

Feature Selection (FS) workflow report

May 18, 2017

Introduction

The report summarizing the Feature Selection pipeline results.

Feature Selection workflow

Univariate canonical correlation (X2) with Recursive Feature Elimination (RFE) wrapped with Random Forest (RF).

Dataset

Analysis of breast cancer tumor samples using 2-color cDNA microarrays (GSE5325).

Summary stats from training phase

Table 1: Best model metrics from 10-folds cross-validation resampling.

Variables	Accuracy	Kappa	AccuracySD	KappaSD
1	0.6	0.1724	0.1127	0.2126
2	0.7	0.3915	0.1421	0.2688
3	0.7571	0.5113	0.1788	0.3498
4	0.7571	0.5155	0.1656	0.3182
5	0.7857	0.5741	0.1934	0.3738
6	0.7714	0.536	0.1536	0.2961
7	0.7714	0.5433	0.1536	0.2898
8	0.8	0.607	0.1677	0.3147
9	0.7857	0.5765	0.1543	0.2859
10	0.7857	0.5765	0.1543	0.2859
15	0.7857	0.5769	0.1934	0.3699
20	0.7857	0.5702	0.1543	0.3004
25	0.8286	0.6587	0.1475	0.2865
30	0.8143	0.6284	0.1513	0.2924
35	0.8	0.6005	0.1536	0.2985
40	0.8	0.5958	0.138	0.271
45	0.7857	0.5701	0.1388	0.2671
50	0.7857	0.5654	0.1214	0.2353
60	0.8143	0.6285	0.1656	0.3231
70	0.8143	0.6262	0.1355	0.2656
80	0.8429	0.68	0.1421	0.2826
90	0.8286	0.652	0.1313	0.2603
100	0.8429	0.6848	0.171	0.3369
1697	0.7857	0.5701	0.1388	0.2671

Summary stats from testing phase

Table 2: Classification metrics from twenty class-balanced and randomized runs.

run	Variables	Accuracy	Kappa	AccuracyPValue
1	25	0.8571	0.7107	0.0003014
2	5	0.8286	0.65	0.001202
3	35	0.8	0.595	0.003999
4	45	0.8857	0.7667	6.136e-05
5	1697	0.8571	0.7009	0.0003014
6	80	0.9143	0.8264	9.733e-06
7	6	0.8857	0.7667	6.136e-05
8	35	0.8286	0.65	0.001202
9	10	0.9143	0.8235	9.733e-06
10	1	0.6857	0.3937	0.115
11	9	0.8857	0.7627	6.136e-05
12	30	0.8857	0.7667	6.136e-05
13	25	0.8	0.5882	0.003999
14	1697	0.8286	0.6379	0.001202
15	35	0.7714	0.5333	0.01134
16	25	0.8857	0.7705	6.136e-05
17	20	0.8571	0.7107	0.0003014
18	80	0.9429	0.8814	1.128e-06
19	70	0.8	0.608	0.003999
20	15	0.8286	0.65	0.001202

Accuracy_Mean	Accuracy_SD	Accuracy_Max
0.8471	0.05883	0.9429

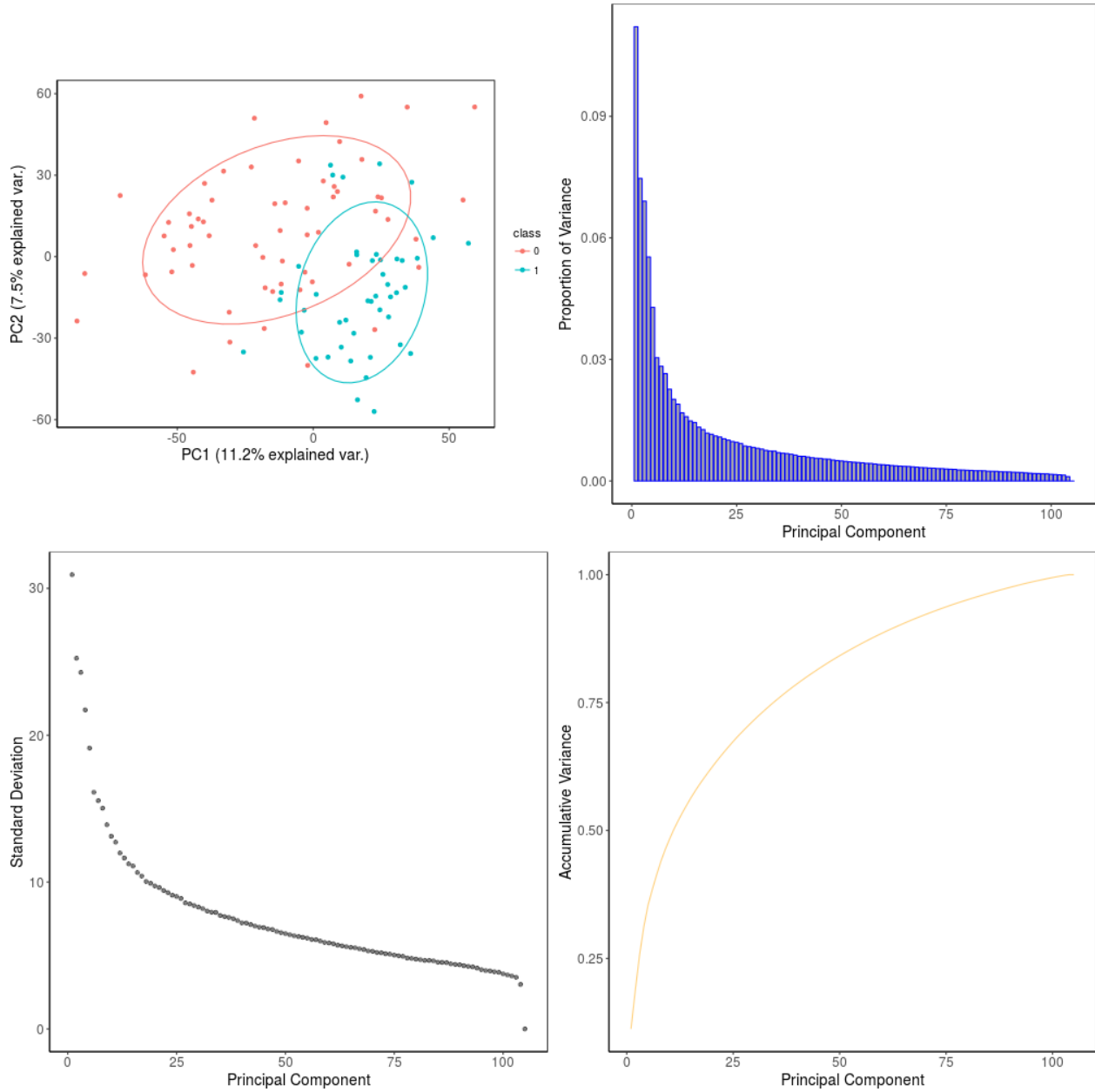
Workflow runtime

8.178 minutes

Plots

Visualization of the classification using PCA

- Groups distribution on the first two Principal Components (PC1 and PC2) from the original data (without apply any FS method).



- Groups distribution on the first two Principal Components (PC1 and PC2) after to apply the FS workflow.

