Evaluate testing data (survival) - lasso

Andrew Chang

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Contents

Labels: os_time

```
## user input
project_home <- "~/EVE/tests"
project_name <- "lasso_survival_outCV_test"
```

0. Load Data

```
300 of samples were used

100 of full features

2 runs, each run contains 3 CVs.
```
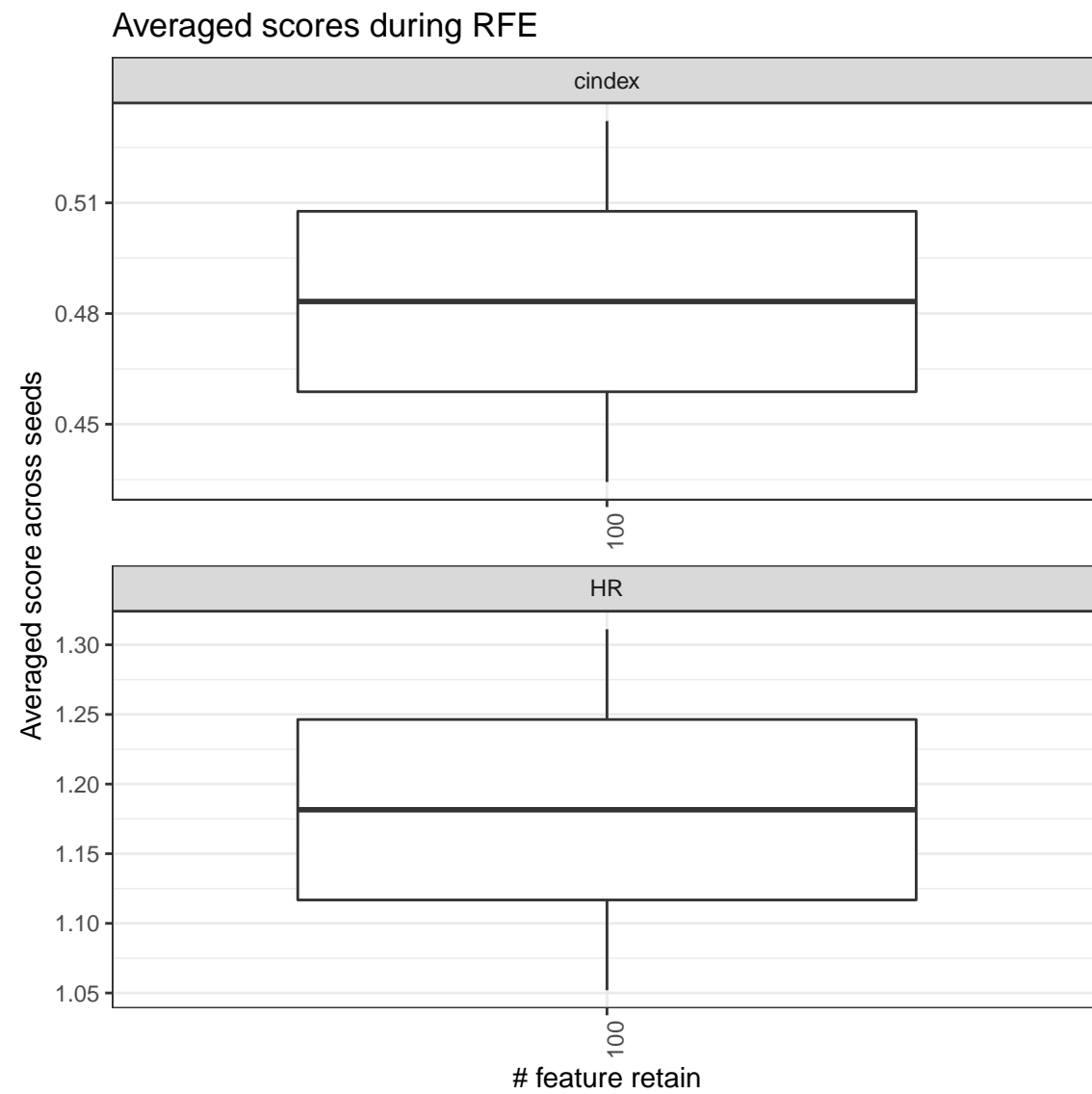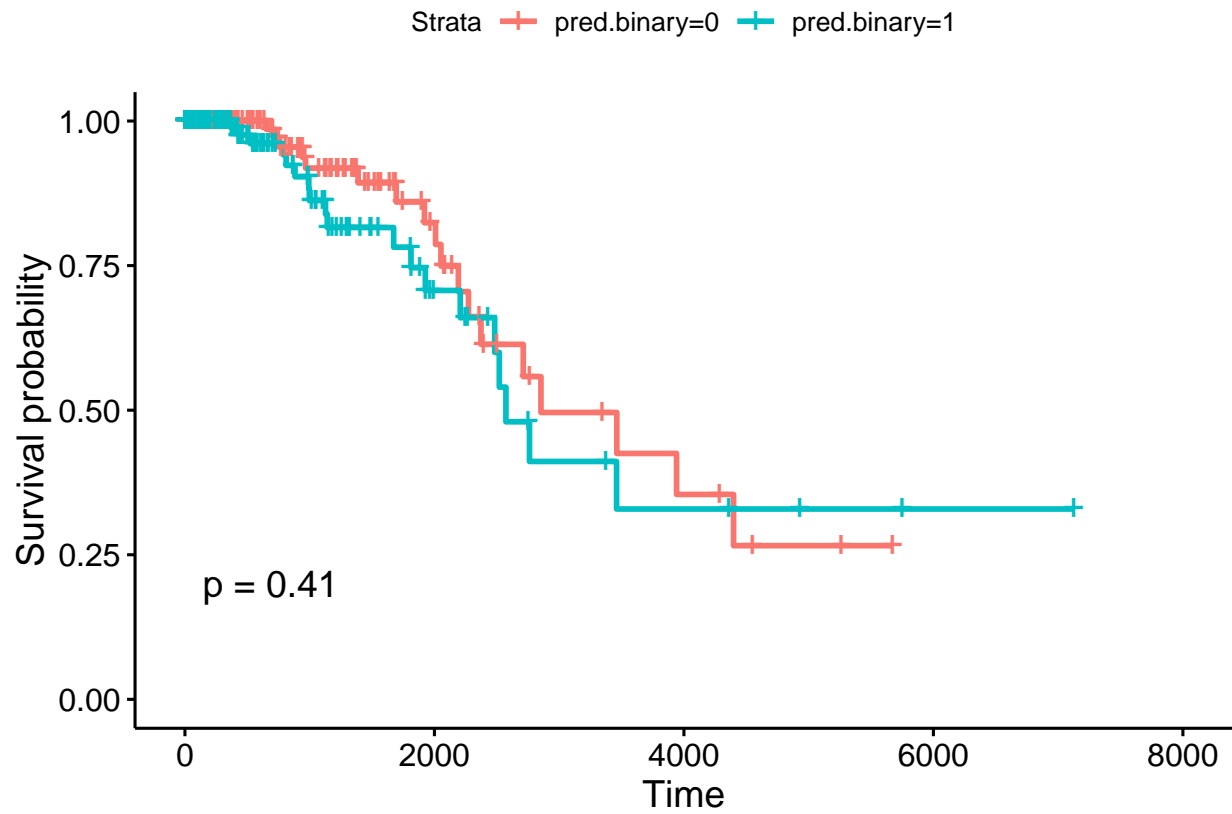
1. Scores

`average = T`: scores are based on pre-validation (combine predictions from all CVs per job and then calculate a single score per job).

`average = F`: report all the scores from each CV across entire jobs.

1. Scores

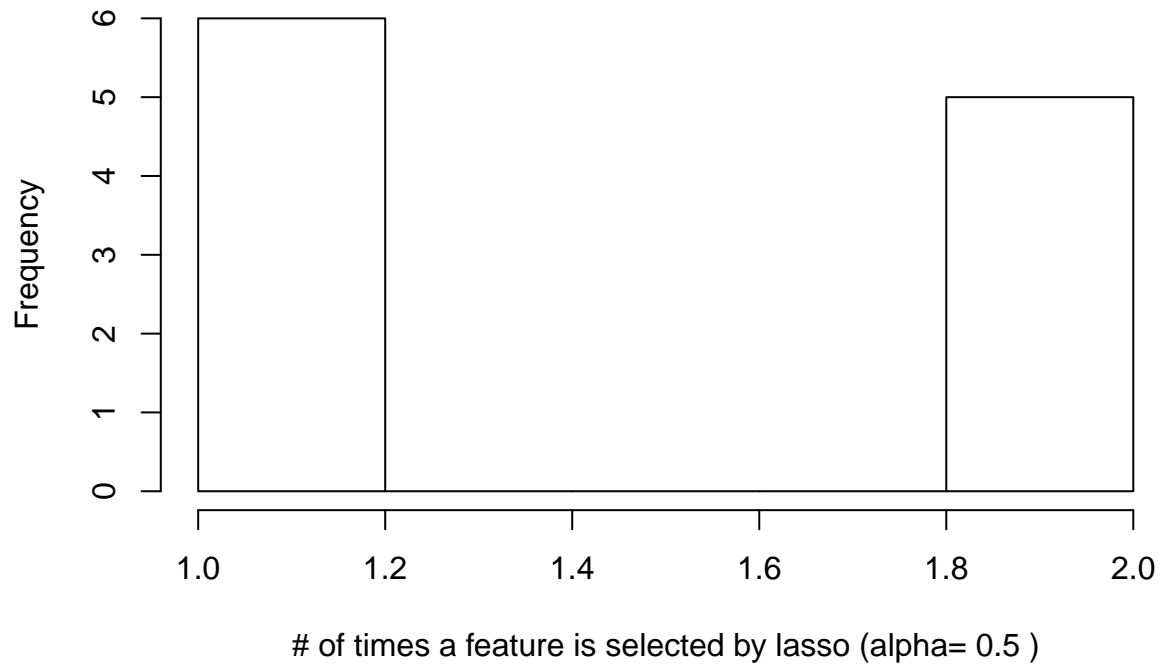## 1.1 uncalibrated average

Averaged scores during RFE

2. Important Features

For Lasso, we calculate how many times a given features is being used in all the runs.

distribution across 2 seed x 3 CV

# of times a feature is selected by lasso (alpha= 0.5)

most used features

GeneID.9635

GeneID.8424

GeneID.3872

GeneID.342667

GeneID.29127

GeneID.89846

GeneID.8914

GeneID.5366

GeneID.3852

GeneID.3838

GeneID.2173

0.0   0.5   1.0   1.5   2.0