**[Name]: A Container-based Reproducible Untargeted Metabolomics Data Processing Pipeline for Cloud Server**

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**ABSTRACT**

***Background:***Untargeted metabolomics data processing remains a challenge due to the large size of metabolomics data files as well as the extensive steps and computational power required to process the data. Developed tools for metabolomics data processing are either local-based or web-based. Nevertheless, Local machines do not have enough computational power to process large dataset and web-based platforms needs tons of time to upload and process large data without closing the browser. Fortunately, cloud servers are available nowadays, whose configuration is good enough to process large data and all the jobs can be submitted to the server and kept running remotely even when our local machine is turned off. However, none of the untargeted metabolomics data processing tools was designed for cloud servers. Before using cloud servers, people need to specify a configuration, and better configuration costs more. Therefore, it is necessary to develop a tool specifically for cloud servers meanwhile able to guide the choice of configurations so that it will be easier for researchers to make budgets and process metabolomics data efficiently.

***Findings****:* Nextflow, a pipeline development tool supporting containerization, parallelization and high performance computing, was used to build the pipeline. Several tools and codes were connected and incorporated into Nextflow. Suggestions regarding the choice of cloud server configurations were made based on Nextflow report.

***Conclusions****:* An containerized untargeted metabolomics data processing tool specifically for cloud server was developed to facilitate metabolomics data analysis and collaborative research.

**1.1 Background**

**1.2 Findings**

**1.4 Methods**

Relationship between:

Input size and memory need

Input number of samples and memory need

**1.5 Results**

**1.5 Discussions**

**1.6 Conclusions**

**1.7 Acknowledgements**

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**1.8 Authors’ Contributions**

**1.9 Statement on conflict of interest**

The authors have no financial or personal relationships with other people or organizations that could inappropriately influence (bias) their work.

**1.9 SOURCES CITED**

**FIGURES**

Screenshot of log file

Relationship between input size and memory cost

Relationship between input number of samples and memory cost

**TABLES**

**SUPPLEMENTARY MATERIAL**