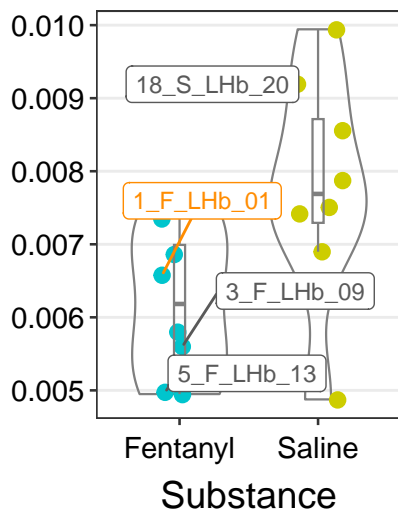
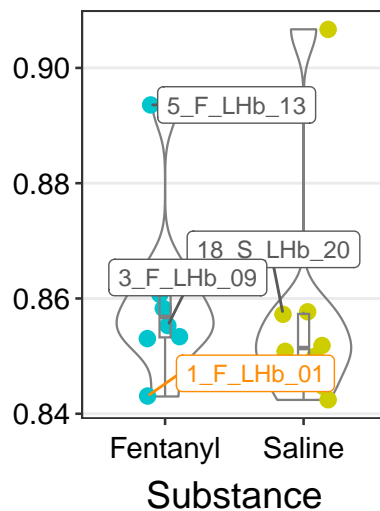


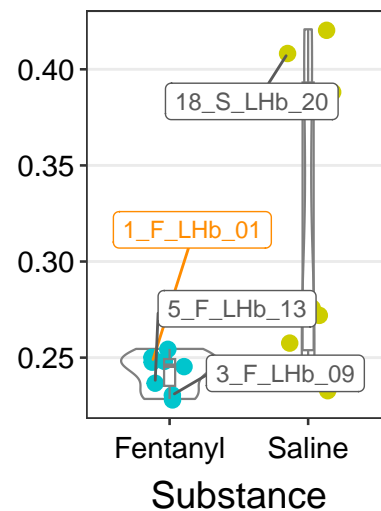
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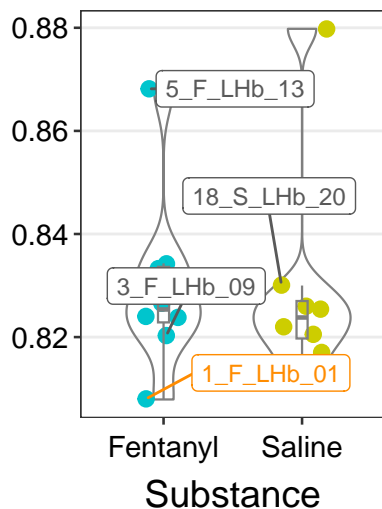
overallMapRate



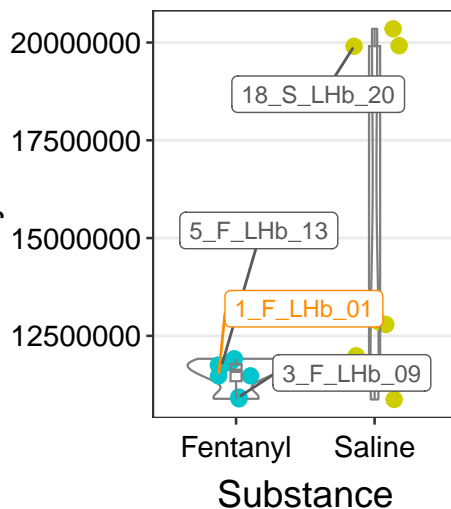
totalAssignedGene



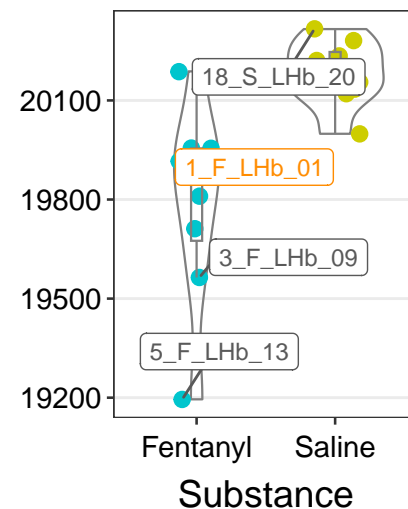
concordMapRate



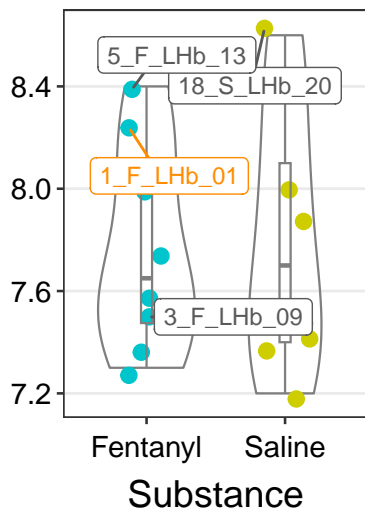
library size



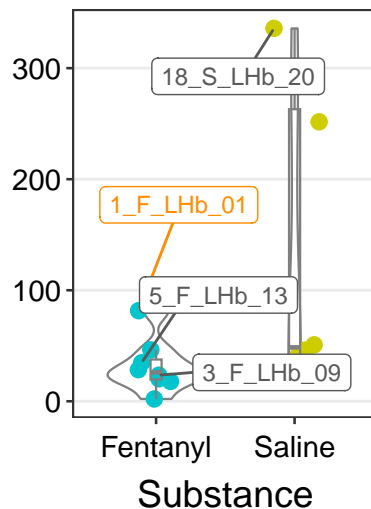
detected num genes



RIN



RNA concentration



Total RNA amount

