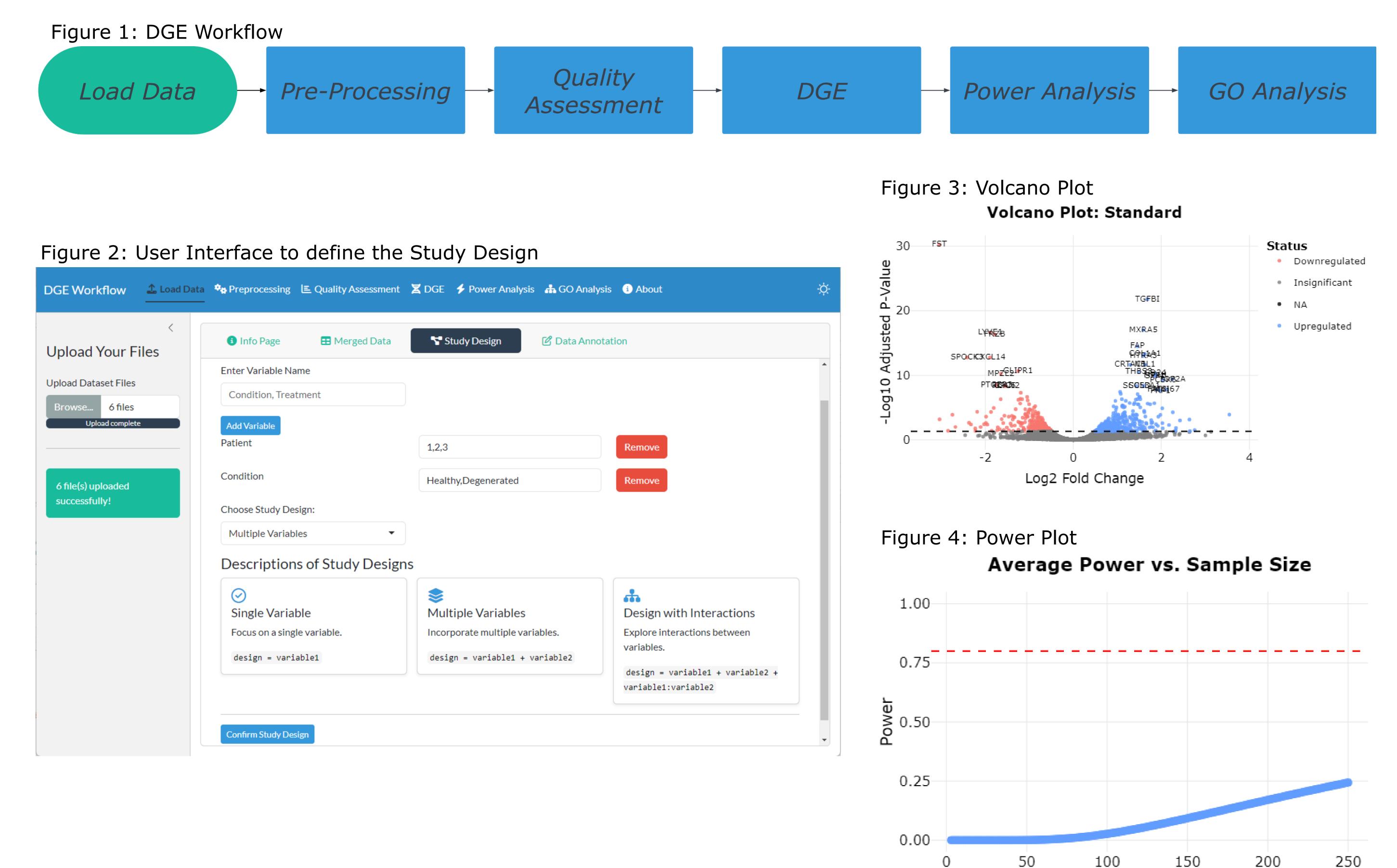


**Technik & Architektur** Master' Thesis Medical Engineering

**Master's Thesis Medical Engineering** 

# Implementation of DGE Workflow using R Shiny



#### 50 100 150 200

Sample Size (n)

#### **Problem Statement**

Differential Gene Expression (DGE) analysis is a key bioinformatics technique for unravelling biological processes. At the CC-BME, a project is dedicated to investigating gene expression differences in human knee cartilage tissue. As part of this effort, a workflow for bulk RNA-seq analysis in R was developed.

However, the current workflow is static and restricted to processing a single dataset. This limitation necessitates a more dynamic solution that empowers researchers and clinicians to analyse multiple datasets and adjust outputs interactively based on user inputs.

### **Solution Design**

An R Shiny app was developed to address the outlined problem by offering a dynamic solution. To implement this application, the existing workflow was modularized, dividing the functionality into distinct UI and server components. The core concept of reactivity was integrated, allowing for dynamic adjustments based on user inputs. Finally, the app was tested with several datasets to ensure its robustness and functionality in real-world scenarios.

### Results

The developed R Shiny application is divided into six modules, guiding users through the entire DGE analysis process. The app enables users to upload their own datasets, annotate them, and apply tailored settings to perform analyses.

Verification confirmed that the app produces accurate and reliable results across different datasets, although currently limited to human data. Moreover, it matches the performance of other similar applications while providing an added feature: a built-in power analysis tool.

## **Simon Rammerstorfer**

# Supervisor Prof Dr Fabian Ille

#### Expert

Prof PD Dr sc. ETH Philipp Stämpfli

Although several open-source applications for dynamic DGE analysis exist, they often enforce rigid workflows incompatible with the one already established. Additionally, integrating a newly developed app into the Clinical Data Cockpit at the CC-BME would enhance its utility for future research applications.

The application is designed to be userfriendly, requiring no programming skills. All visualizations and outputs update dynamically in real-time as options are modified.

**FH Zentralschweiz** 

