A bunch of random studies on cancer

Adrin Jalali

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Introduction

Here we introduce.

Background

- 2.1 Kernel Methods
- 2.2 Boosting and Ensemble Methods
- 2.3 K Best Shortest Paths

Flow Cytometry Analysis

Here we talk about flow cytometry data, how we analyze and visualize it; and how we use that analysis alongside with some machine learning tools to classify samples into cancer subtypes.

- 3.1 Flow Cytometry
- 3.2 Data Preprocessing and Challenges
- 3.3 High Dimensional Analysis and Visualization
- 3.3.1 flowType
- 3.3.2 RchyOptimyx
- 3.4 Lymphoma Diagnosis Quality Checking

Adaptive Learning

Here we talk about adaptive and interpretable methods.

- 4.1 Challenges in Cancer Data
- 4.1.1 Cancer Heterogeneity
- 4.1.2 Batch Effects and Noise
- 4.2 RatBoost
- 4.3 Raccoon

Conclusion