Implementing A Streamlined Metabolomics Data Analysis Workflow (EZMS) Based on R Programming Language*

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Abstract

A streamlined Metabolomics data analysis workflow based on R programming language was built and tested. This new workflow is efficient, all-platform-supported and open-sourced. This may release

1 Introduction

 $^{^{\}ast}$ This is part of Tu Hu's master thesis work, supervised by Lars Ove Dragsted and Gözde Gürdeniz