

## Task 4

**Task 4:** Use 5-fold cross-validation to select the best  $\lambda$ . Compare the prediction performance between the “optimal” model and “full” model.

### 5-fold CV

We write an **R** function `cv.logit.lasso` to conduct 5-fold cross-validation to select the best  $\lambda$ .

```
cv.logit.lasso <- function(x, y, nfolds = 5, lambda) {
  auc <- data.frame(matrix(ncol = 3, nrow = 0))
  folds <- createFolds(y, k = nfolds)
  for (i in 1:nfolds) {
    valid_index <- folds[[i]]
    x_training <- x[-valid_index, ]
    y_training <- y[-valid_index]
    training_dat <- data.frame(cbind(y_training, x_training))
    x_valid <- cbind(rep(1, length(valid_index)), x[valid_index, ])
    y_valid <- y[valid_index]
    res <- LogisticLASSO(dat = training_dat, start = rep(0, ncol(training_dat)), lambda = lambda)
    for (k in 1:nrow(res)) {
      betavec <- res[k, 2:ncol(res)]
      u_valid <- x_valid %*% betavec
      phat_valid <- sigmoid(u_valid)[, 1]
      roc <- roc(response = y_valid, predictor = phat_valid)
      auc <- rbind(auc, c(lambda[k], i, roc$auc[1]))
    }
  }
  colnames(auc) <- c("lambda", "fold", "auc")
  cv_res <- auc %>%
    group_by(lambda) %>%
    summarize(auc_mean = mean(auc)) %>%
    mutate(auc_ranking = min_rank(desc(auc_mean)))
  bestlambda <- min(cv_res$lambda[cv_res$auc_ranking == 1])
  return(cv_res)
}
```

Compare the results of cross-validation using `glmnet` and using our algorithm.

1. Our function `cv.logit.lasso`:

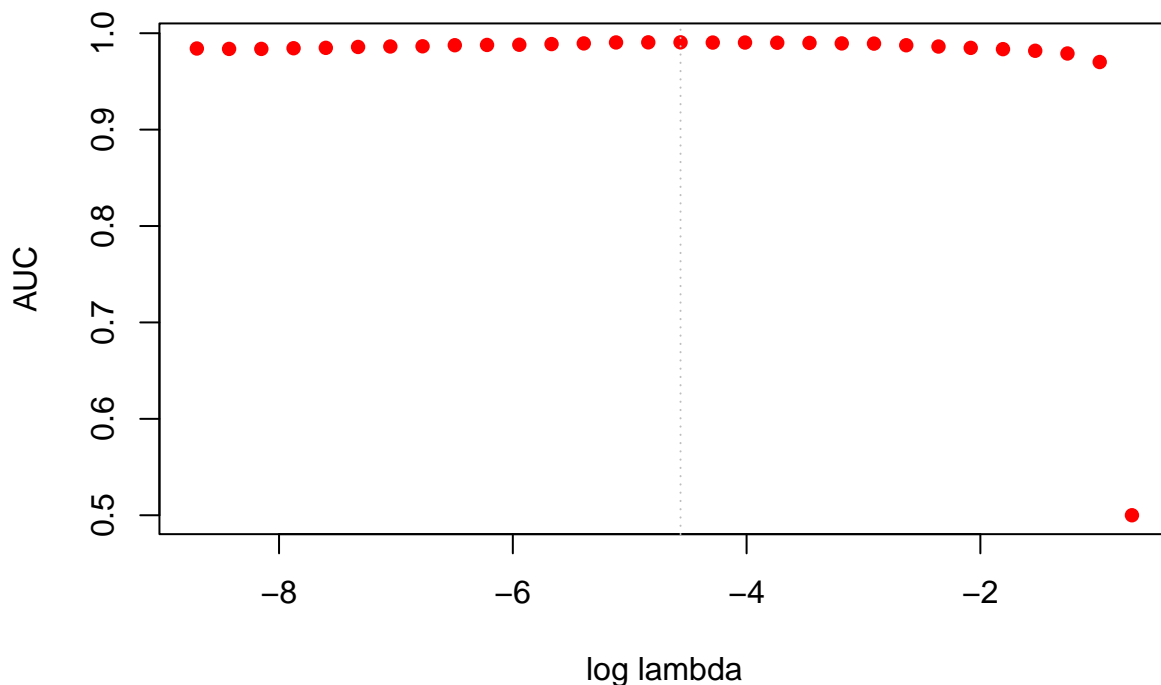
```
lambda_max <- max(abs(t(x) %*% (y - mean(y)))) / floor(length(y) * 4/5) # cv trains model on 4/5 of the
lambdas <- exp(seq(log(lambda_max), log(lambda_max) - 8, length = 30))
set.seed(1)
res_cv = cv.logit.lasso(x, y, nfolds = 5, lambda = lambdas)
as.matrix(res_cv %>% arrange(-lambda))
```

```
##           lambda  auc_mean auc_ranking
## [1,] 0.4950042193 0.5000000          30
## [2,] 0.3756674150 0.9701369          29
## [3,] 0.2851006137 0.9789711          28
## [4,] 0.2163678740 0.9817904          27
## [5,] 0.1642053881 0.9834819          26
## [6,] 0.1246183593 0.9848048          21
## [7,] 0.0945750664 0.9862041          18
## [8,] 0.0717746826 0.9875328          15
## [9,] 0.0544710700 0.9892431          10
## [10,] 0.0413390538 0.9894404           8
## [11,] 0.0313729356 0.9899379           7
## [12,] 0.0238094730 0.9901358           6
## [13,] 0.0180694282 0.9903346           4
## [14,] 0.0137132071 0.9903339           5
## [15,] 0.0104071942 0.9905325           1
## [16,] 0.0078982029 0.9905317           2
## [17,] 0.0059940852 0.9904323           3
## [18,] 0.0045490168 0.9894383           9
## [19,] 0.0034523289 0.9887423          11
## [20,] 0.0026200332 0.9880455          12
## [21,] 0.0019883893 0.9878508          13
## [22,] 0.0015090236 0.9875498          14
## [23,] 0.0011452246 0.9864511          16
## [24,] 0.0008691311 0.9862492          17
## [25,] 0.0006595989 0.9857439          19
## [26,] 0.0005005812 0.9848421          20
## [27,] 0.0003798999 0.9844401          22
## [28,] 0.0002883127 0.9837394          25
## [29,] 0.0002188056 0.9837398          24
## [30,] 0.0001660554 0.9841404          23
```

```
# best lambda
best_lambda <- max(res_cv$lambda[res_cv$auc_ranking == 1])
best_lambda
```

```
## [1] 0.01040719
```

```
plot(log(res_cv$lambda),
     res_cv$auc_mean,
     pch = 16,
     xlab = "log lambda",
     ylab = "AUC",
     col = "red")
abline(v = log(res_cv$lambda[which((res_cv$auc_ranking == 1))]), col = "gray", lty = 3)
```



```
# coefficients of the best model
res_coef <- LogisticLASSO(dat = Training, start = rep(0, ncol(Training)),
                          lambda = lambdas) %>% as.data.frame
res_coef[res_coef$lambda == best_lambda, -1]

## (Intercept) radius_mean texture_mean perimeter_mean area_mean
## 15 -0.6314292      0      0.3808524      0      0
## smoothness_mean compactness_mean concavity_mean concave.points_mean
## 15      0      0      0      0.4863225
## symmetry_mean fractal_dimension_mean radius_se texture_se perimeter_se
## 15      0      0 0.6822552      0      0
## area_se smoothness_se compactness_se concavity_se concave.points_se
## 15      0      0      0      0      0
## symmetry_se fractal_dimension_se radius_worst texture_worst perimeter_worst
## 15      0      -0.0145589      2.534049      0.4297598      0
## area_worst smoothness_worst compactness_worst concavity_worst
## 15      0      0.474963      0      0.1166963
## concave.points_worst symmetry_worst fractal_dimension_worst
## 15      1.000594      0.3539412      0
```

## 2. glmnet from R package caret

```
set.seed(1)
fit.logit.lasso <- cv.glmnet(x, y,
                             nfolds = 5, alpha = 1,
```

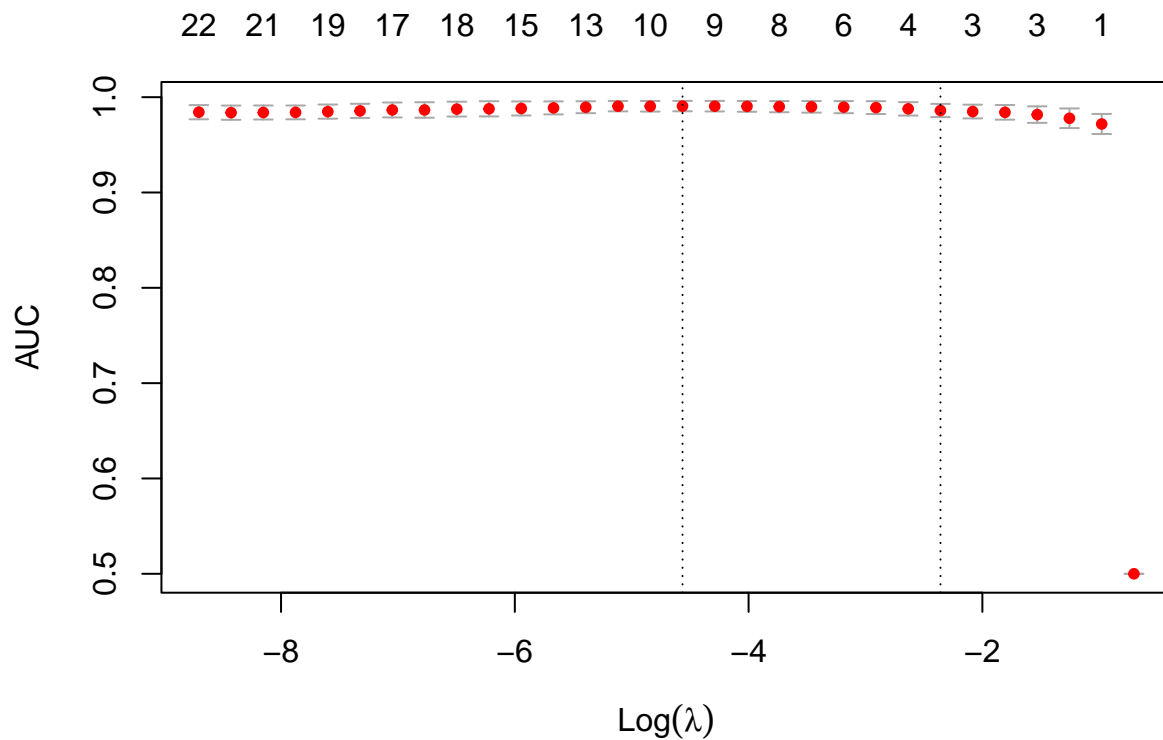
```

lambda = lambdas,
family = "binomial", type.measure = "auc")
# best lambda
fit.logit.lasso$lambda.min

```

```
## [1] 0.01040719
```

```
plot(fit.logit.lasso)
```



```

# coefficients of the best model
coef(fit.logit.lasso, fit.logit.lasso$lambda.min)

```

```

## 31 x 1 sparse Matrix of class "dgCMatrix"
##                               s1
## (Intercept)                 -0.6155278
## radius_mean                  .
## texture_mean                 0.3680397
## perimeter_mean               .
## area_mean                    .
## smoothness_mean              .
## compactness_mean             .
## concavity_mean               .
## concave.points_mean          0.4396486
## symmetry_mean                .

```

```
## fractal_dimension_mean .
## radius_se 0.7840213
## texture_se .
## perimeter_se .
## area_se .
## smoothness_se .
## compactness_se .
## concavity_se .
## concave.points_se .
## symmetry_se .
## fractal_dimension_se -0.0167962
## radius_worst 2.5469005
## texture_worst 0.4561346
## perimeter_worst .
## area_worst .
## smoothness_worst 0.4813888
## compactness_worst .
## concavity_worst 0.1024629
## concave.points_worst 1.0701080
## symmetry_worst 0.3587025
## fractal_dimension_worst .
```

The results are slightly different (mean AUC values).

```
tibble(
  lambda = lambdas,
  ours_AUC = res_cv %>% arrange(-lambda) %>% .$auc_mean,
  cv.glmnet_AUC = fit.logit.lasso$cvm
) %>%
  knitr::kable()
```

lambda	ours_AUC	cv.glmnet_AUC
0.4950042	0.5000000	0.5000000
0.3756674	0.9701369	0.9718839
0.2851006	0.9789711	0.9778577
0.2163679	0.9817904	0.9817045
0.1642054	0.9834819	0.9840974
0.1246184	0.9848048	0.9849193
0.0945751	0.9862041	0.9860165
0.0717747	0.9875328	0.9877492
0.0544711	0.9892431	0.9890586
0.0413391	0.9894404	0.9895565
0.0313729	0.9899379	0.9898538
0.0238095	0.9901358	0.9900528
0.0180694	0.9903346	0.9903509
0.0137132	0.9903339	0.9905500
0.0104072	0.9905325	0.9906491
0.0078982	0.9905317	0.9904501
0.0059941	0.9904323	0.9905479
0.0045490	0.9894383	0.9894559
0.0034523	0.9887423	0.9887615
0.0026200	0.9880455	0.9881670

lambda	ours_AUC	cv.glmnet_AUC
0.0019884	0.9878508	0.9877718
0.0015090	0.9875498	0.9874650
0.0011452	0.9864511	0.9865655
0.0008691	0.9862492	0.9865631
0.0006596	0.9857439	0.9856522
0.0005006	0.9848421	0.9848468
0.0003799	0.9844401	0.9840349
0.0002883	0.9837394	0.9839369
0.0002188	0.9837398	0.9837380
0.0001661	0.9841404	0.9842386

The best  $\lambda$ 's are the same, and the coefficients are very similar.

```
# our best lambda
best_lambda
```

```
## [1] 0.01040719
```

```
# cv.glmnet's best lambda
fit.logit.lasso$lambda.min
```

```
## [1] 0.01040719
```

```
tibble(
  predictor = c("(Intercept)", names(Training)[-1]),
  ours_coef = res_coef[res_coef$lambda == best_lambda, -1] %>% as.vector %>% as.numeric,
  cv.glmnet_coef = coef(fit.logit.lasso, fit.logit.lasso$lambda.min) %>% as.vector
) %>%
  knitr::kable()
```

predictor	ours_coef	cv.glmnet_coef
(Intercept)	-0.6314292	-0.6155278
radius_mean	0.0000000	0.0000000
texture_mean	0.3808524	0.3680397
perimeter_mean	0.0000000	0.0000000
area_mean	0.0000000	0.0000000
smoothness_mean	0.0000000	0.0000000
compactness_mean	0.0000000	0.0000000
concavity_mean	0.0000000	0.0000000
concave.points_mean	0.4863225	0.4396486
symmetry_mean	0.0000000	0.0000000
fractal_dimension_mean	0.0000000	0.0000000
radius_se	0.6822552	0.7840213
texture_se	0.0000000	0.0000000
perimeter_se	0.0000000	0.0000000
area_se	0.0000000	0.0000000
smoothness_se	0.0000000	0.0000000
compactness_se	0.0000000	0.0000000

predictor	ours_coef	cv.glmnet_coef
concavity_se	0.0000000	0.0000000
concave.points_se	0.0000000	0.0000000
symmetry_se	0.0000000	0.0000000
fractal_dimension_se	-0.0145589	-0.0167962
radius_worst	2.5340492	2.5469005
texture_worst	0.4297598	0.4561346
perimeter_worst	0.0000000	0.0000000
area_worst	0.0000000	0.0000000
smoothness_worst	0.4749630	0.4813888
compactness_worst	0.0000000	0.0000000
concavity_worst	0.1166963	0.1024629
concave.points_worst	1.0005941	1.0701080
symmetry_worst	0.3539412	0.3587025
fractal_dimension_worst	0.0000000	0.0000000

## Prediction performance comparison

Below is the prediction performance on the test data.

```
# test data
X_test <- cbind(rep(1, nrow(Test)), model.matrix(diagnosis ~ ., Test)[, -1])
y_test <- Test$diagnosis

# logistic model
res_logit <- NewtonRaphson(dat = Training, func = logisticstuff, start = rep(0, ncol(Training)))

## Warning in NewtonRaphson(dat = Training, func = logisticstuff, start = rep(0, :
## Complete separation occurs. Algorithm does not converge.

betavec_logit <- res_logit[nrow(res_logit), 3:ncol(res_logit)]
u <- X_test %*% betavec_logit
phat <- sigmoid(u)[, 1]
roc_logit <- roc(response = y_test, predictor = phat)

# logistic LASSO model
betavec_logit.lasso <- res_coef[res_coef$lambda == best_lambda, -c(1, 2)]
col_nonzero <- names(betavec_logit.lasso)[betavec_logit.lasso != 0]
df_nonzero <- Training[c("diagnosis", col_nonzero)]
refit_logit <- NewtonRaphson(dat = df_nonzero, func = logisticstuff, start = rep(0, ncol(df_nonzero)))
betavec_lasso.refit <- refit_logit[nrow(refit_logit), 3:ncol(refit_logit)]
betavec_lasso.refit <- bind_rows(betavec_logit.lasso, betavec_lasso.refit)[2,] %>% select("(Intercept)")
betavec_lasso.refit[is.na(betavec_lasso.refit)] <- 0
betavec_lasso.refit <- as.numeric(betavec_lasso.refit)
u <- X_test %*% betavec_lasso.refit
phat <- sigmoid(u)[, 1]
roc_logitlasso <- roc(response = y_test, predictor = phat)

# logistic LASSO model (cv.glmnet)
betavec_logit.lasso.glm_temp <- coef(fit_logit.lasso, fit_logit.lasso$lambda.min)
betavec_logit.lasso.glm <- betavec_logit.lasso.glm_temp %>% as.vector
```

```

names(betavec_logit.lasso.glm) <- betavec_logit.lasso.glm_temp@Dimnames[[1]]
col_nonzero <- names(betavec_logit.lasso.glm)[betavec_logit.lasso.glm != 0][-1]
df_nonzero <- Training[c("diagnosis", col_nonzero)]
refit_logit.glm <- NewtonRaphson(dat = df_nonzero, func = logisticstuff, start = rep(0, ncol(df_nonzero)))
betavec_lasso.refit.glm <- refit_logit.glm[nrow(refit_logit.glm), 3:ncol(refit_logit.glm)]
betavec_lasso.refit.glm <- bind_rows(betavec_logit.lasso, betavec_lasso.refit.glm)[2,] %>% select("(Int",
betavec_lasso.refit.glm[is.na(betavec_lasso.refit.glm)] <- 0
betavec_lasso.refit.glm <- as.numeric(betavec_lasso.refit.glm)
u <- X_test %*% betavec_lasso.refit.glm
phat <- sigmoid(u)[, 1]
roc.logitlasso.glm <- roc(response = y_test, predictor = phat)

# draw rocs
auc <- c(roc.logit$auc[1], roc.logitlasso$auc[1], roc.logitlasso.glm$auc[1])
auc

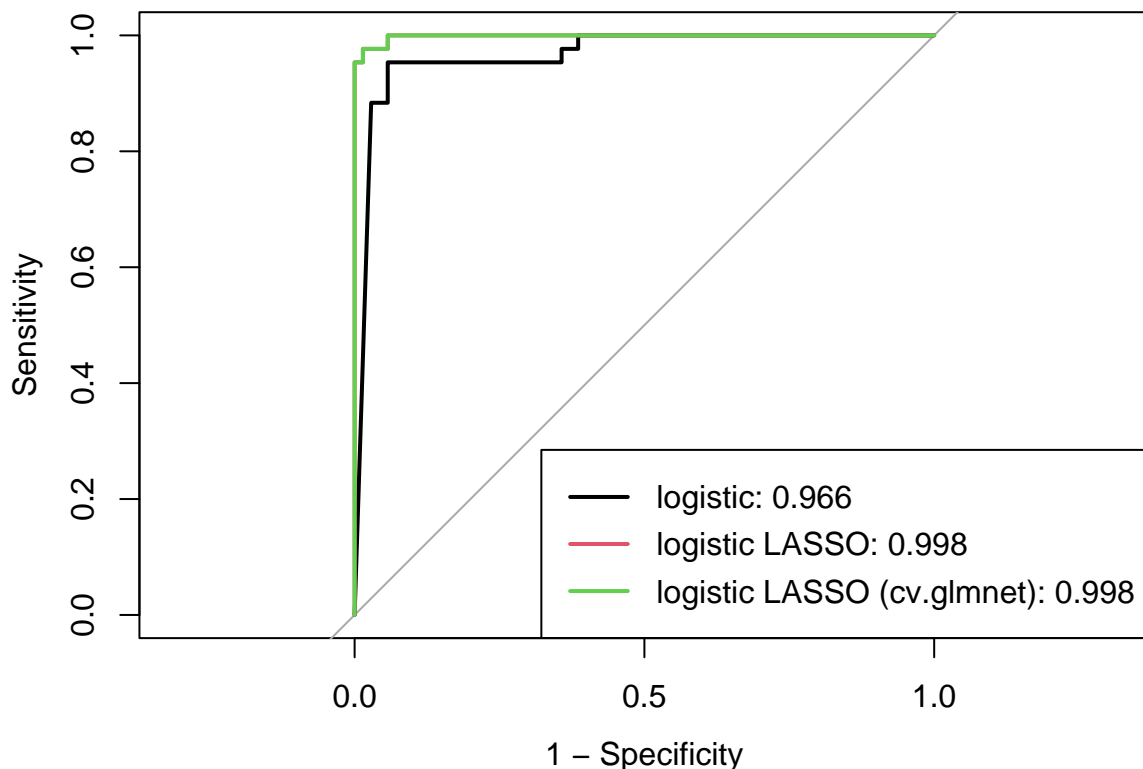
```

```
## [1] 0.9661130 0.9983389 0.9983389
```

```

plot(roc.logit, legacy.axes = TRUE)
plot(roc.logitlasso, col = 2, add = TRUE)
plot(roc.logitlasso.glm, col = 3, add = TRUE)
modelNames <- c("logistic", "logistic LASSO", "logistic LASSO (cv.glmnet)")
legend("bottomright", legend = paste0(modelNames, ": ", round(auc, 3)),
col = 1:3, lwd = 2)

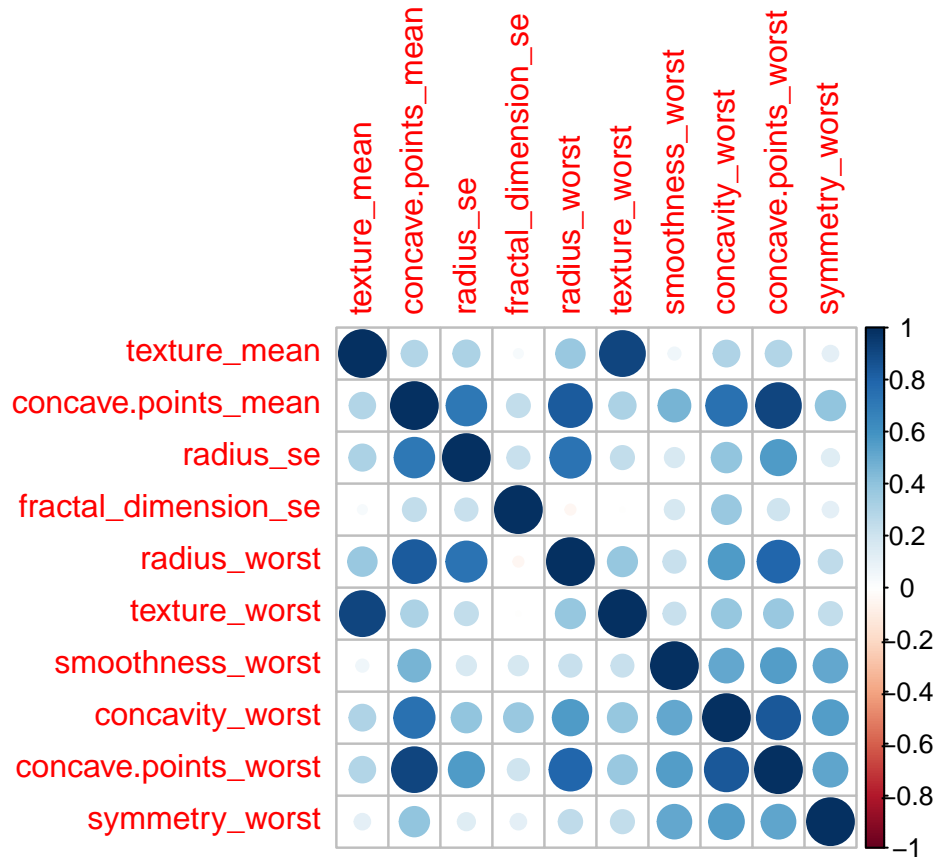
```





## Correlation Plot

```
corrplot::corrplot(cor(Training[names(betavec_logit.lasso)[betavec_logit.lasso != 0]]))
```



## LASSO model coefficients

Re-fit the logistic regression with the predictors selected by LASSO.

```
refit_logit[nrow(refit_logit), 3:ncol(refit_logit)]
```

```
##      (Intercept)      texture_mean  concave.points_mean
##      -0.09092737      0.94392378      0.84754704
##      radius_se fractal_dimension_se      radius_worst
##      3.77040802      -1.12476354      6.01510669
##      texture_worst  smoothness_worst  concavity_worst
##      1.23805580      1.67539455      1.19425400
## concave.points_worst  symmetry_worst
##      1.87249528      0.69899461
```