

Module 2: Supervised Learning

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```
# If a package is installed, it will be loaded. If any
## are not, the missing package(s) will be installed
## from CRAN and then loaded.

## First specify the packages of interest
packages <- c(
  "dplyr", "PheCAP", "glmnet", "randomForestSRC", "PheNorm",
  "MAP", "pROC", "mltools", "data.table", "ggplot2", "parallel"
)

## Now load or install&load all
package.check <- lapply(
  packages,
  FUN = function(x) {
    if (!require(x, character.only = TRUE)) {
      install.packages(x, dependencies = TRUE)
      library(x, character.only = TRUE)
    }
  }
)

# load environment from example 1
load("../data/CAD_norm_pub.rda")
```

Prepare data for algorithm development

- Split data into training and testing set
- Training 106, Testing 75

```
ehr_data <- cbind(1:nrow(x), y, x)
colnames(ehr_data) <- c("patient_id", "label", colnames(x))
data <- PhecapData(ehr_data, "healthcare_utilization", "label", 75,
  patient_id = "patient_id", seed = 1234
)

# Transform Features log(x + 1)
labeled_data <- ehr_data %>% dplyr::filter(!is.na(label))

# All Features
all_x <- ehr_data %>% dplyr::select(
  starts_with("COD"), starts_with("NLP"),
```

```

    surrogate, healthcare_utilization
  )
health_count <- ehr_data$healthcare_utilization

# Training Set
train_data <- ehr_data %>% dplyr::filter(patient_id %in% data$training_set)
train_x <- train_data %>%
  dplyr::select(
    starts_with("COD"), starts_with("NLP"),
    surrogate, healthcare_utilization
  ) %>%
  as.matrix()
train_y <- train_data %>%
  dplyr::select(label) %>%
  pull()

# Testing Set
test_data <- ehr_data %>% dplyr::filter(patient_id %in% data$validation_set)
test_x <- test_data %>%
  dplyr::select(
    starts_with("COD"), starts_with("NLP"),
    surrogate, healthcare_utilization
  ) %>%
  as.matrix()
test_y <- test_data %>%
  dplyr::select(label) %>%
  pull()

```

Penalized logistic regression

- Fit LASSO and Adaptive LASSO(ALASSO)

```

# Choose best lambda using CV
beta.lasso <- lasso_fit(x = train_x, y = train_y,
  tuning = "cv", family = "binomial")

# Features Selected
names(beta.lasso[abs(beta.lasso)>0])[-1]

## [1] "NLP20" "NLP288" "NLP304"
## [4] "NLP405" "surrogate" "healthcare_utilization"

# prediction on testing set
y_hat.lasso <- linear_model_predict(beta = beta.lasso, x = test_x,
  probability = TRUE)

# Fit Adaptive LASSO
beta.alasso <- adaptive_lasso_fit(x = train_x, y = train_y,
  tuning = "cv", family = "binomial")
y_hat.alasso <- linear_model_predict(beta = beta.alasso, x = test_x,
  probability = TRUE)

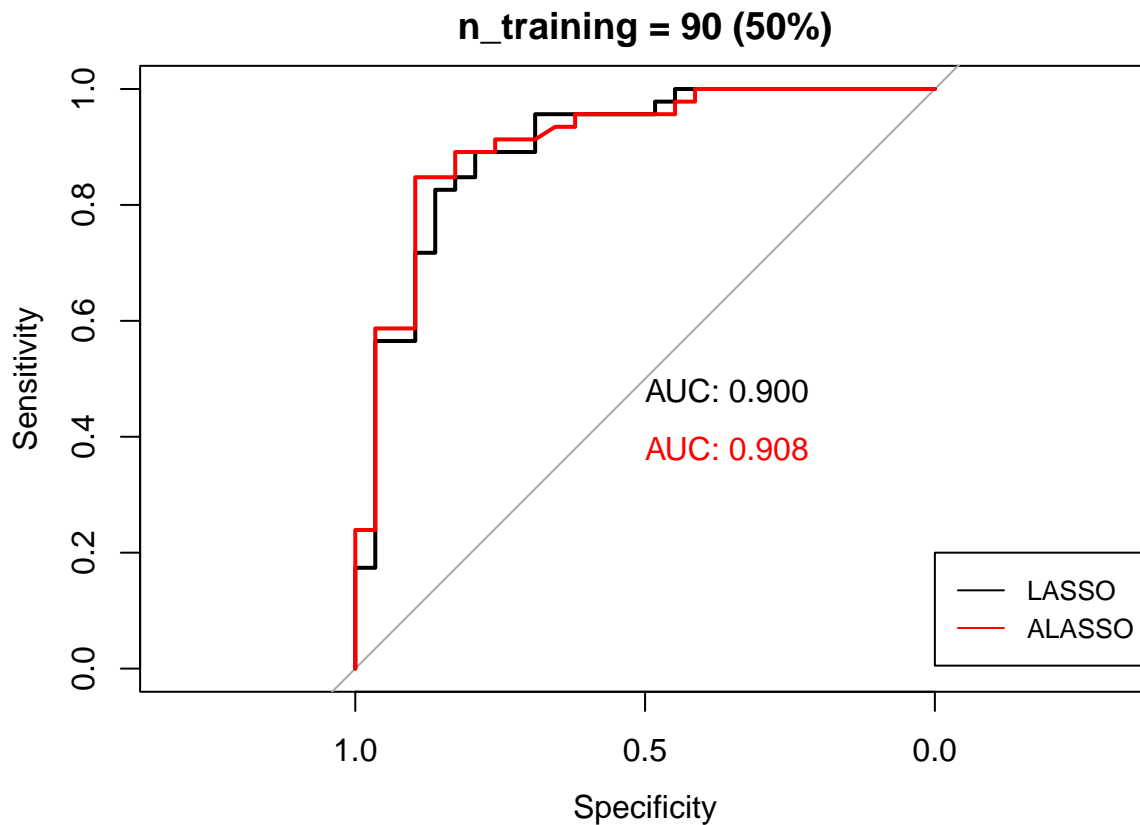
# Features Selected
names(beta.alasso[abs(beta.alasso)>0])[-1]

```

```
## [1] "NLP304"                "surrogate"                "healthcare_utilization"

roc.lasso <- roc(test_y, y_hat.lasso)
roc.lasso <- roc(test_y, y_hat.lasso)

plot(roc.lasso,
     print.auc = TRUE, main = "n_training = 90 (50%)")
)
plot(roc.lasso,
     print.auc = TRUE, col = 'red', add = TRUE, print.auc.y = 0.4
)
legend(0, 0.2, legend = c("LASSO", "ALASSO"), col = c("black", "red"),
      lty = 1, cex = 0.8)
```



```
roc_full.lasso <- get_roc(y_true = test_y, y_score = y_hat.lasso)
head(roc_full.lasso, 10)
```

##	cutoff	pos.rate	FPR	TPR	PPV	NPV	F1
## [1,]	0.9469064	0.006666667	0.00000000	0.07729469	1.0000000	0.4059098	0.1434978
## [2,]	0.9246943	0.08000000	0.00000000	0.13333333	1.0000000	0.4211036	0.2352941
## [3,]	0.9080636	0.12000000	0.03448276	0.20521739	0.9042146	0.4337051	0.3345145
## [4,]	0.9060840	0.12000000	0.03448276	0.31869565	0.9361430	0.4718571	0.4755109
## [5,]	0.8782036	0.25333333	0.03448276	0.43217391	0.9521073	0.5173688	0.5944976
## [6,]	0.8292790	0.34666667	0.03448276	0.54565217	0.9616858	0.5725971	0.6962552
## [7,]	0.8202000	0.37333333	0.06896552	0.56521739	0.9285714	0.5744681	0.7027027
## [8,]	0.8194223	0.37333333	0.06896552	0.56521739	0.9285714	0.5744681	0.7027027
## [9,]	0.8186445	0.37333333	0.06896552	0.56521739	0.9285714	0.5744681	0.7027027
## [10,]	0.8139544	0.38666667	0.10344828	0.58195652	0.8992274	0.5748397	0.7066121

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