

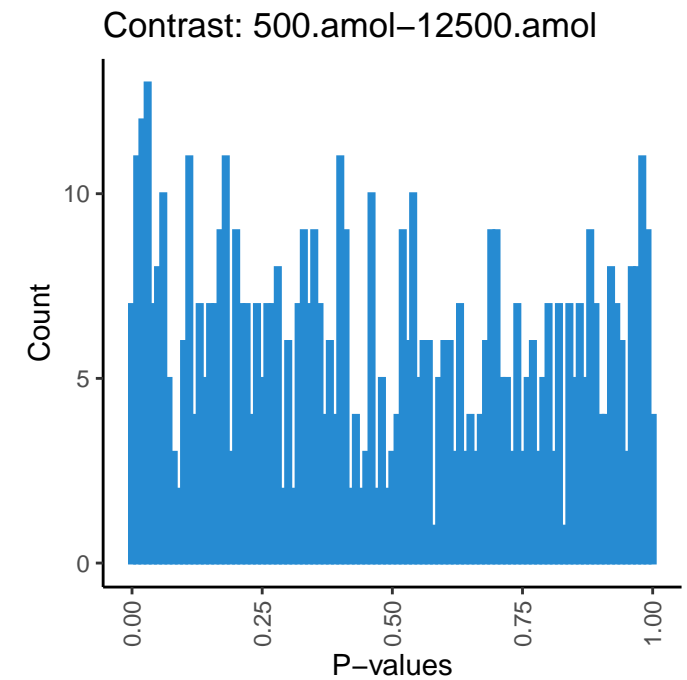
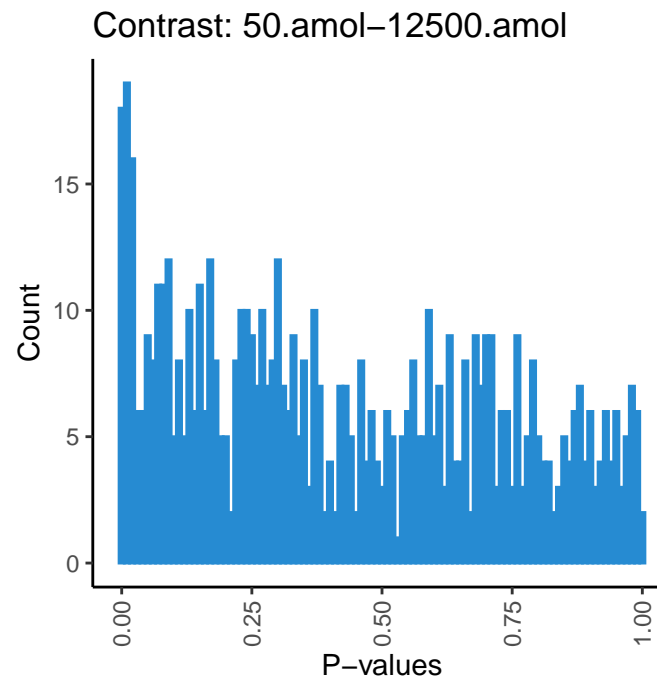
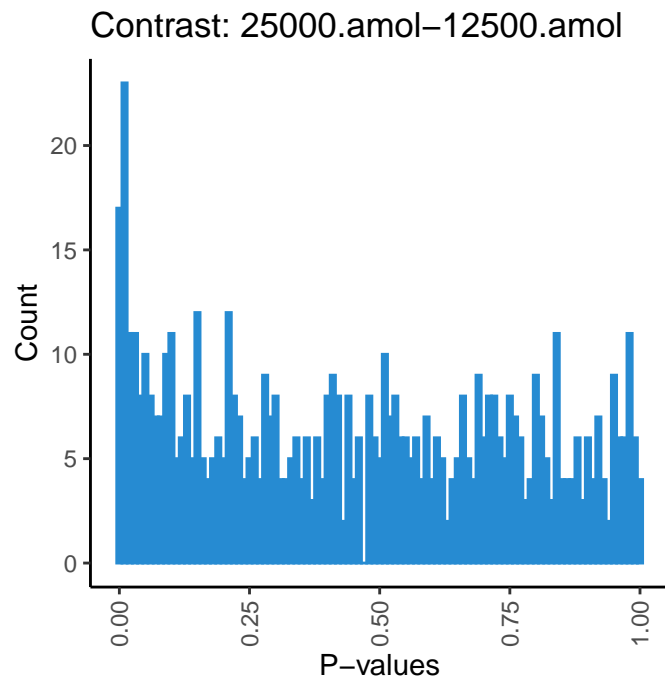
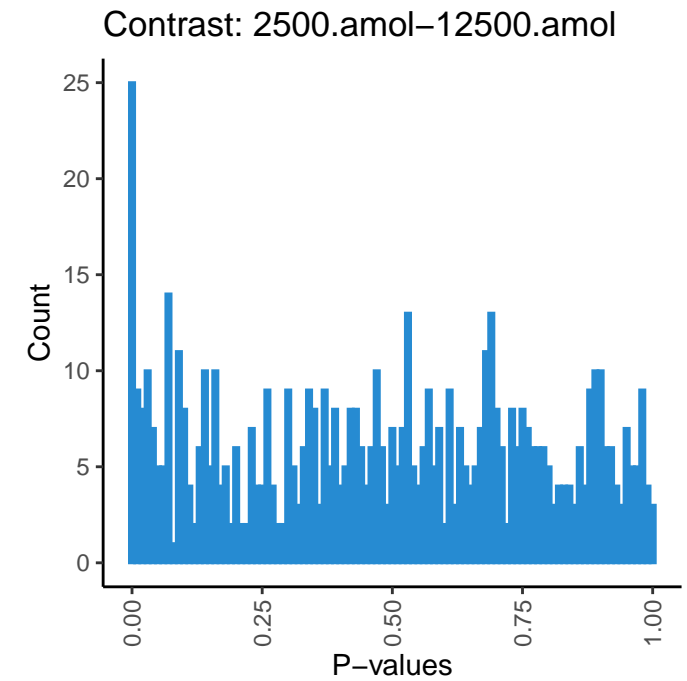
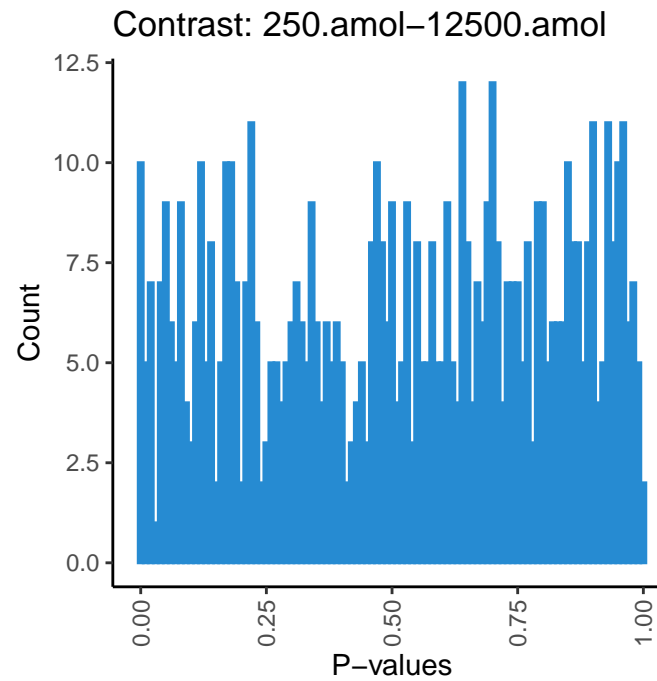
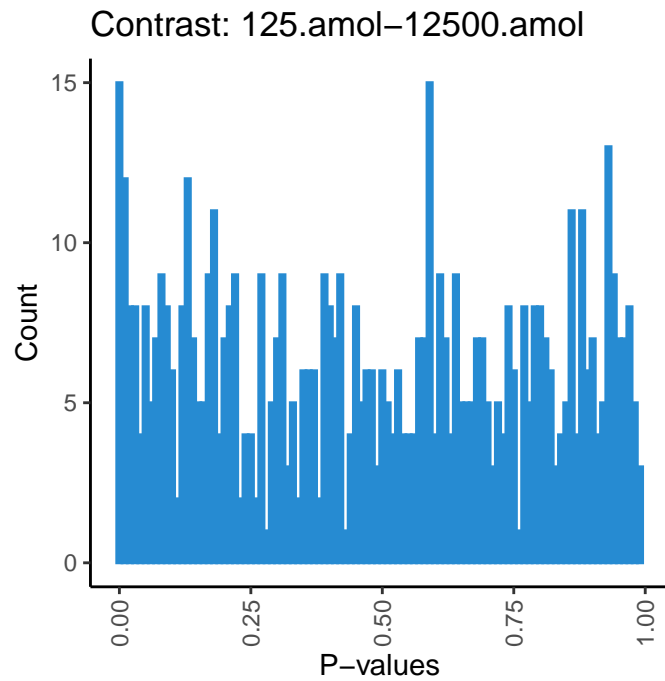
Project Name: NormalyzerProteins

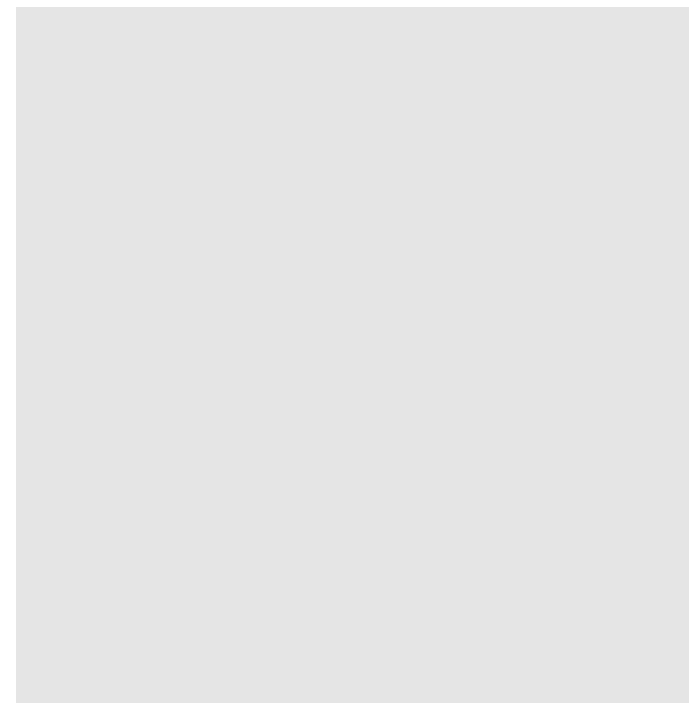
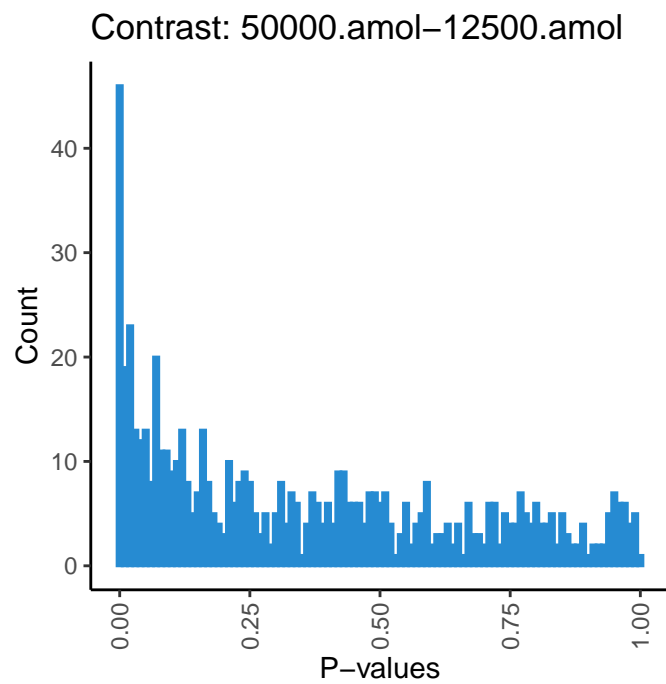
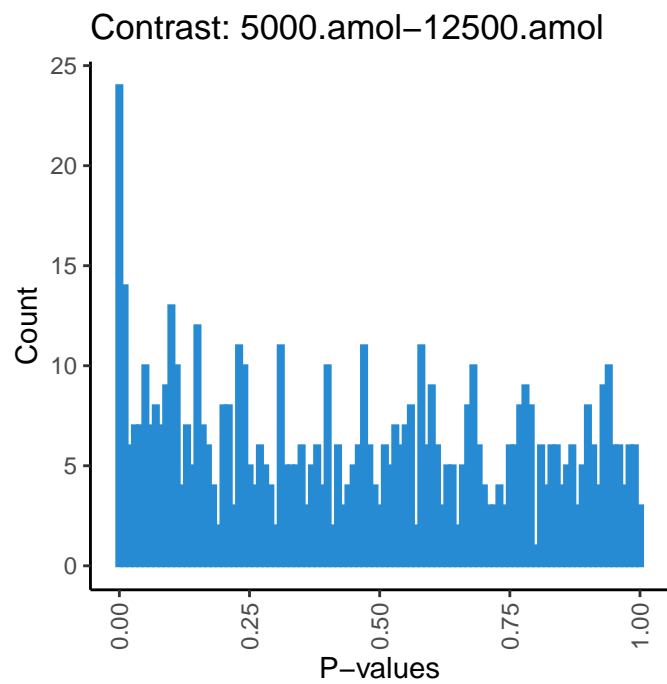
# NormalyzerDE (ver 1.14.0 )

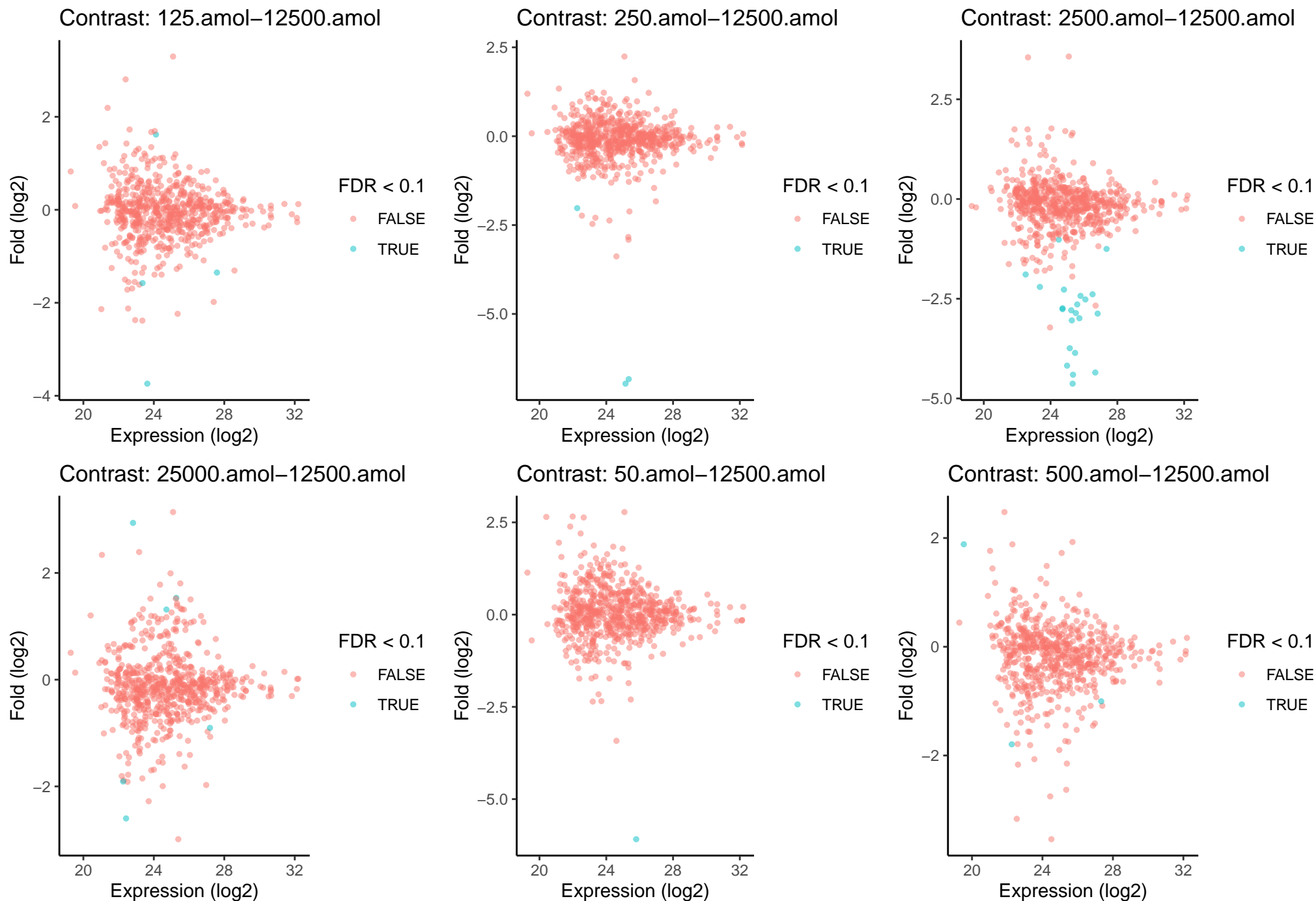
Report created on: 2022-07-12

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

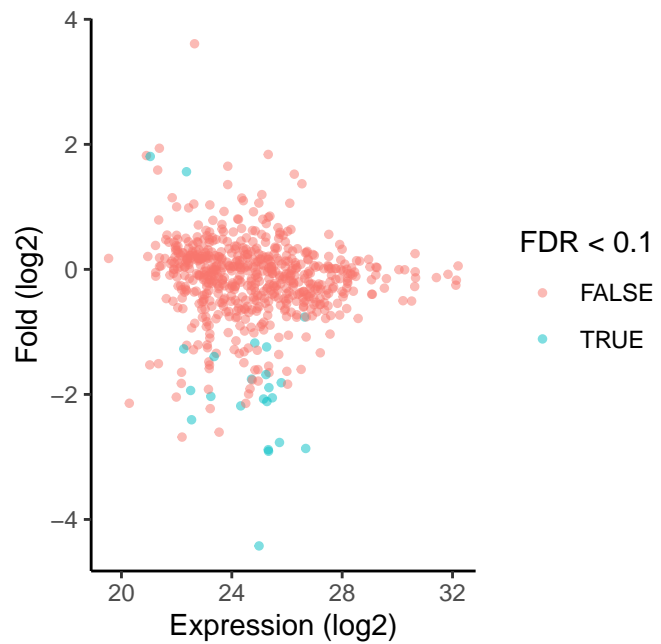
Documentation for analyzing this report can be found at <http://quantitativeproteomics.org/normalyzer/help.php>



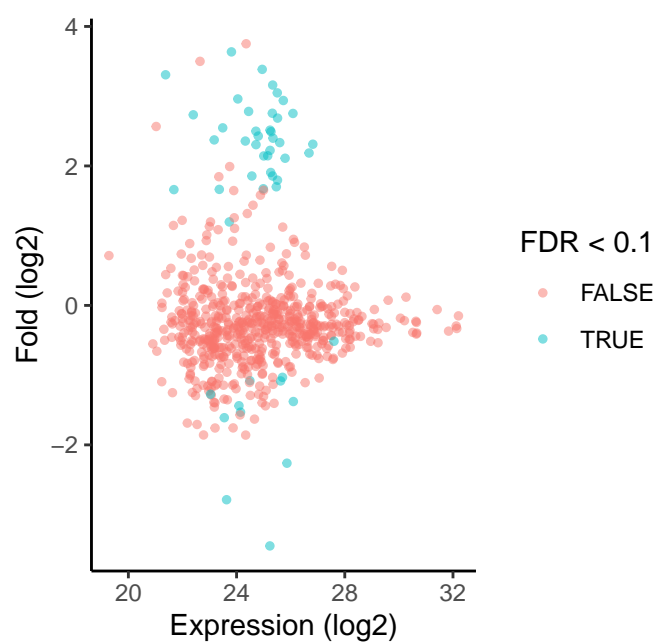


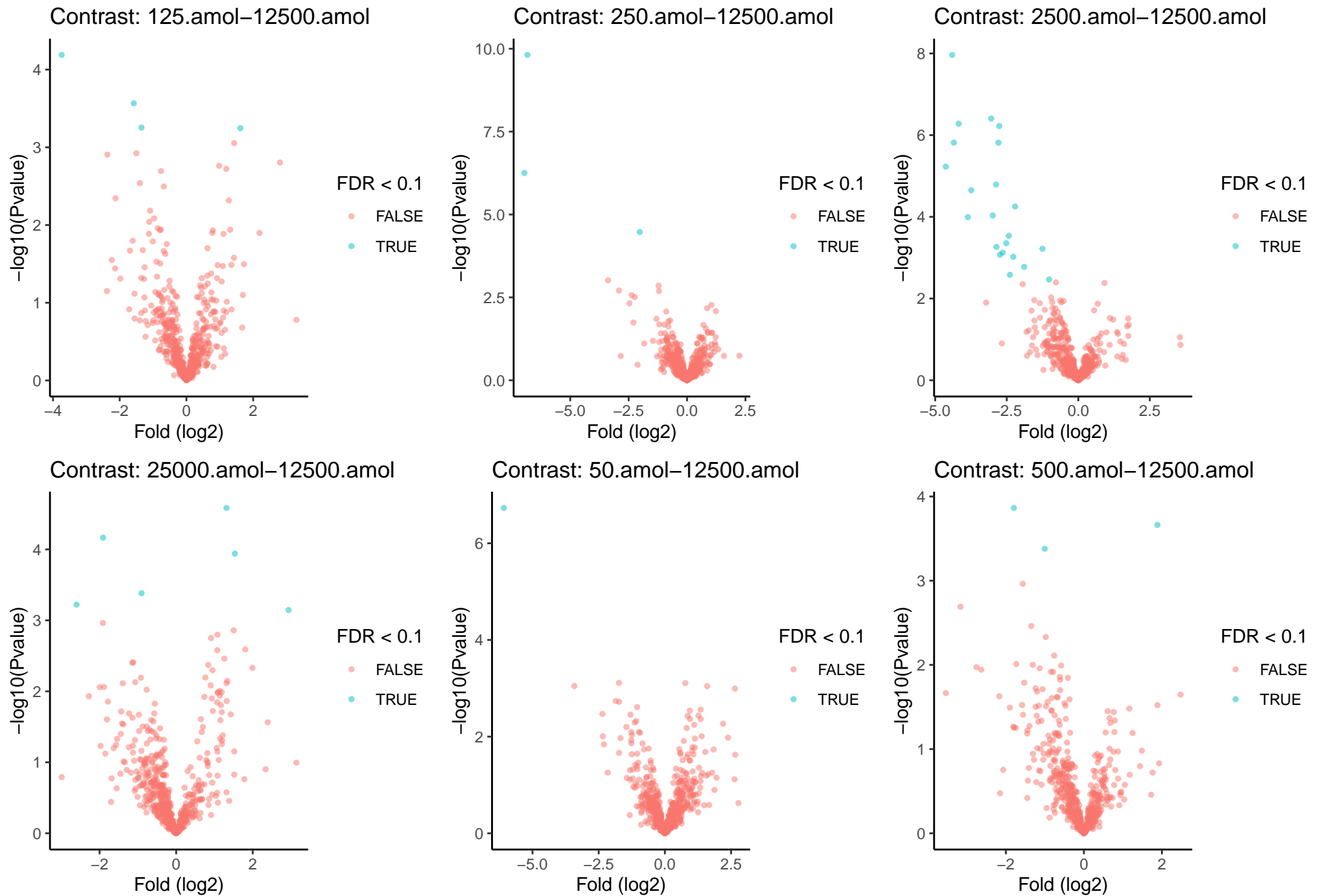


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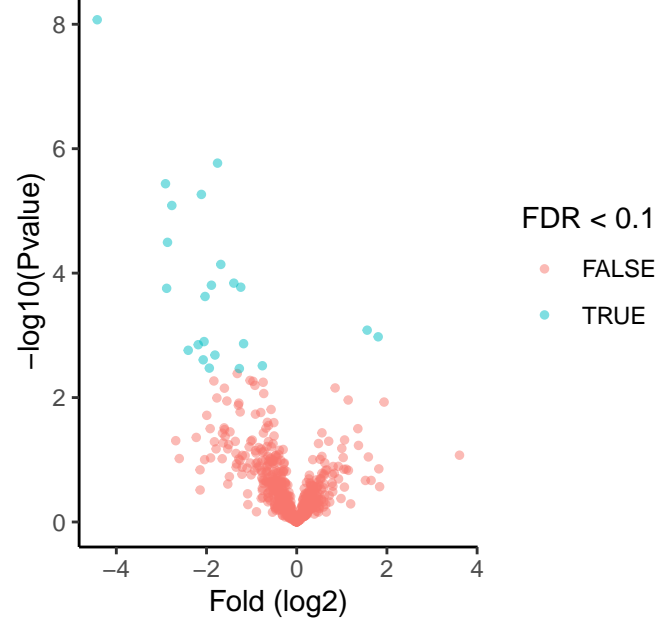


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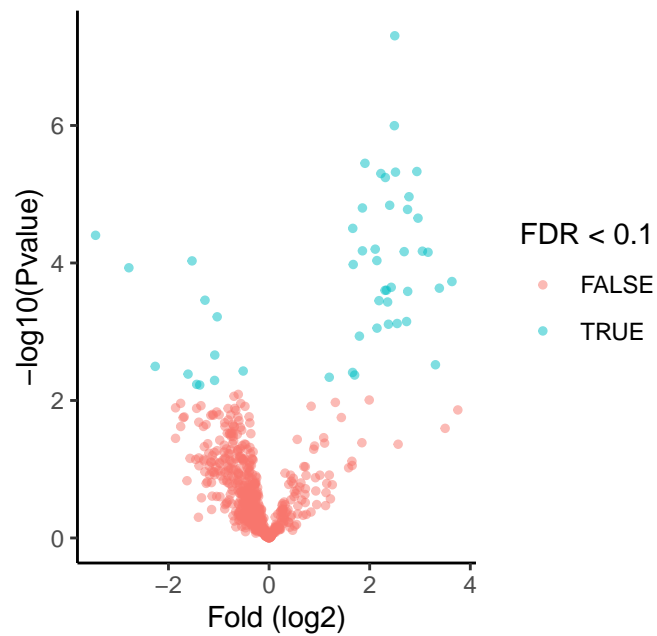


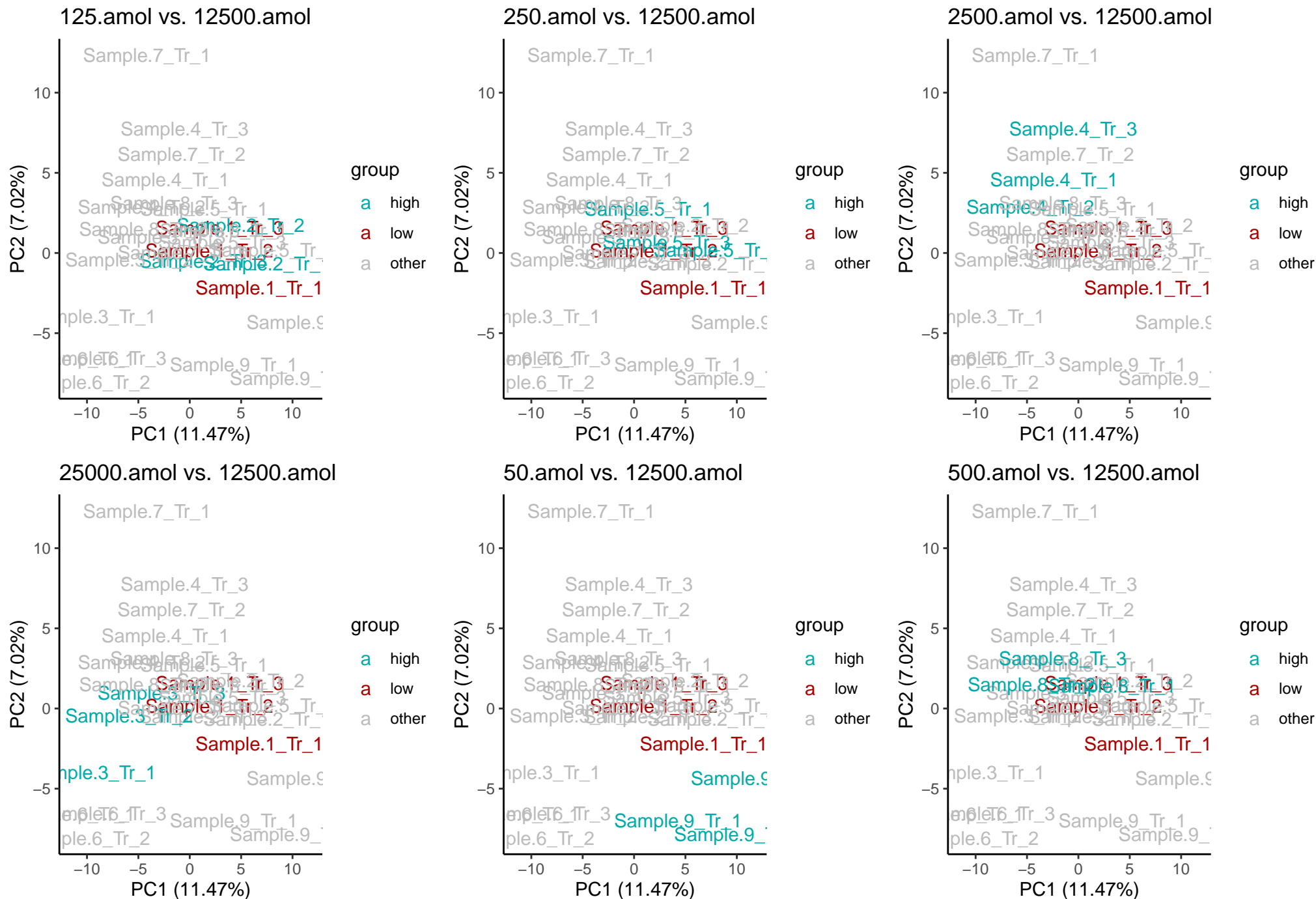


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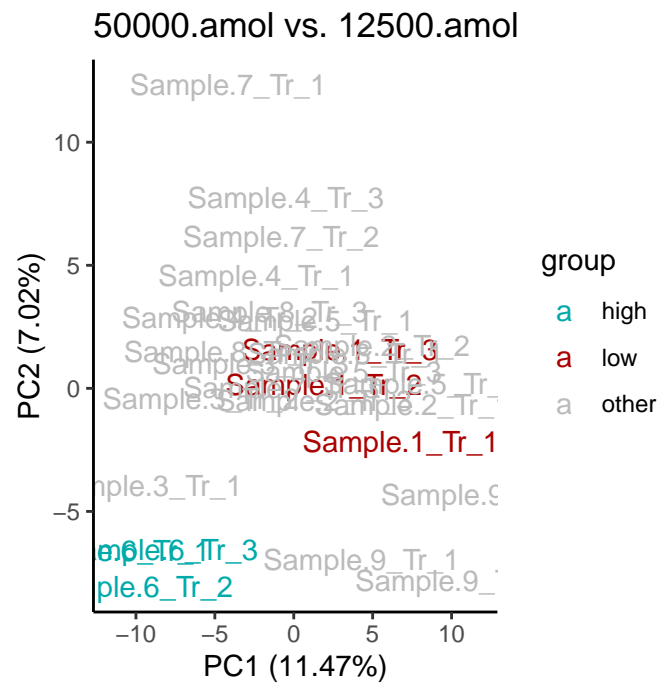
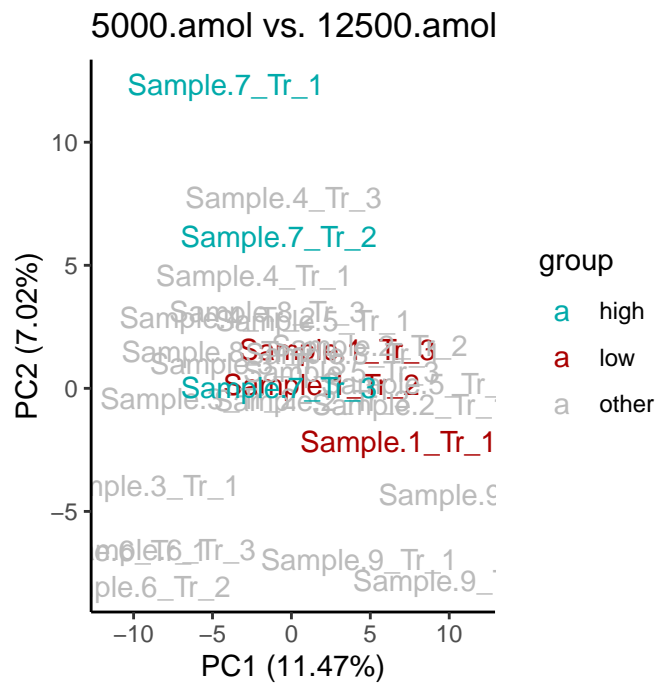


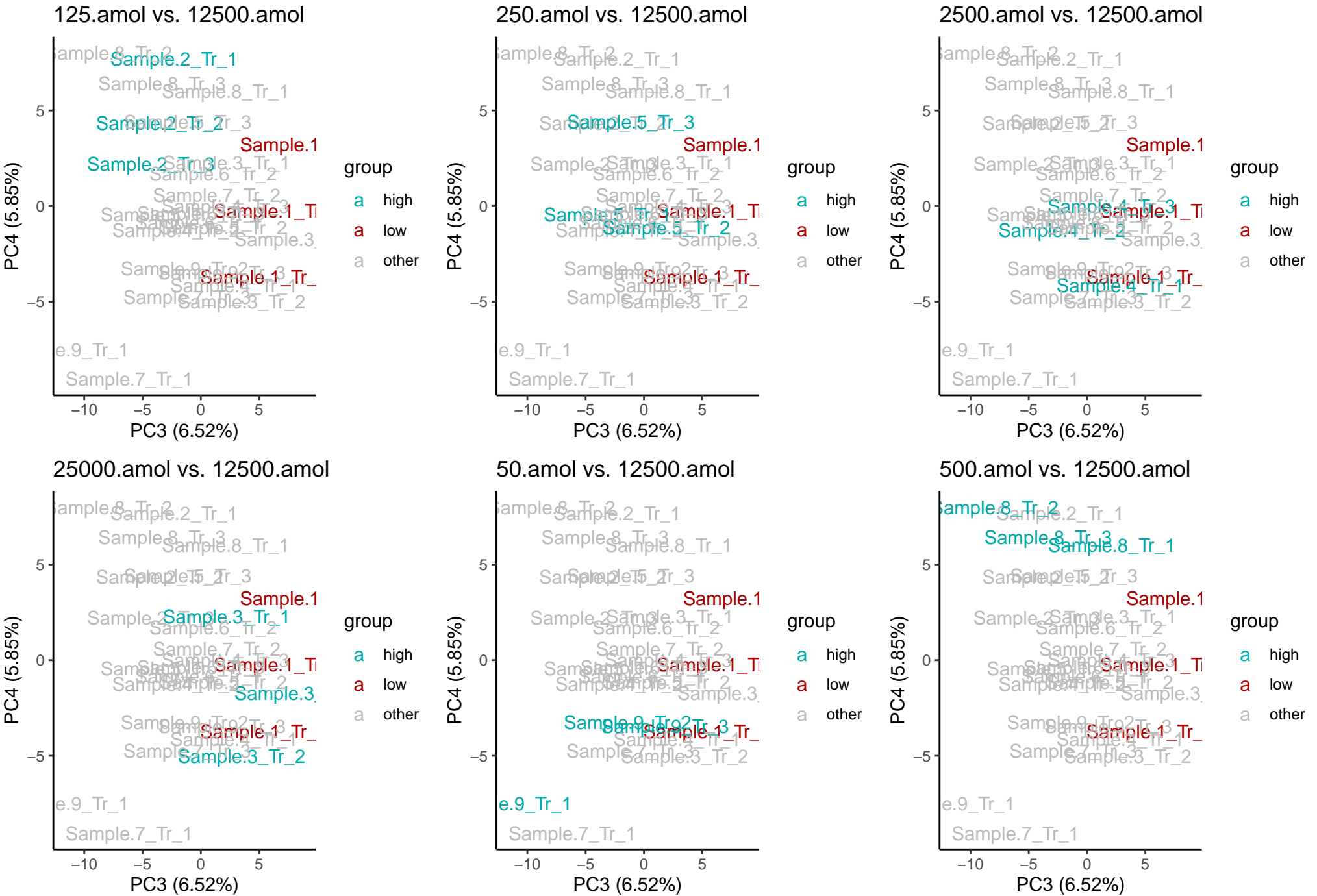
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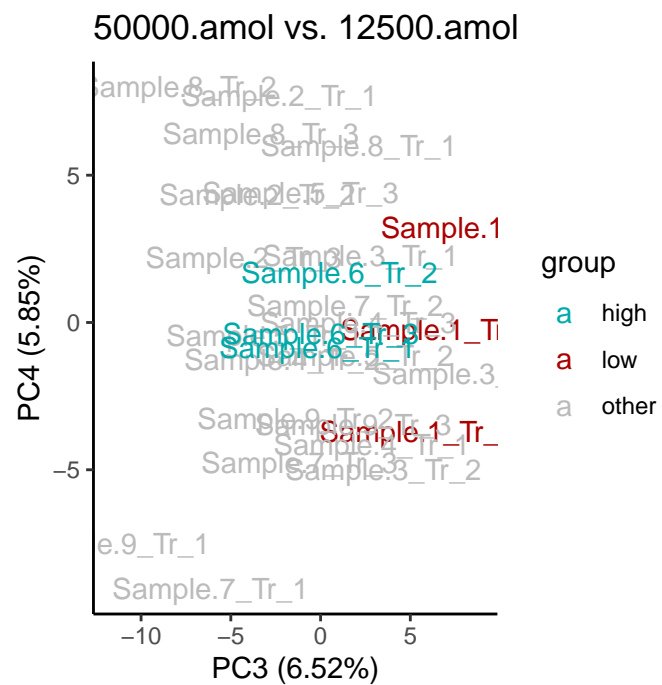
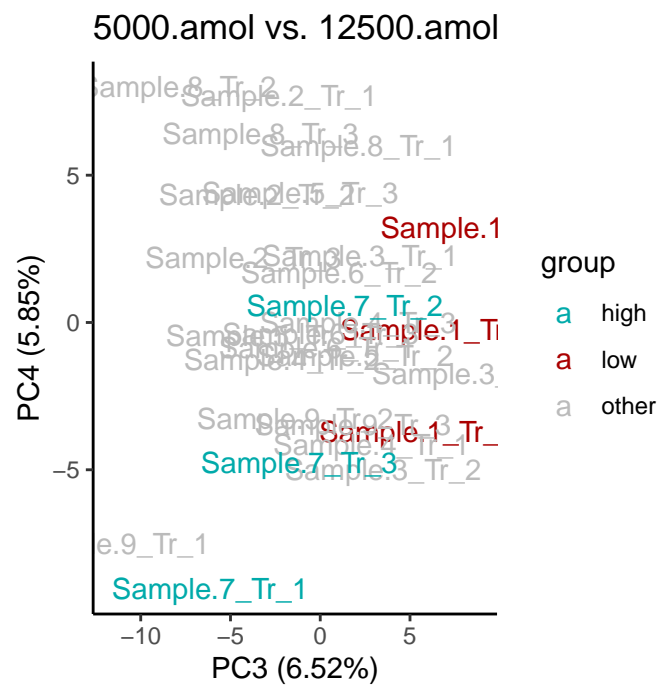




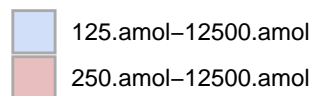
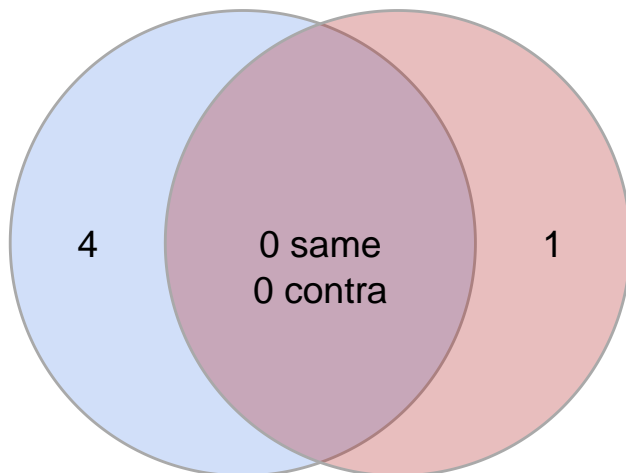




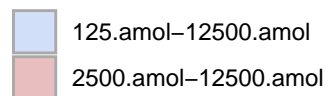
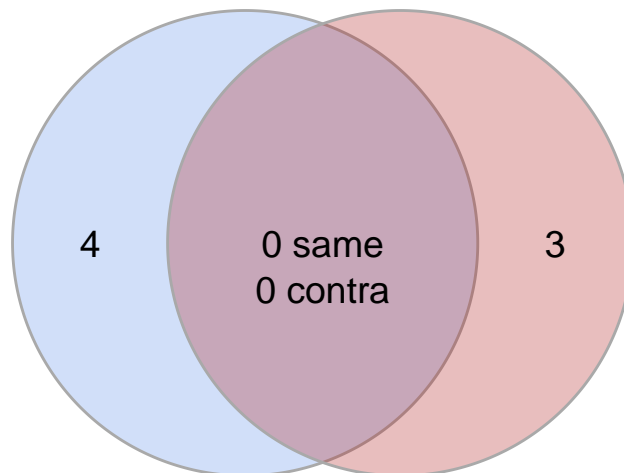




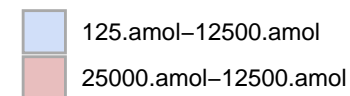
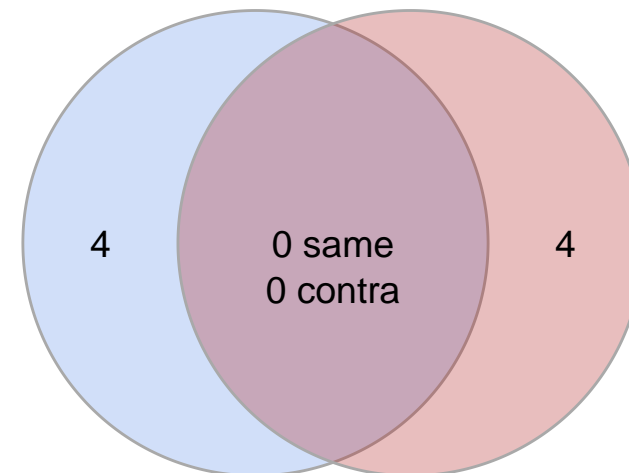
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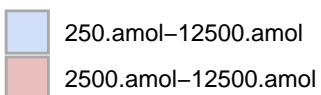
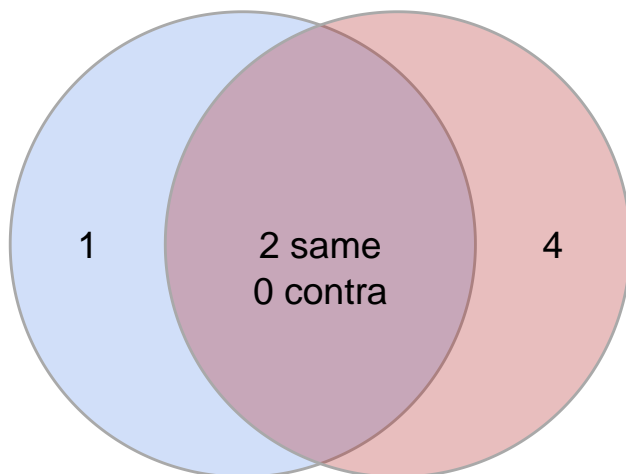
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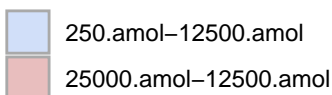
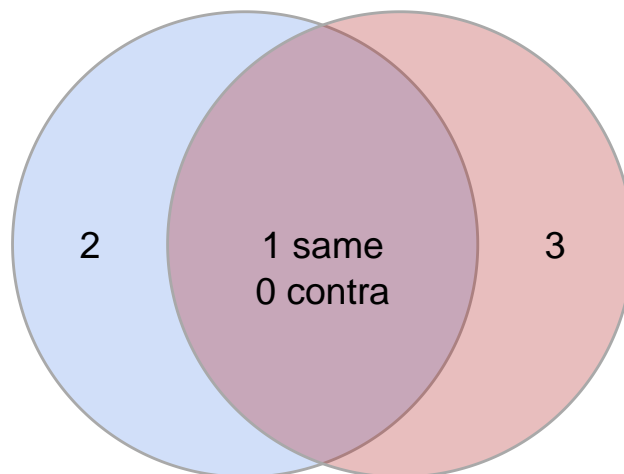
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fdr &lt; 0.1, |log2 fold| &gt;= 0



fdr &lt; 0.1, |log2 fold| &gt;= 0



fdr &lt; 0.1, |log2 fold| &gt;= 0

