



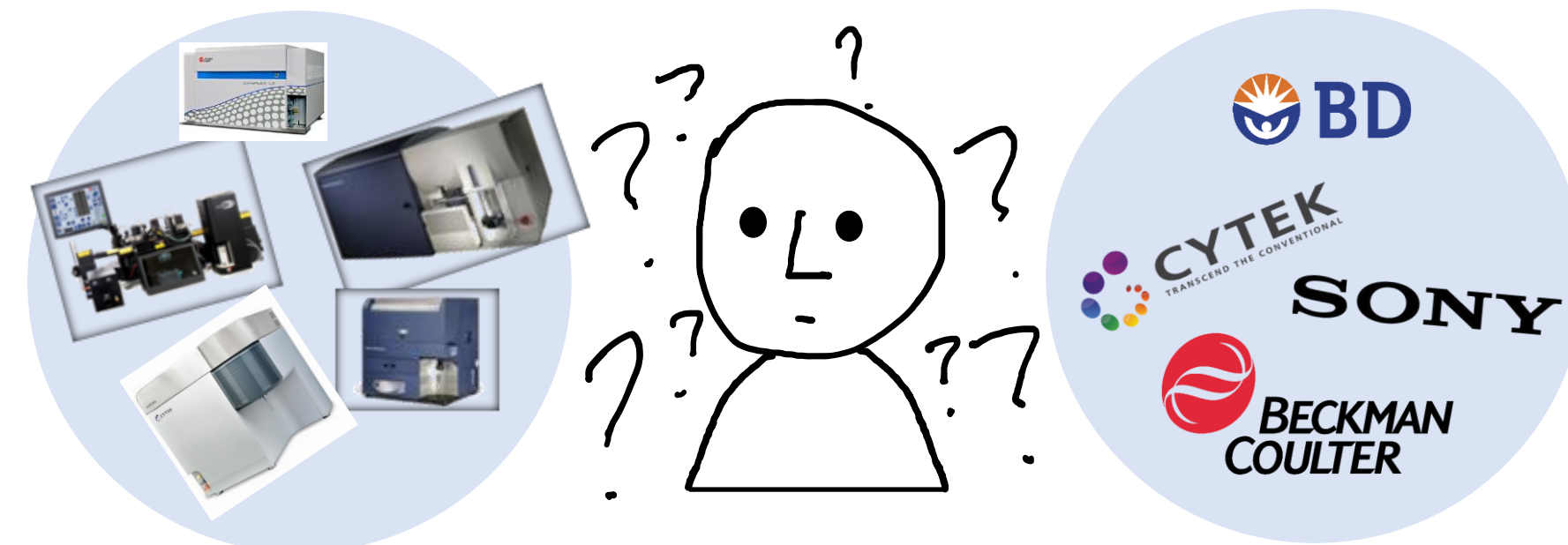
Objectives

- Develop a universal, open-source QC platform in R
- Automatically gate and calculate QC statistics from FCS files
- Display QC statistics with interactive plots to generate predictive outcomes of maintenance or repair

Why?

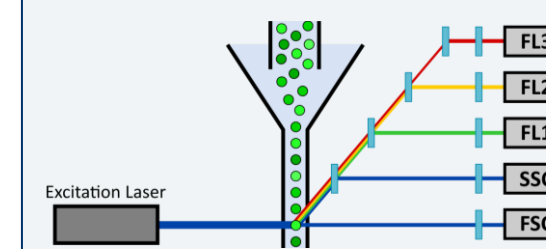
There are many different:

- Flow cytometer manufacturers
- Instruments, lasers, and detectors
- QC Beads and acquisition software



Difficult to track performance and predict problems in labs with multiple flow cytometers

Pipeline



- Run QC beads
- Export FCS files



- Read FCS files from cloud/server
- Perform k-means clustering
- Calculate summary statistics for each autogated peak
- Export raw clustered data and summary statistics



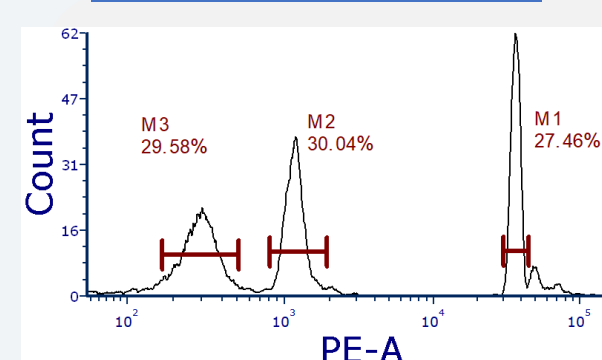
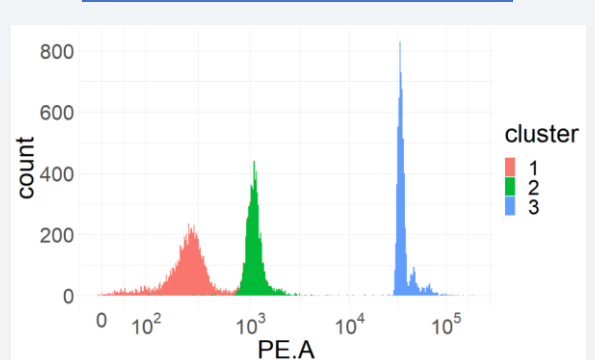
- Pull statistics and clustered raw data from cloud/server
- Create interactive plots using Plotly and Shiny.
- Export to shinyapps.io, website, or run Shiny App locally in R.

Validation

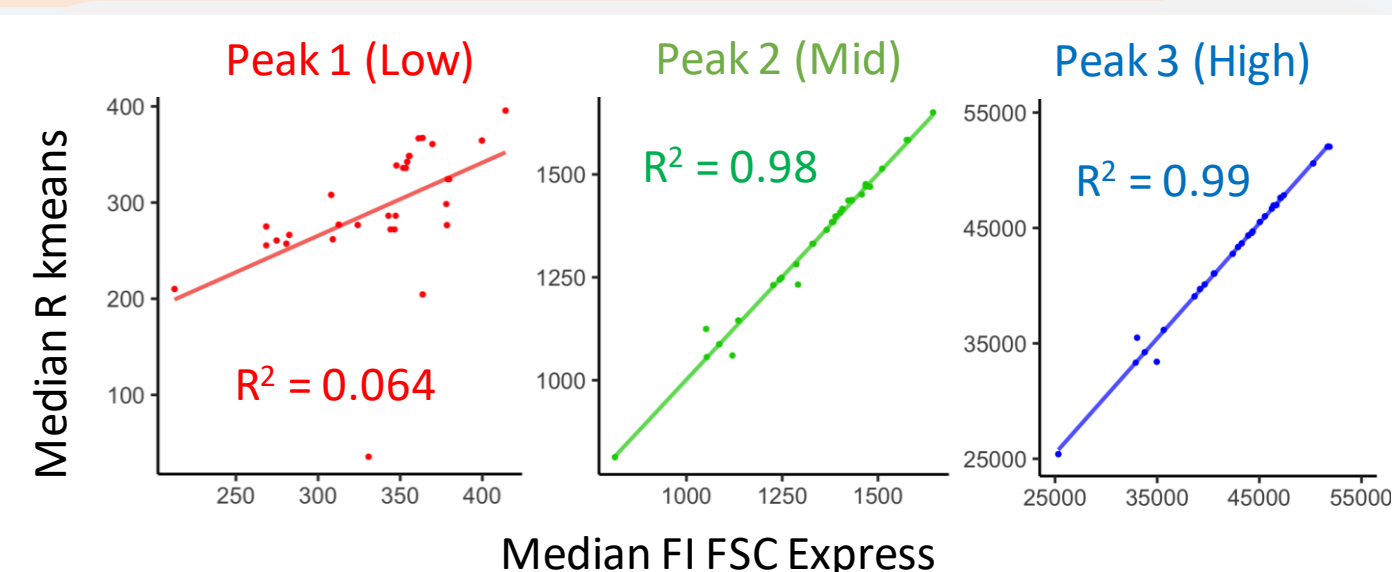
Automated vs. Manual Gating

R k-means

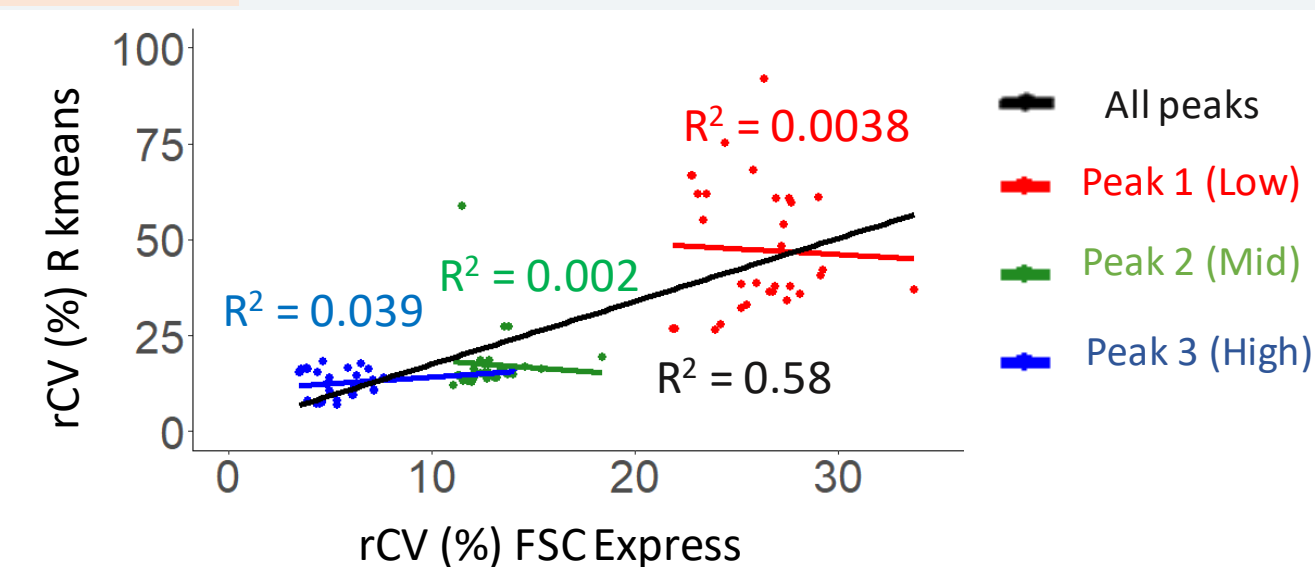
FCS Express



Median Fluorescence Intensity

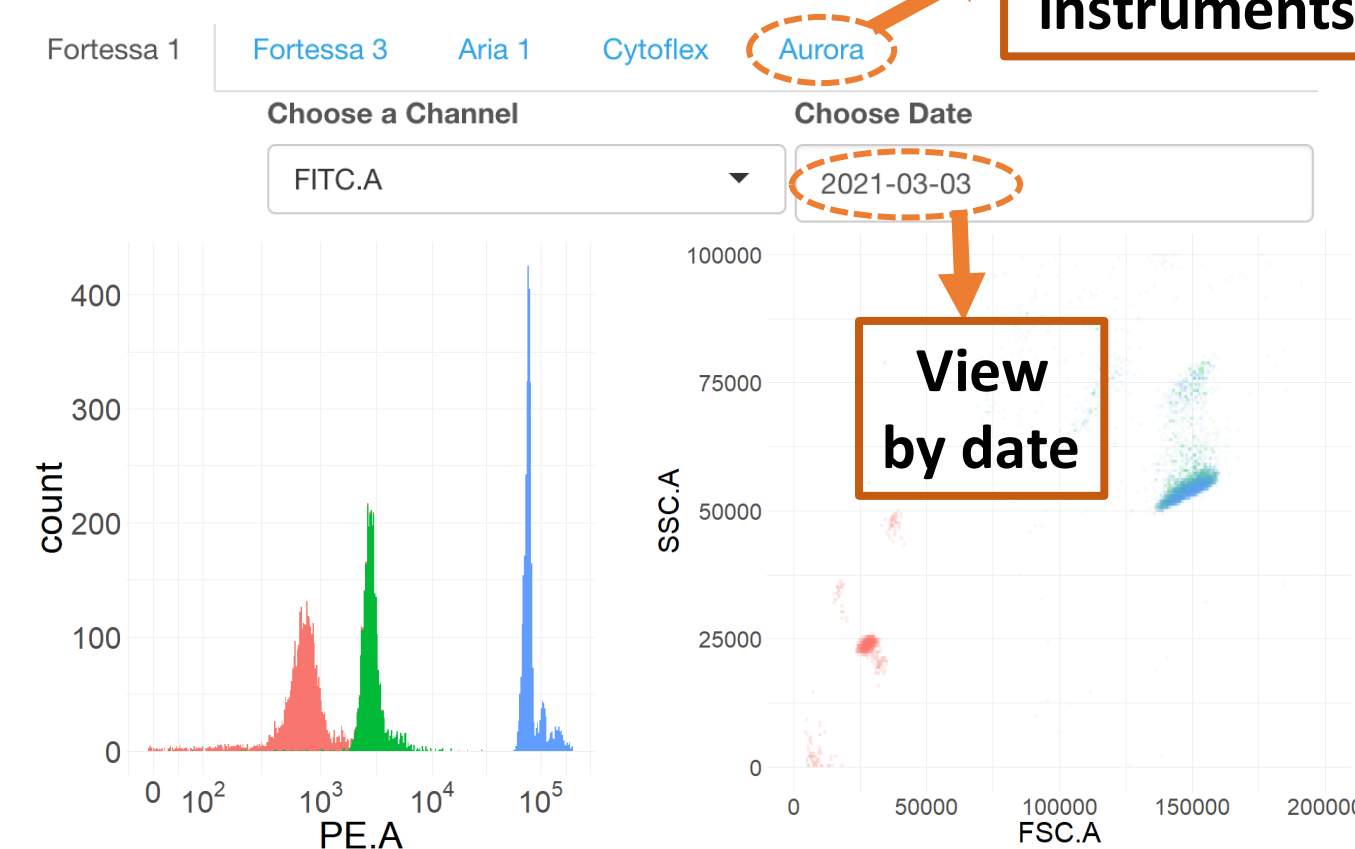


rCVs

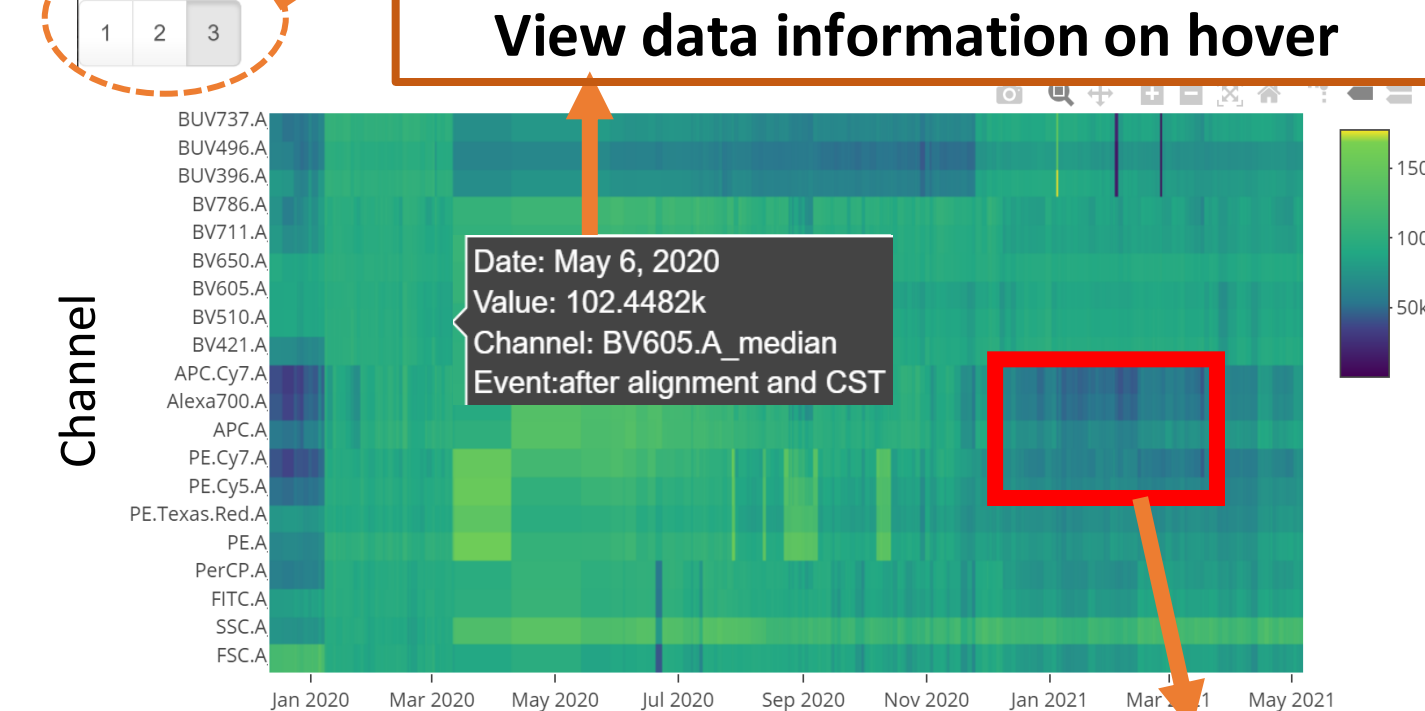


Shiny App Output

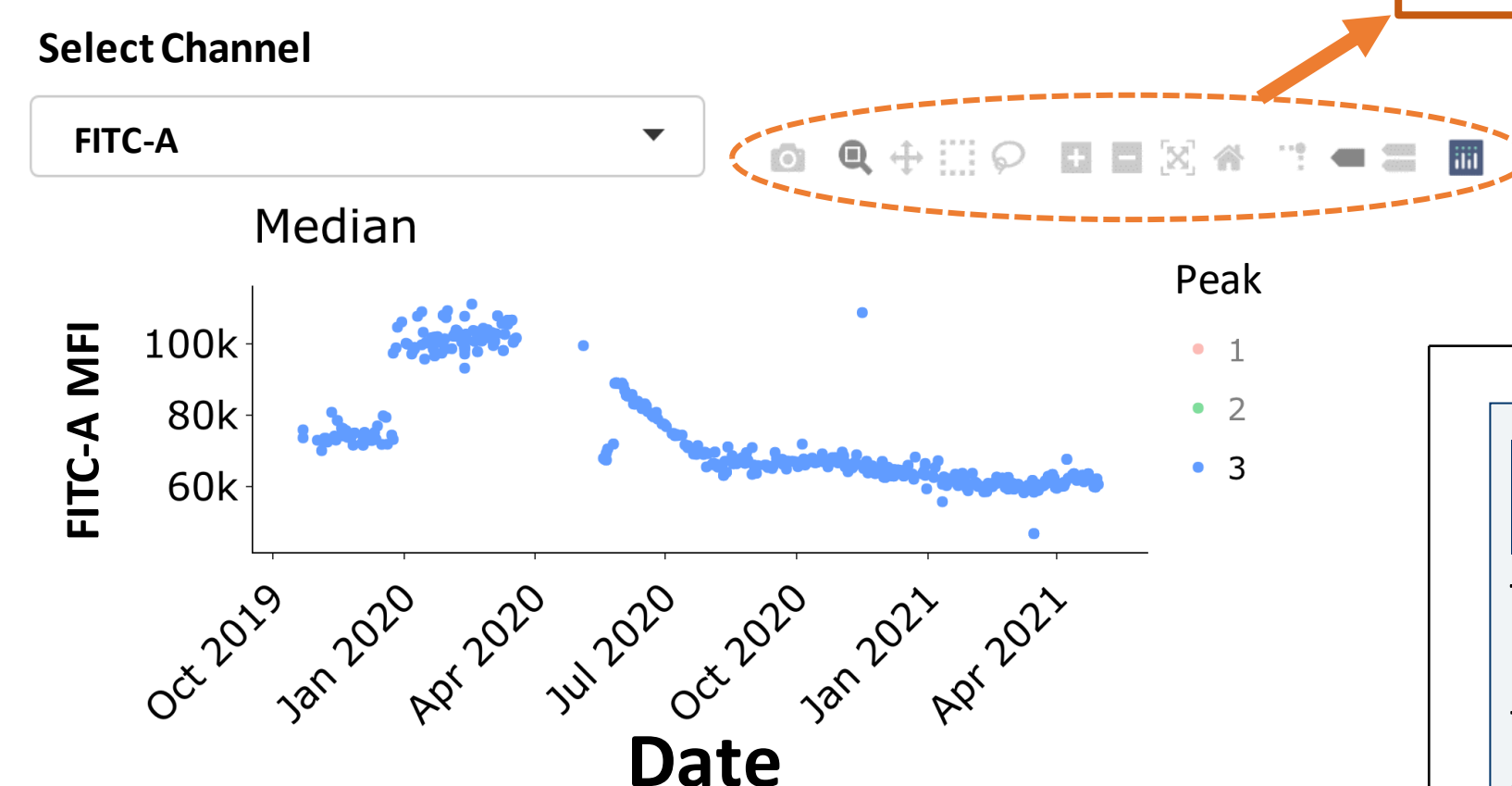
Clustered Peaks



Heatmaps



Levey-Jennings Plots



Interactive Plotly plot. Can zoom, pan, screenshot, etc.

Red laser failed and replaced, note decrease MFIs

Demo App, Poster, More info



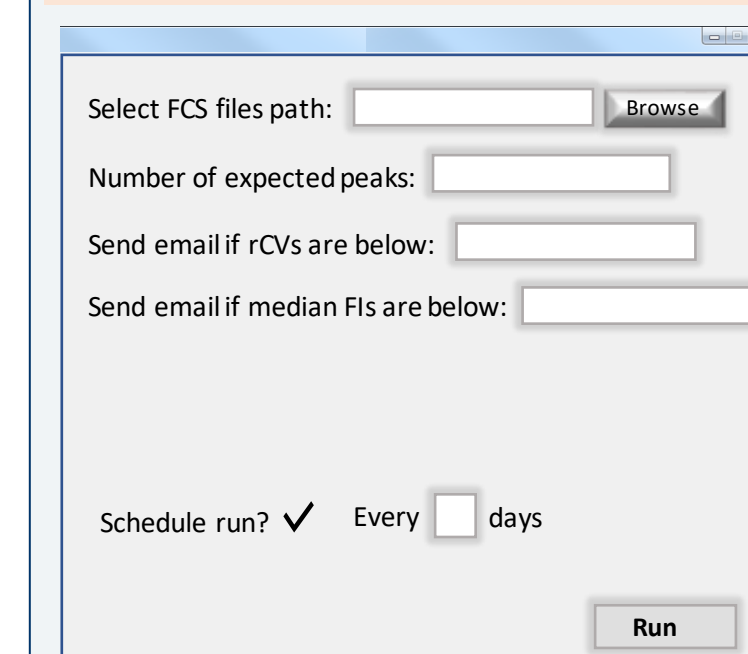
Acknowledgments

Thank you to the amazing Flow Cytometry Core team at MSK for their feedback on this project and poster



Future Work

User Friendliness



Demo model of application GUI

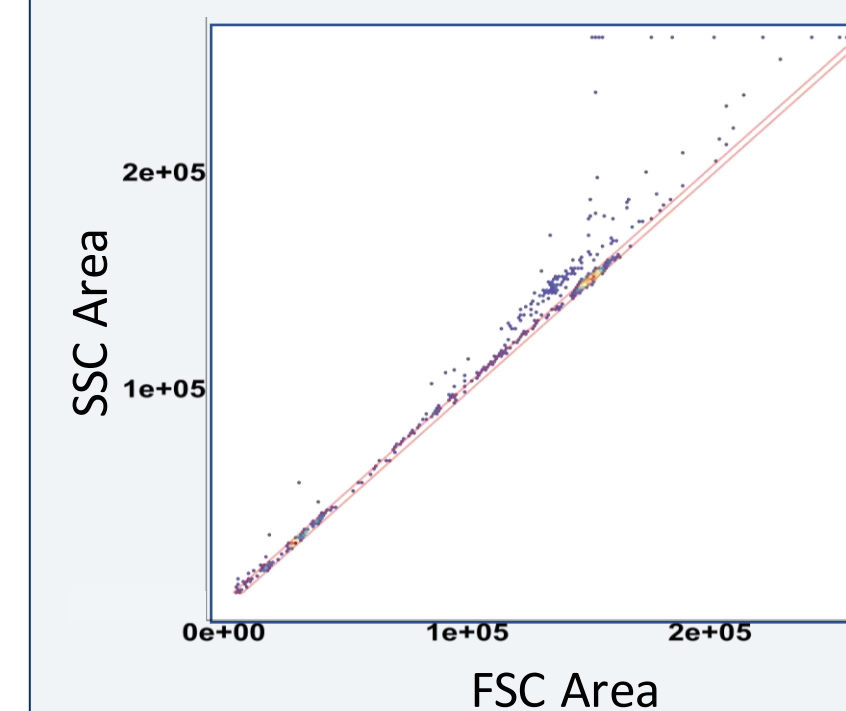
Automatic alerts for when specs (ie. rCVs, MFIs) are out of range.

Build GUI as an alternative to hard coding in R for customizing clustering and data analysis

Automate based on QC schedule

Improve Clustering

Singlet gate: flowStats package



Mixture Model Clustering: flowClust package

Assigns probability that datapoint belongs to cluster

