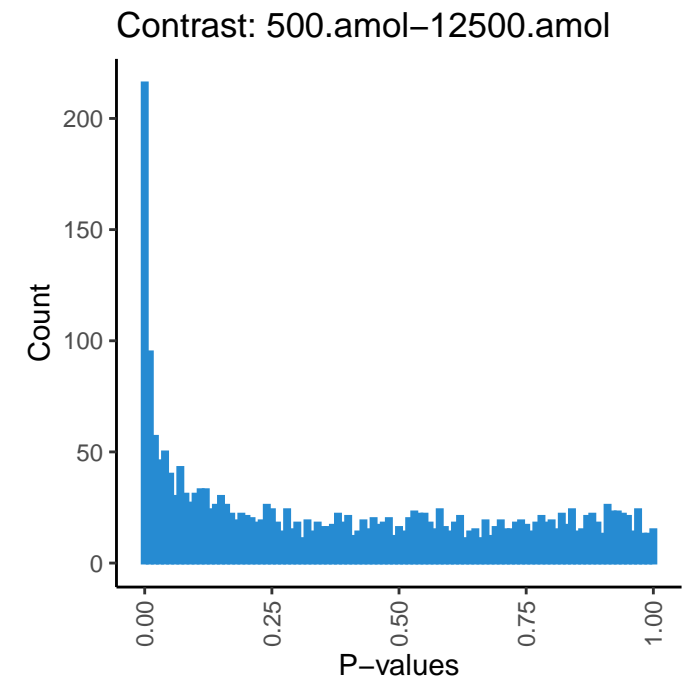
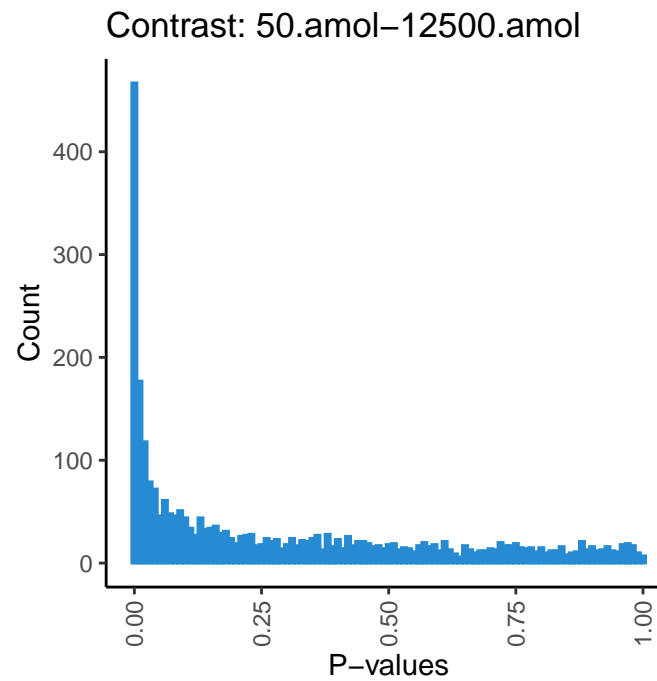
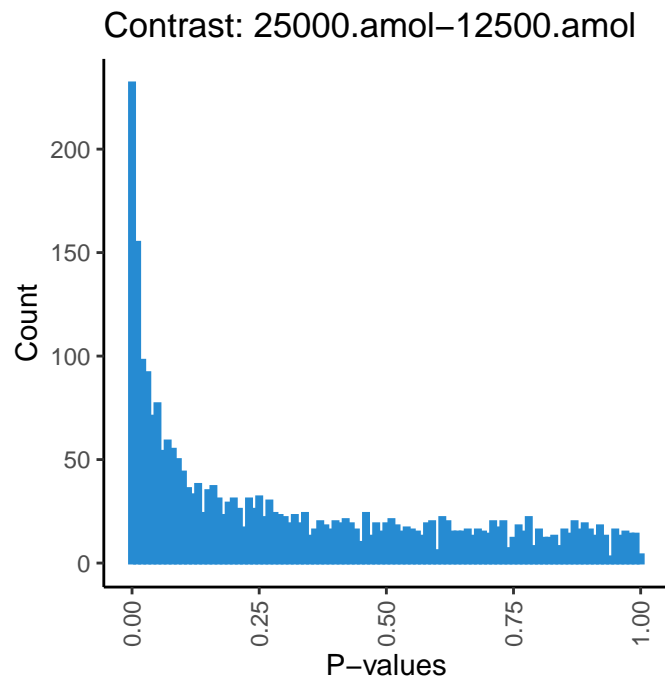
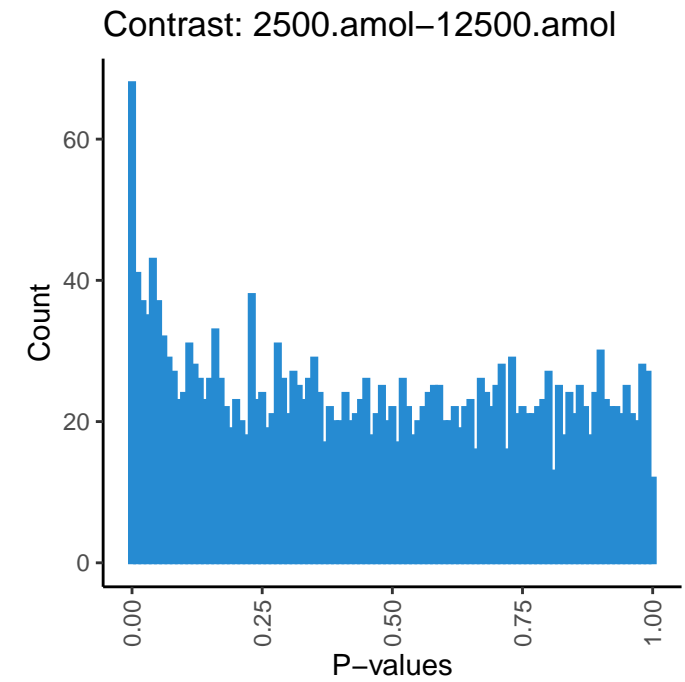
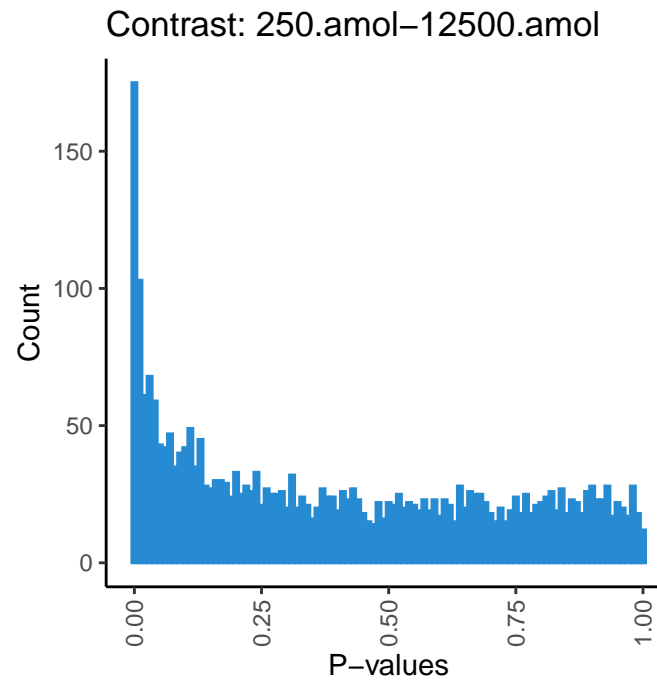
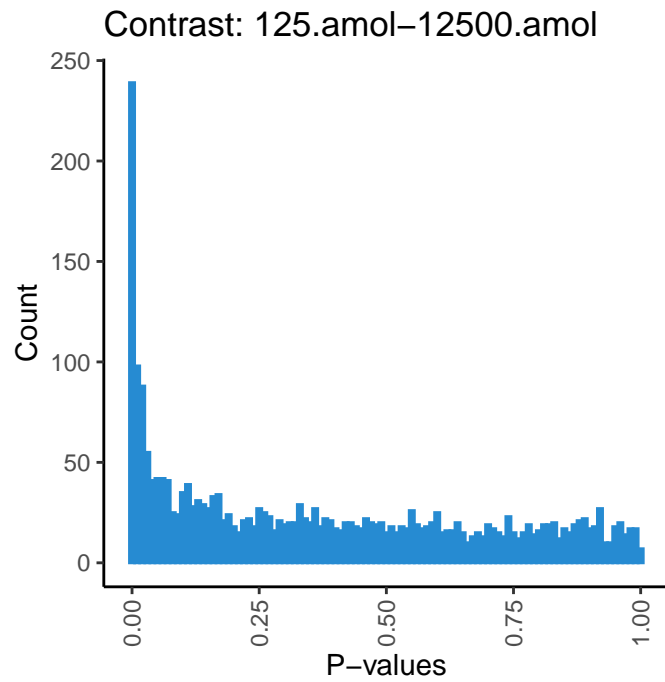
Project Name: NormalyzerPeptides

NormalyzerDE (ver 1.14.0)

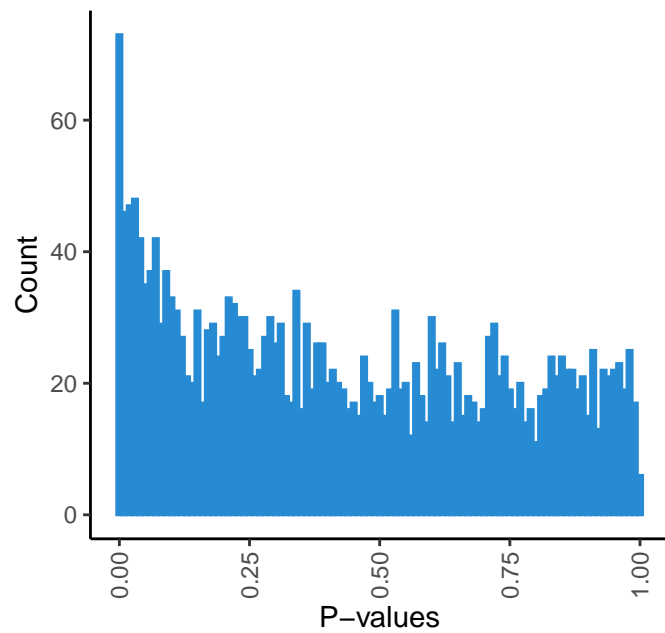
Report created on: 2022-07-12

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

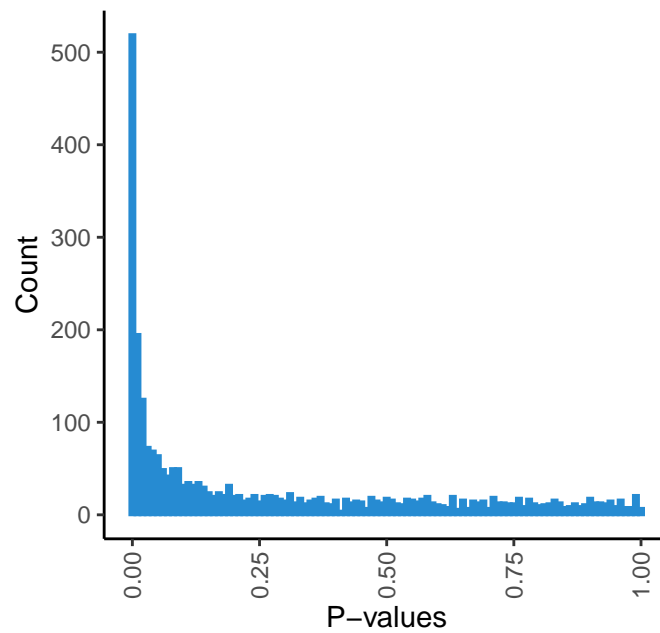
Documentation for analyzing this report can be found at <http://quantitativeproteomics.org/normalyzer/help.php>

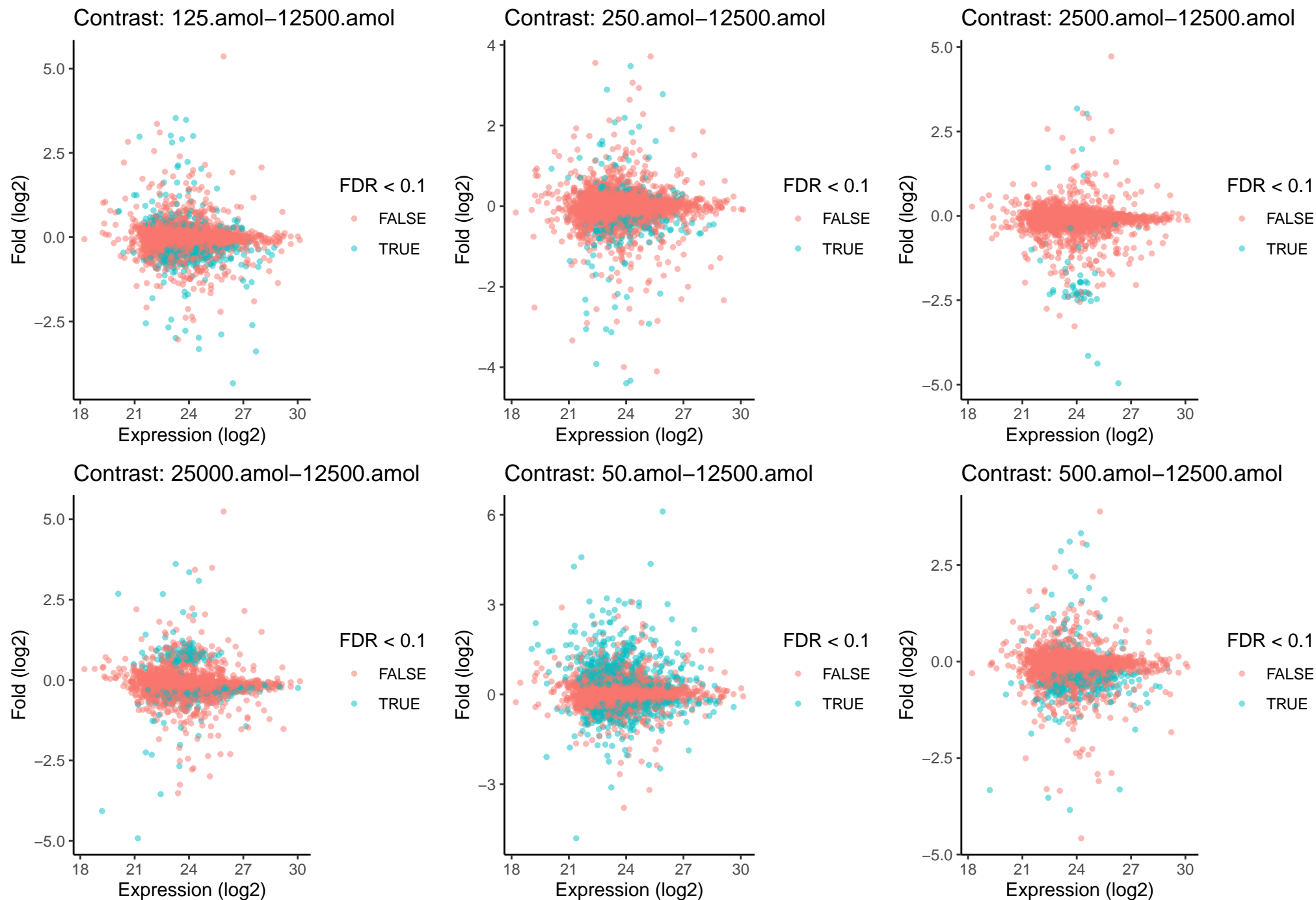


Contrast: 5000.amol–12500.amol

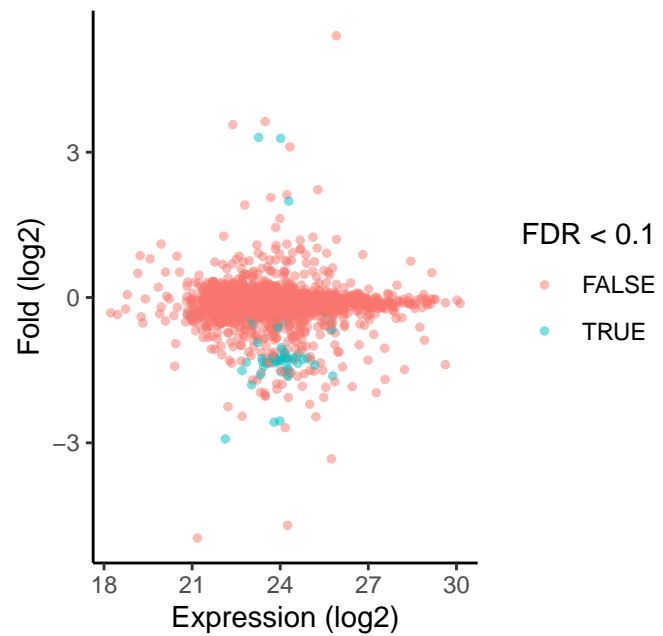


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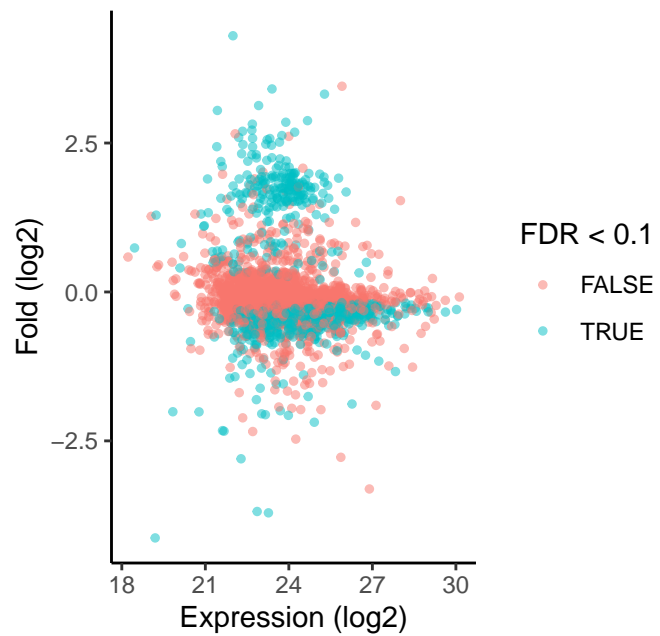


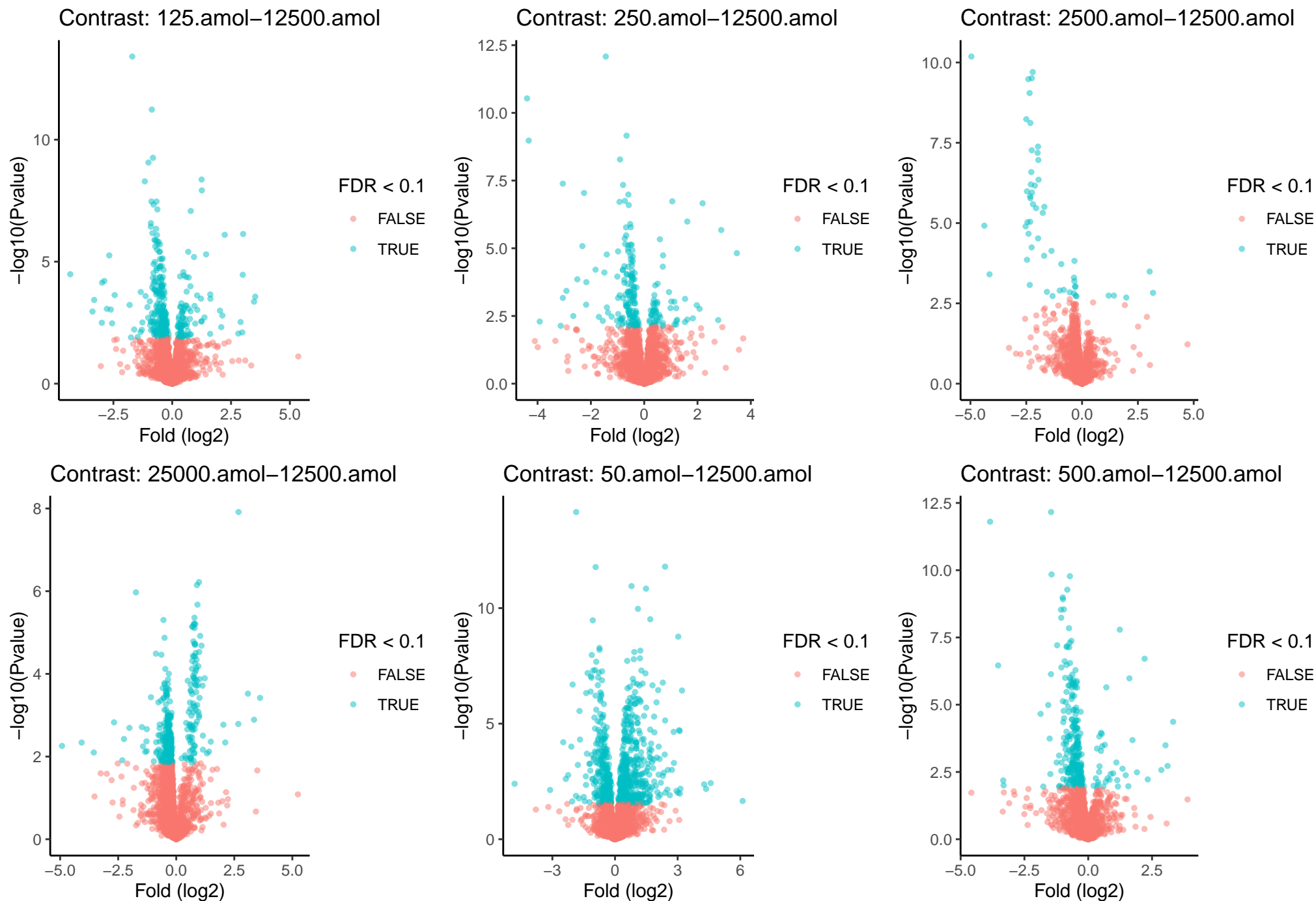


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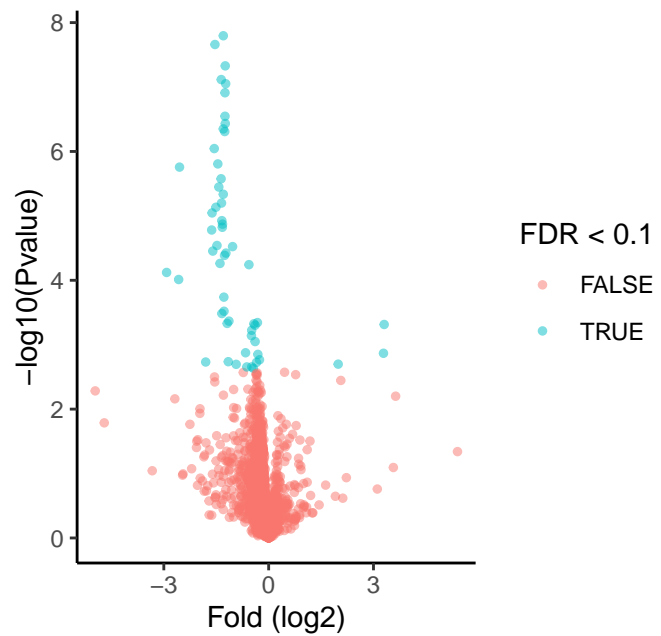


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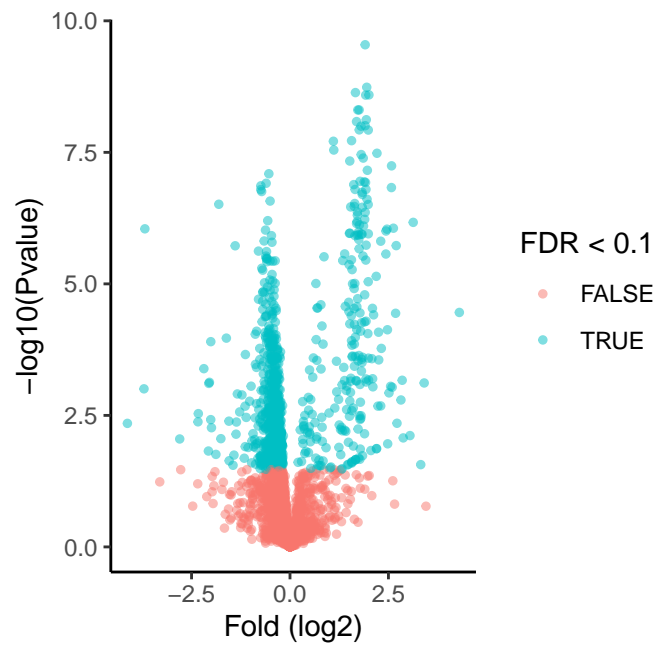


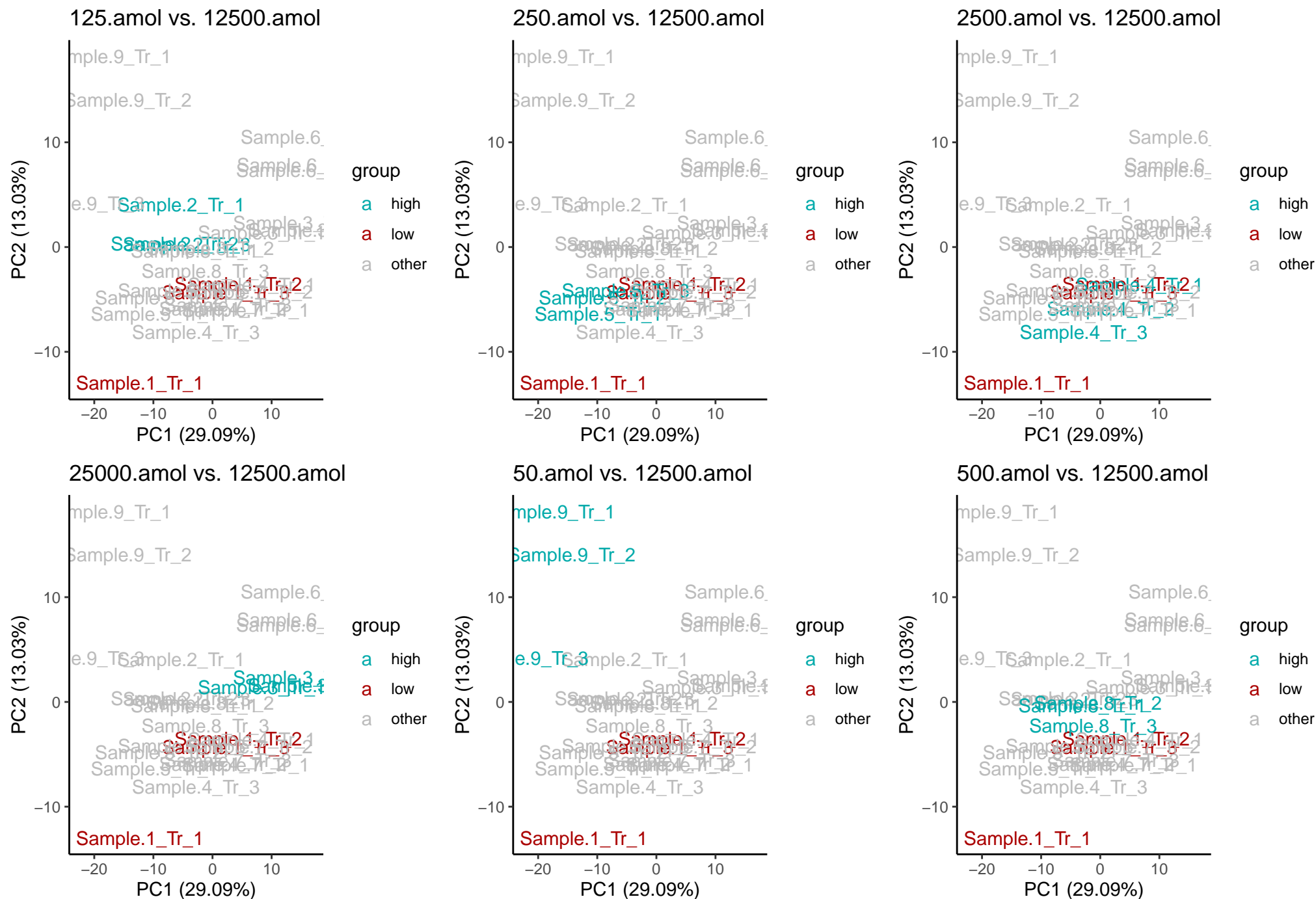


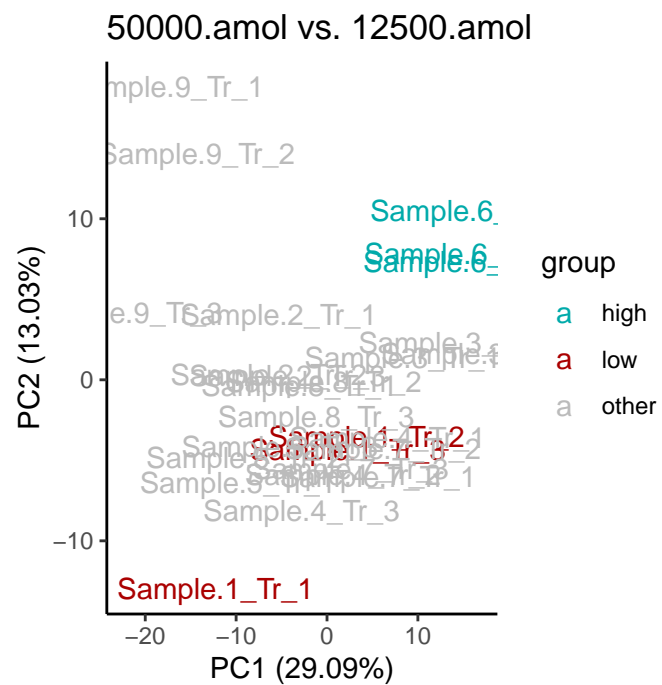
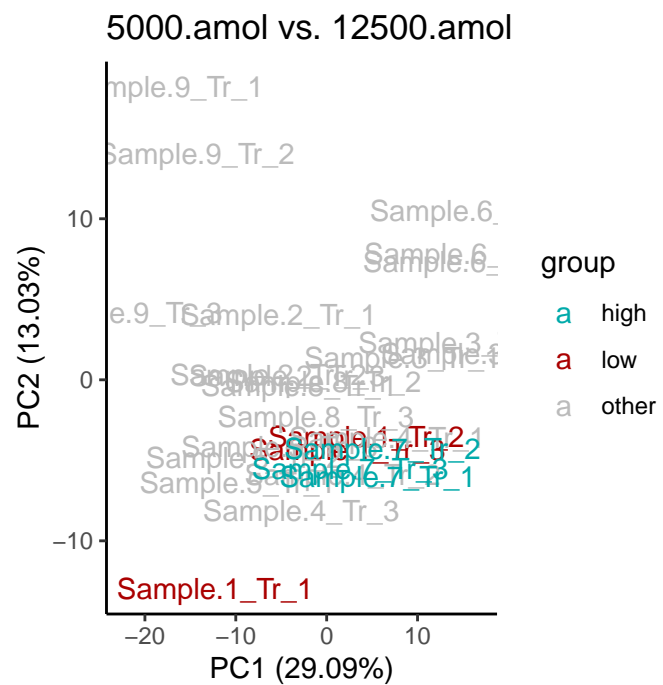
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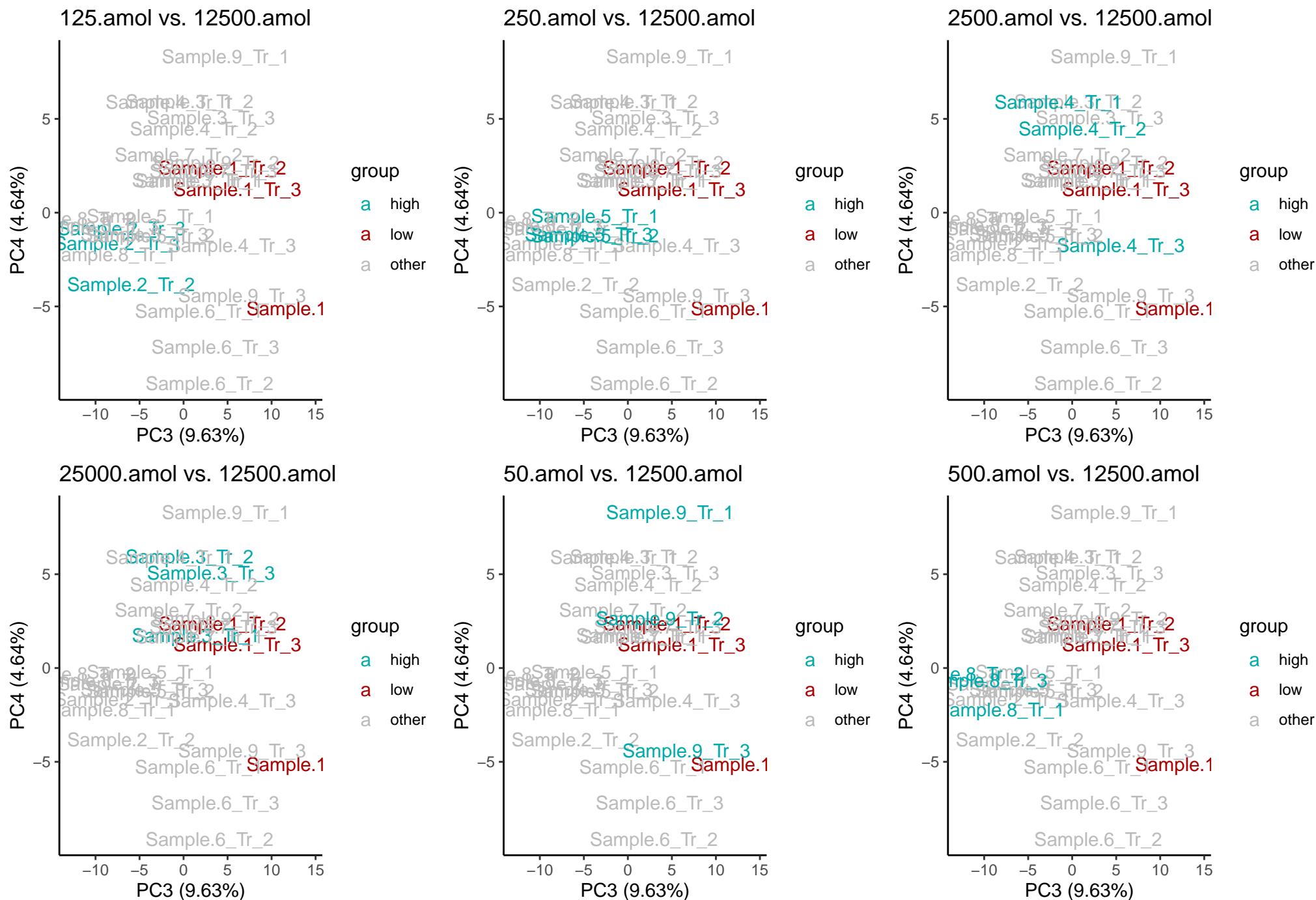


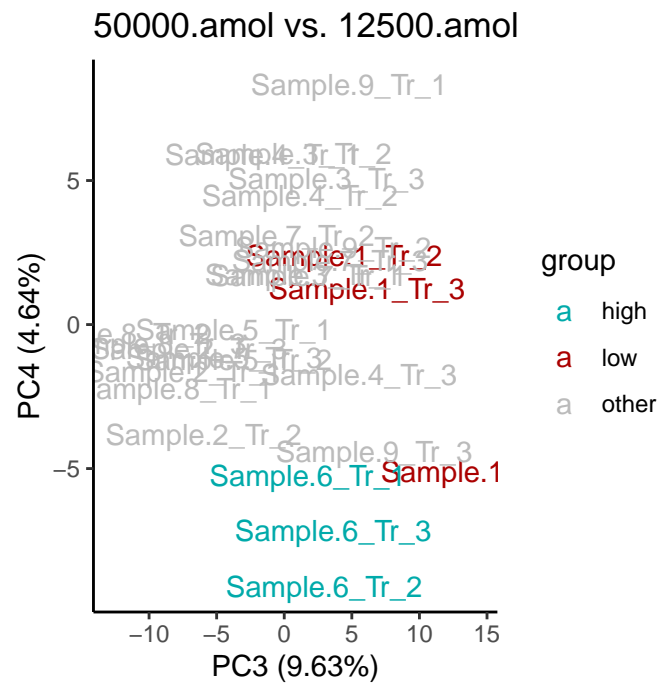
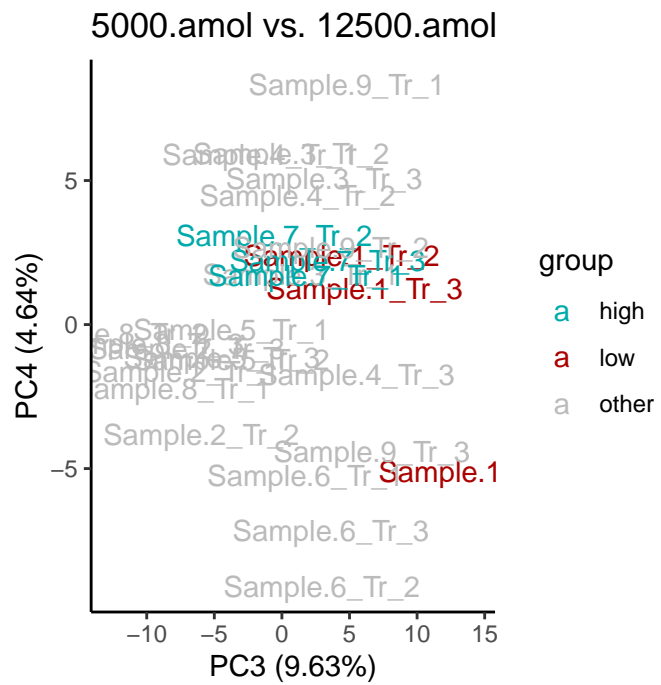
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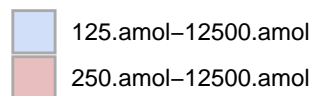
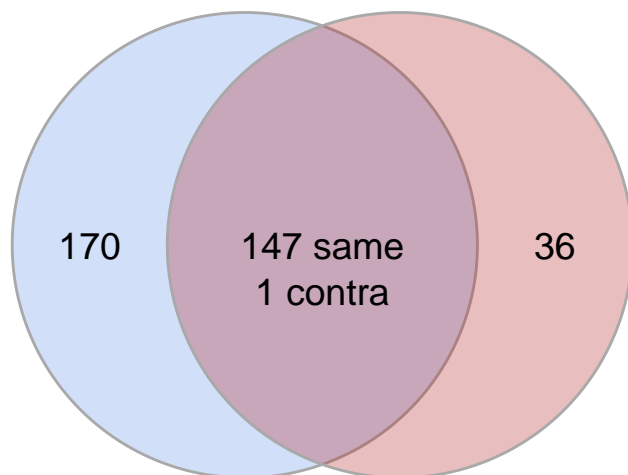




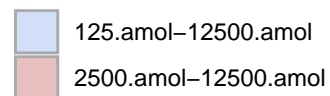
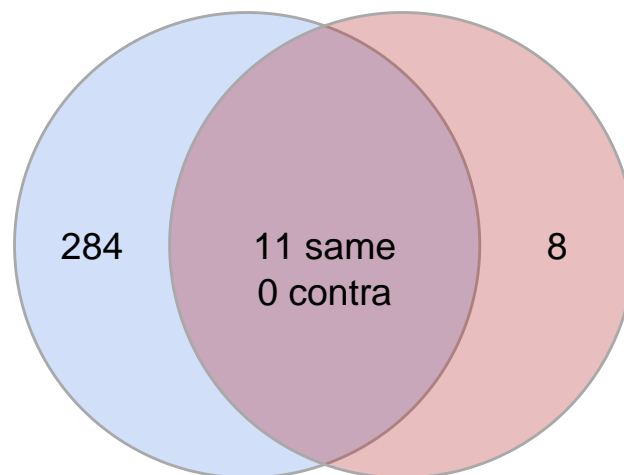




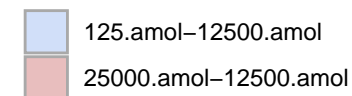
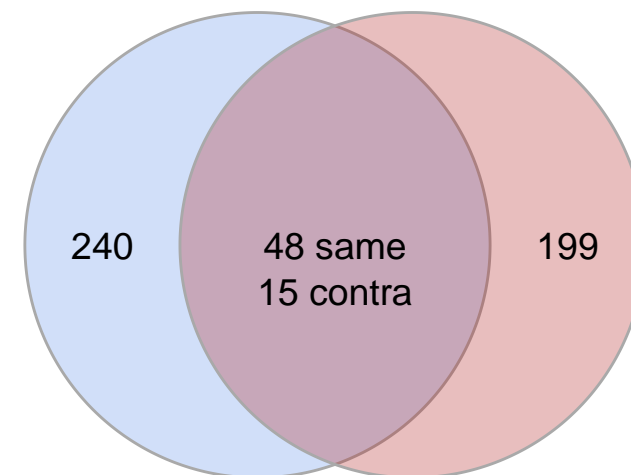
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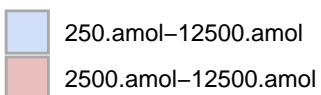
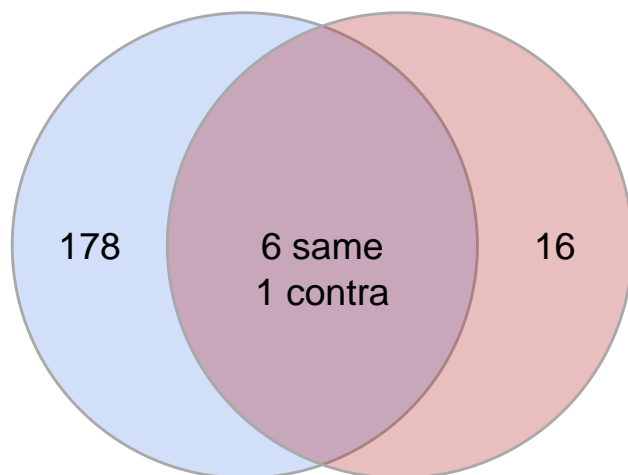
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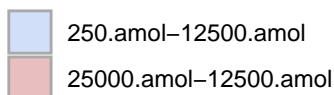
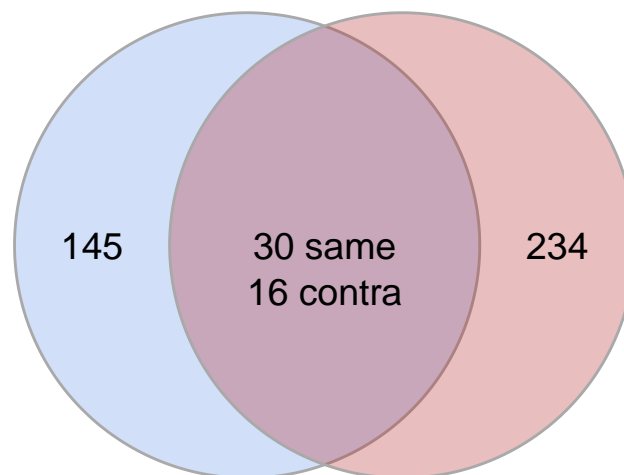
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fdr < 0.1, |log2 fold| >= 0



fdr < 0.1, |log2 fold| >= 0



fdr < 0.1, |log2 fold| >= 0

