

Risk-Evaluation: Breast Cancer Royston-Altman

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1 Evaluation of RISK survival models

This document highlights the use of

- `RRPlot()`,
- `CoxRiskCalibration()`, and
- `CalibrationProbPoissonRisk()`,

for the evaluation (`RRPlot`), and calibration of cox models (`CoxRiskCalibration`) or logistic models (`CalibrationProbPoissonRisk`) of survival data.

Furthermore, it can be used to evaluate any Risk index that reruns the probability of a future event on external data-set.

This document will use the `survival::rotterdam`, and `survival::gbsg` data-sets to train and predict the risk of cancer recurrence after surgery. Both Cox and Logistic models will be trained and evaluated.

Here are some sample plots returned by the evaluated functions:

1.1 The libraries

```
library(survival)
library(FRESA.CAD)
```

```
## Loading required package: Rcpp
## Loading required package: stringr
## Loading required package: miscTools
## Loading required package: Hmisc
```

```
##
## Attaching package: 'Hmisc'

## The following objects are masked from 'package:base':
##
##      format.pval, units

## Loading required package: pROC

## Type 'citation("pROC")' for a citation.

##
## Attaching package: 'pROC'

## The following objects are masked from 'package:stats':
##
##      cov, smooth, var

source("~/GitHub/FRESA.CAD/R/RRPlot.R")
source("~/GitHub/FRESA.CAD/R/PoissonEventRiskCalibration.R")
op <- par(no.readonly = TRUE)
pander::panderOptions('digits', 3)
#pander::panderOptions('table.split.table', 400)
pander::panderOptions('keep.trailing.zeros', TRUE)
```

1.2 Breast Cancer Royston-Altman data

1.2.1 data(gbsg, package="survival") and data(rotterdam, package="survival")

```
gbsgdata <- gbsg
rownames(gbsgdata) <- gbsgdata$pid
gbsgdata$pid <- NULL

odata <- rotterdam
rownames(odata) <- odata$pid
odata$pid <- NULL
odata$rfstime <- odata$rtime
odata$status <- odata$recur
odata$rtime <- NULL
odata$recur <- NULL

odata <- odata[,colnames(odata) %in% colnames(gbsgdata)]

odata$size <- 10*(odata$size=="<=20") +
  35*(odata$size=="20-50") +
  60*(odata$size==">50")

data <- as.data.frame(model.matrix(Surv(rfstime,status)~.*.,odata))

data$`(Intercept)` <- NULL

dataBrestCancerTrain <- cbind(time=odata[rownames(data),"rfstime"],status=odata[rownames(data),"status"])

colnames(dataBrestCancerTrain) <- str_replace_all(colnames(dataBrestCancerTrain),":","_")
colnames(dataBrestCancerTrain) <- str_replace_all(colnames(dataBrestCancerTrain)," ","")
colnames(dataBrestCancerTrain) <- str_replace_all(colnames(dataBrestCancerTrain),"\\.", "_")
colnames(dataBrestCancerTrain) <- str_replace_all(colnames(dataBrestCancerTrain),"-", "_")
```

```
colnames(dataBrestCancerTrain) <-str_replace_all(colnames(dataBrestCancerTrain), ">", "_")
dataBrestCancerTrain$time <- dataBrestCancerTrain$time/365 ## To years

pander::pander(table(odata[rownames(data), "status"]), caption="rotterdam")
```

Table 1: rotterdam

| 0 | 1 |
|------|------|
| 1464 | 1518 |

1.2.2 data(gbsg, package="survival") data conditioning

```
gbsgdata <- gbsgdata[, colnames(odata)]
data <- as.data.frame(model.matrix(Surv(rfstime, status) ~ .*, gbsgdata))

data$`(Intercept)` <- NULL

dataBrestCancerTest <- cbind(time=gbsgdata[rownames(data), "rfstime"], status=gbsgdata[rownames(data), "status"])

colnames(dataBrestCancerTest) <-str_replace_all(colnames(dataBrestCancerTest), ":", "_")
colnames(dataBrestCancerTest) <-str_replace_all(colnames(dataBrestCancerTest), " ", "_")
colnames(dataBrestCancerTest) <-str_replace_all(colnames(dataBrestCancerTest), "\\.", "_")
colnames(dataBrestCancerTest) <-str_replace_all(colnames(dataBrestCancerTest), "-", "_")
colnames(dataBrestCancerTest) <-str_replace_all(colnames(dataBrestCancerTest), ">", "_")
dataBrestCancerTest$time <- dataBrestCancerTest$time/365

pander::pander(table(odata[rownames(data), "status"]), caption="gbsg")
```

Table 2: gbsg

| 0 | 1 |
|-----|-----|
| 499 | 183 |

1.3 Cox Modeling

```
ml <- BSWiMS.model(Surv(time, status) ~ ., data=dataBrestCancerTrain, loops=1, NumberofRepeats = 5)

sm <- summary(ml)
pander::pander(sm$coefficients)
```

Table 3: Table continues below

| | Estimate | lower | HR | upper | u.Accuracy | r.Accuracy |
|------------|-----------|-------|-------|-------|------------|------------|
| age_nodes | 7.76e-04 | 1.001 | 1.001 | 1.001 | 0.626 | 0.601 |
| size_grade | 5.02e-03 | 1.005 | 1.005 | 1.006 | 0.598 | 0.624 |
| nodes | 8.25e-02 | 1.077 | 1.086 | 1.095 | 0.637 | 0.642 |
| size | 6.77e-03 | 1.005 | 1.007 | 1.009 | 0.595 | 0.641 |
| size_nodes | -3.61e-04 | 1.000 | 1.000 | 1.000 | 0.624 | 0.643 |

| | Estimate | lower | HR | upper | u.Accuracy | r.Accuracy |
|-------------|-----------|-------|-------|-------|------------|------------|
| age_pgr | -4.02e-06 | 1.000 | 1.000 | 1.000 | 0.548 | 0.631 |
| grade | 1.97e-01 | 1.138 | 1.218 | 1.303 | 0.565 | 0.638 |
| age_size | -1.23e-04 | 1.000 | 1.000 | 1.000 | 0.567 | 0.629 |
| age | -2.97e-03 | 0.996 | 0.997 | 0.998 | 0.513 | 0.628 |
| grade_nodes | -1.31e-02 | 0.982 | 0.987 | 0.992 | 0.635 | 0.645 |
| grade_pgr | 4.96e-05 | 1.000 | 1.000 | 1.000 | 0.541 | 0.633 |
| size_pgr | 1.22e-06 | 1.000 | 1.000 | 1.000 | 0.490 | 0.633 |
| meno_nodes | -2.80e-03 | 0.996 | 0.997 | 0.999 | 0.580 | 0.635 |
| meno_pgr | 7.13e-05 | 1.000 | 1.000 | 1.000 | 0.527 | 0.634 |
| age_grade | -8.79e-05 | 1.000 | 1.000 | 1.000 | 0.508 | 0.632 |

Table 4: Table continues below

| | full.Accuracy | u.AUC | r.AUC | full.AUC | IDI |
|-------------|---------------|-------|-------|----------|----------|
| age_nodes | 0.633 | 0.630 | 0.602 | 0.634 | 0.028330 |
| size_grade | 0.633 | 0.599 | 0.627 | 0.635 | 0.017876 |
| nodes | 0.643 | 0.640 | 0.644 | 0.644 | 0.007344 |
| size | 0.643 | 0.595 | 0.642 | 0.644 | 0.013742 |
| size_nodes | 0.643 | 0.629 | 0.644 | 0.644 | 0.003445 |
| age_pgr | 0.635 | 0.544 | 0.634 | 0.637 | 0.009210 |
| grade | 0.643 | 0.561 | 0.639 | 0.644 | 0.009051 |
| age_size | 0.633 | 0.568 | 0.632 | 0.634 | 0.005990 |
| age | 0.643 | 0.513 | 0.628 | 0.644 | 0.004165 |
| grade_nodes | 0.643 | 0.639 | 0.646 | 0.644 | 0.002028 |
| grade_pgr | 0.635 | 0.537 | 0.636 | 0.637 | 0.004752 |
| size_pgr | 0.635 | 0.494 | 0.635 | 0.637 | 0.002102 |
| meno_nodes | 0.635 | 0.584 | 0.637 | 0.637 | 0.001735 |
| meno_pgr | 0.635 | 0.522 | 0.636 | 0.637 | 0.002519 |
| age_grade | 0.635 | 0.509 | 0.634 | 0.637 | 0.000891 |

| | NRI | z.IDI | z.NRI | Delta.AUC | Frequency |
|-------------|----------|-------|--------|-----------|-----------|
| age_nodes | 0.42972 | 12.26 | 13.415 | 3.21e-02 | 1.0 |
| size_grade | 0.37163 | 9.49 | 10.720 | 7.67e-03 | 1.0 |
| nodes | 0.06897 | 8.25 | 2.000 | 6.04e-05 | 1.0 |
| size | 0.33845 | 7.79 | 9.406 | 1.39e-03 | 1.0 |
| size_nodes | 0.34249 | 7.24 | 9.559 | -3.56e-04 | 1.0 |
| age_pgr | 0.18985 | 6.33 | 5.482 | 3.16e-03 | 0.2 |
| grade | 0.20188 | 5.83 | 6.161 | 5.02e-03 | 1.0 |
| age_size | 0.18848 | 5.63 | 5.214 | 2.53e-03 | 1.0 |
| age | 0.09238 | 5.27 | 2.525 | 1.55e-02 | 1.0 |
| grade_nodes | -0.09046 | 4.95 | -2.532 | -2.68e-03 | 1.0 |
| grade_pgr | 0.26528 | 4.83 | 7.348 | 1.01e-03 | 0.2 |
| size_pgr | 0.01627 | 3.82 | 0.452 | 1.62e-03 | 0.2 |
| meno_nodes | 0.00739 | 3.49 | 0.205 | -5.37e-04 | 0.2 |
| meno_pgr | 0.06788 | 3.30 | 1.862 | 4.64e-04 | 0.2 |
| age_grade | 0.07139 | 2.42 | 1.951 | 2.66e-03 | 0.2 |

1.4 Cox Model Performance

Here we evaluate the model using the `RRPlot()` function.

1.4.1 The evaluation of the raw Cox model with RRPlot()

Here we will use the predicted event probability assuming a baseline hazard for events withing 5 years

```
timeinterval <- 5 # Five years
```

```
h0 <- sum(dataBrestCancerTrain$status & dataBrestCancerTrain$time <= timeinterval)
```

```
h0 <- h0/sum((dataBrestCancerTrain$time > timeinterval) | (dataBrestCancerTrain$status==1))
```

```
pander::pander(t(c(h0=h0,timeinterval=timeinterval)),caption="Initial Parameters")
```

Table 6: Initial Parameters

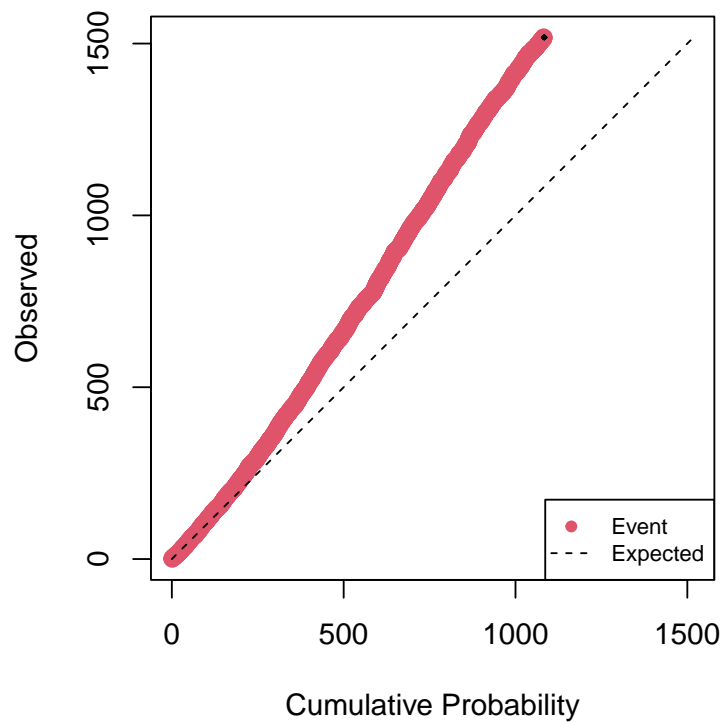
| h0 | timeinterval |
|-------|--------------|
| 0.429 | 5 |

```
index <- predict(ml,dataBrestCancerTrain)
```

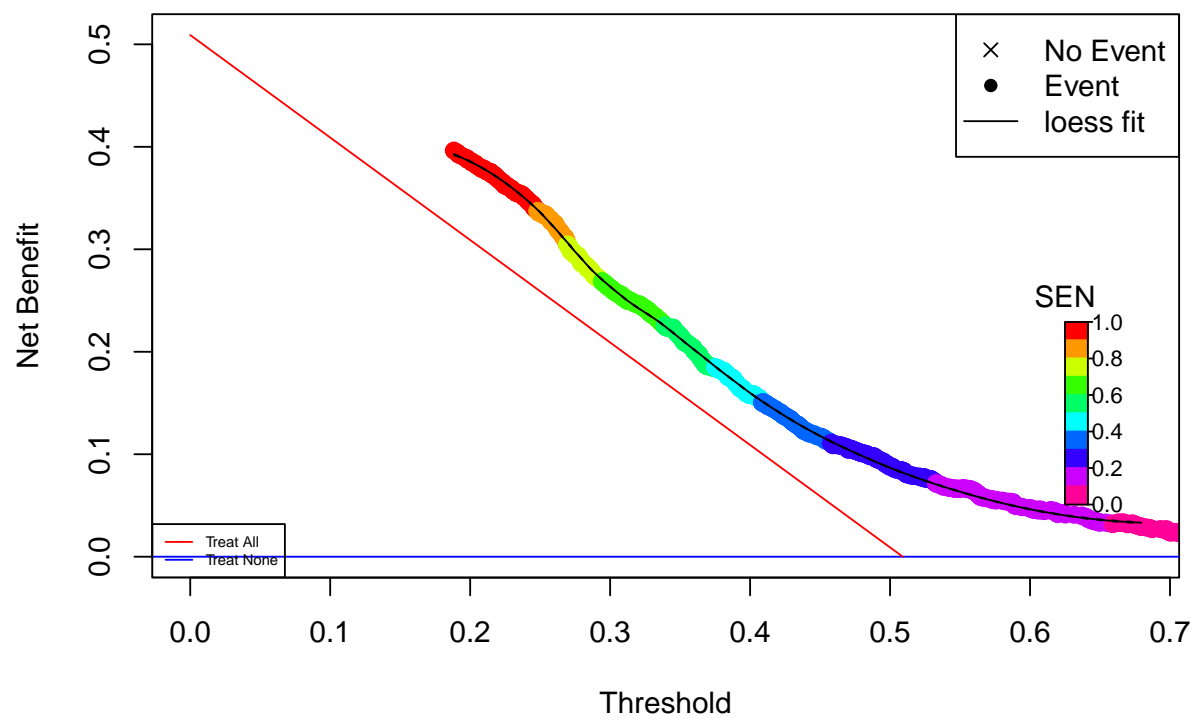
```
rdata <- cbind(dataBrestCancerTrain$status,ppoisGzero(index,h0))
```

```
rrAnalysisTrain <- RRPlot(rdata,atProb=c(0.90,0.80),  
  timetoEvent=dataBrestCancerTrain$time,  
  title="Train: Breast Cancer",  
  ysurvlim=c(0.00,1.0),  
  riskTimeInterval=timeinterval)
```

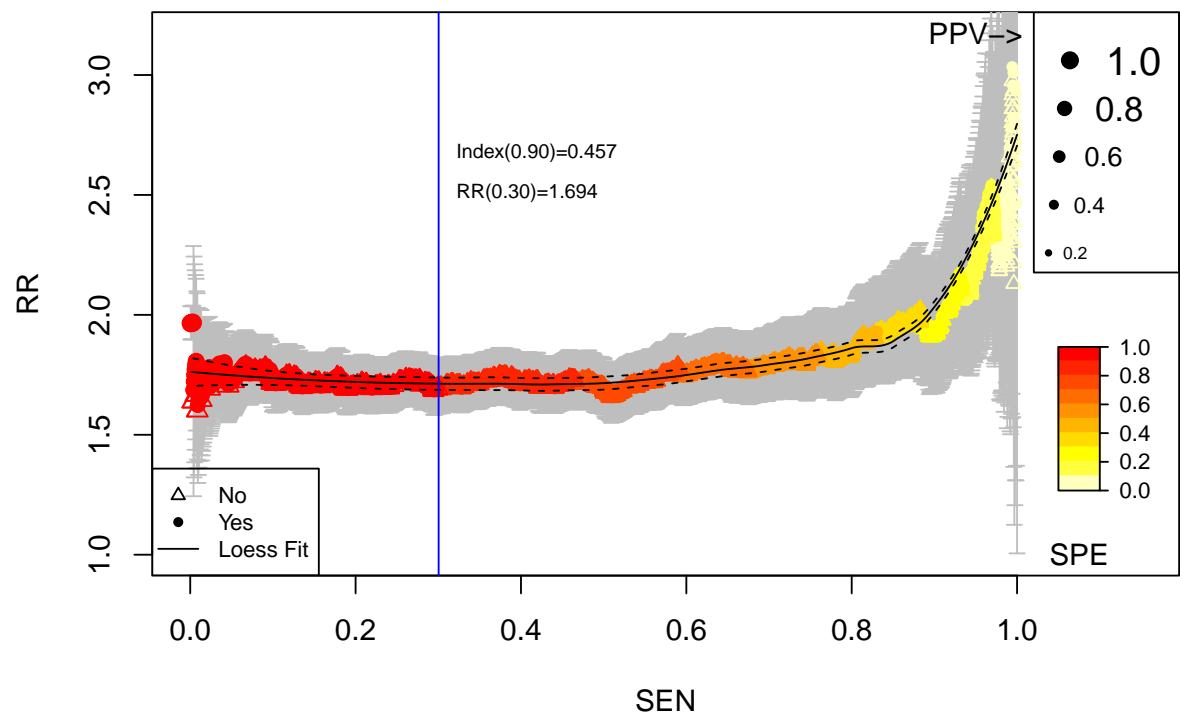
Cumulative vs. Observed: Train: Breast Cancer

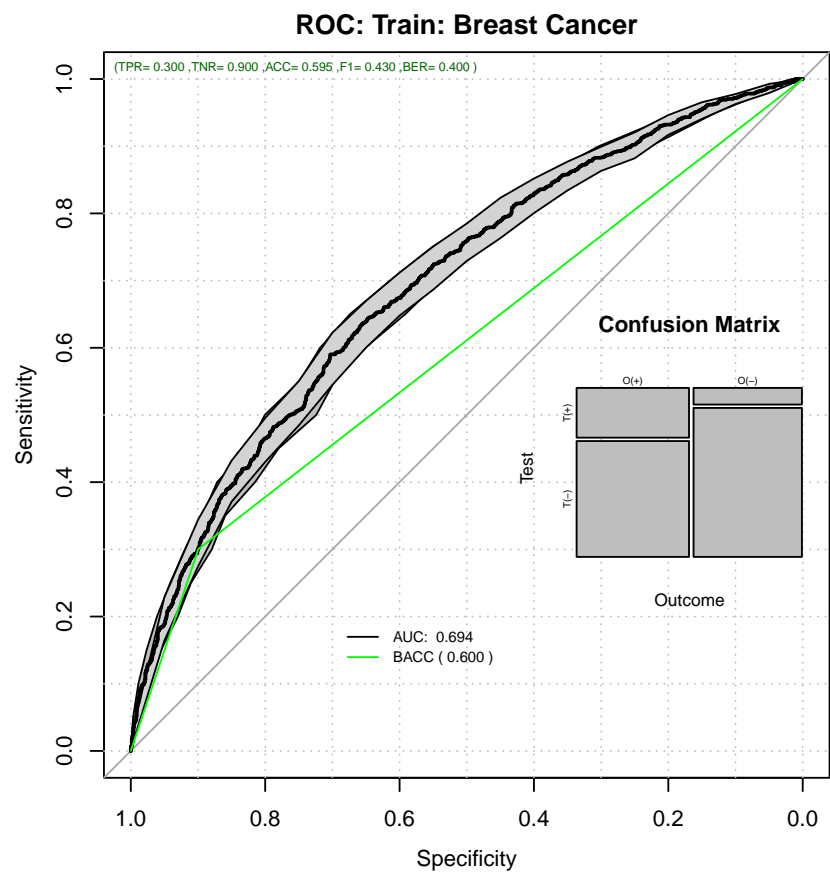


Decision Curve Analysis: Train: Breast Cancer

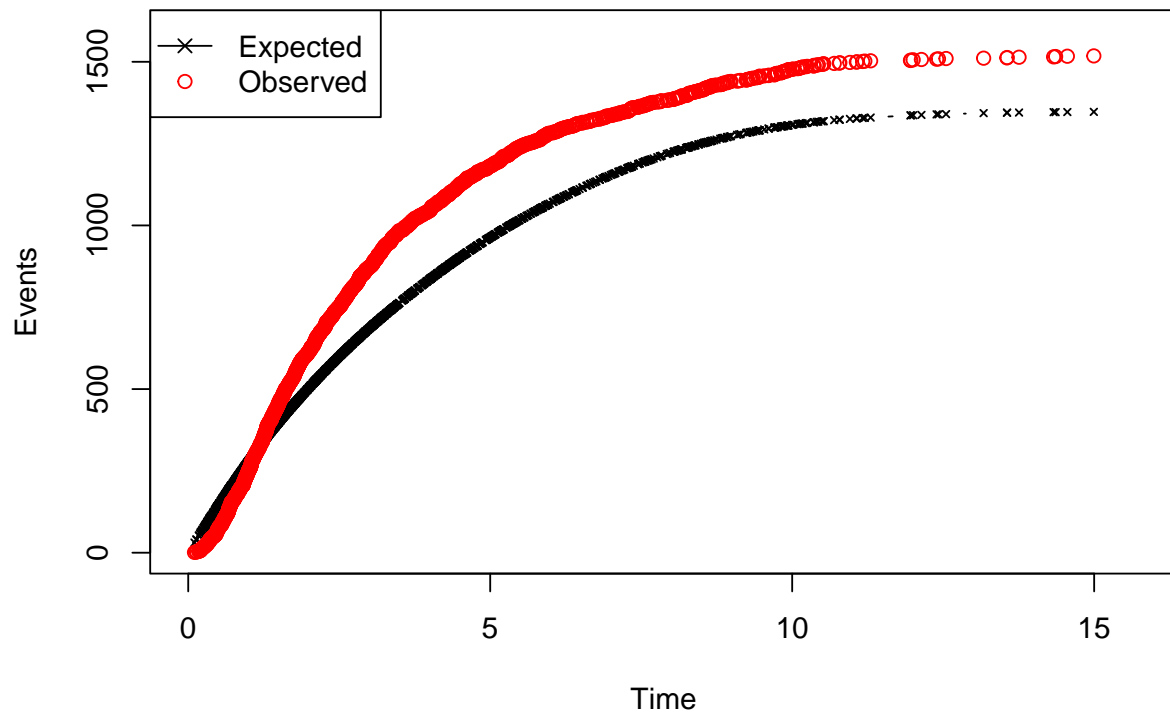


Relative Risk: Train: Breast Cancer

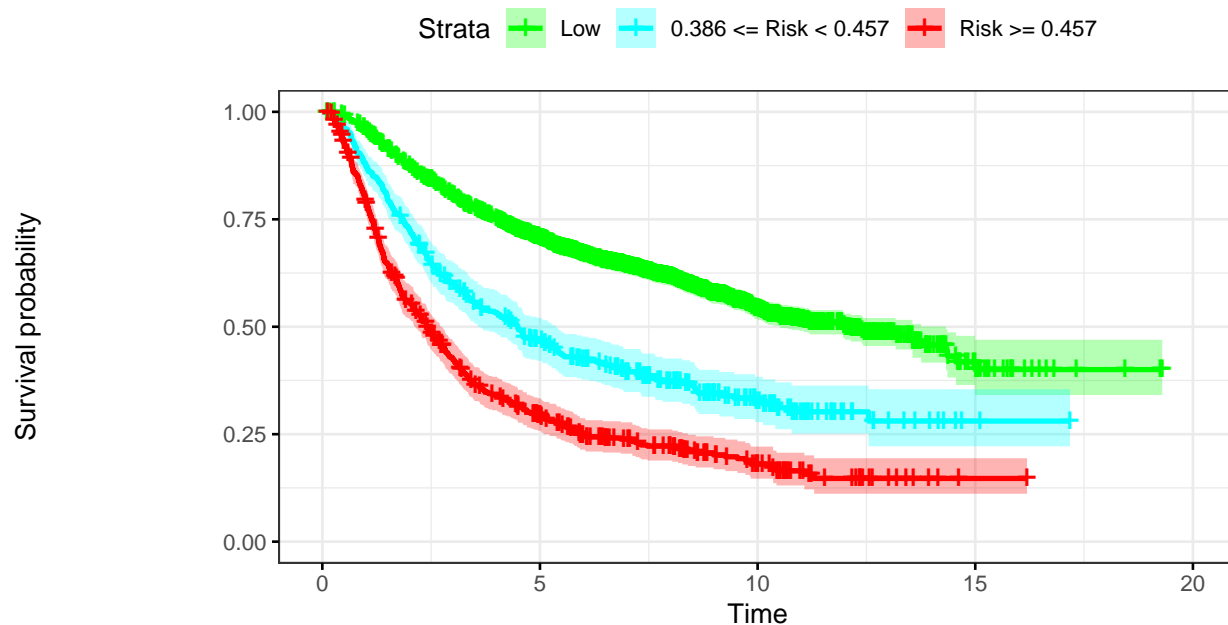




Time vs. Events: Train: Breast Cancer



Kaplan–Meier: Train: Breast Cancer



Number at risk

| | | | | | |
|----------------------------------|------|------|-----|----|---|
| Low | 1983 | 1263 | 396 | 23 | 0 |
| $0.386 \leq \text{Risk} < 0.457$ | 396 | 165 | 49 | 2 | 0 |
| $\text{Risk} \geq 0.457$ | 603 | 143 | 38 | 1 | 0 |

As we can see the Observed probability as well as the Time vs. Events are not calibrated.

1.4.2 Uncalibrated Performance Report

```
pander::pander(t(rrAnalysisTrain$OERatio),caption="O/E Ratio")
```

Table 7: O/E Ratio

| est | lower | upper |
|------|-------|-------|
| 1.13 | 1.07 | 1.19 |

```
pander::pander(t(rrAnalysisTrain$OE95ci),caption="O/E Ratio")
```

Table 8: O/E Ratio

| mean | 50% | 2.5% | 97.5% |
|------|------|------|-------|
| 1.13 | 1.13 | 1.12 | 1.14 |

```
pander::pander(t(rrAnalysisTrain$OAcum95ci),caption="O/Acum Ratio")
```

Table 9: O/Acum Ratio

| mean | 50% | 2.5% | 97.5% |
|------|------|------|-------|
| 1.34 | 1.34 | 1.34 | 1.35 |

```
pander::pander(rrAnalysisTrain$c.index$cstatCI,caption="C. Index")
```

| mean.C Index | median | lower | upper |
|--------------|--------|-------|-------|
| 0.677 | 0.677 | 0.663 | 0.69 |

```
pander::pander(t(rrAnalysisTrain$ROCAalysis$aucs),caption="ROC AUC")
```

Table 11: ROC AUC

| est | lower | upper |
|-------|-------|-------|
| 0.694 | 0.676 | 0.713 |

```
pander::pander((rrAnalysisTrain$ROCAalysis$sensitivity),caption="Sensitivity")
```

Table 12: Sensitivity

| est | lower | upper |
|-----|-------|-------|
| 0.3 | 0.277 | 0.324 |

```
pander::pander((rrAnalysisTrain$ROCAalysis$specificity),caption="Specificity")
```

Table 13: Specificity

| est | lower | upper |
|-----|-------|-------|
| 0.9 | 0.883 | 0.915 |

```
pander::pander(t(rrAnalysisTrain$thr_atP),caption="Probability Thresholds")
```

Table 14: Probability Thresholds

| 90% | 80% |
|-------|-------|
| 0.457 | 0.386 |

```
pander::pander(t(rrAnalysisTrain$RR_atP),caption="Risk Ratio")
```

Table 15: Risk Ratio

| est | lower | upper |
|------|-------|-------|
| 1.69 | 1.59 | 1.81 |

```
pander::pander(rrAnalysisTrain$surdif, caption="Logrank test")
```

Table 16: Logrank test Chisq = 479.123919 on 2 degrees of freedom,
p = 0.000000

| | N | Observed | Expected | (O-E)^2/E | (O-E)^2/V |
|----------------|------|----------|----------|-----------|-----------|
| class=0 | 1983 | 812 | 1145 | 96.7 | 398.2 |
| class=1 | 396 | 250 | 177 | 29.6 | 33.6 |
| class=2 | 603 | 456 | 196 | 345.4 | 401.8 |

1.4.3 Cox Calibration

```
op <- par(no.readonly = TRUE)

calprob <- CoxRiskCalibration(ml, dataBrestCancerTrain, "status", "time")

pander::pander(c(h0=calprob$h0,
  Gain=calprob$hazardGain,
  DeltaTime=calprob$timeInterval),
  caption="Cox Calibration Parameters")
```

| h0 | Gain | DeltaTime |
|-------|------|-----------|
| 0.698 | 1.35 | 6.96 |

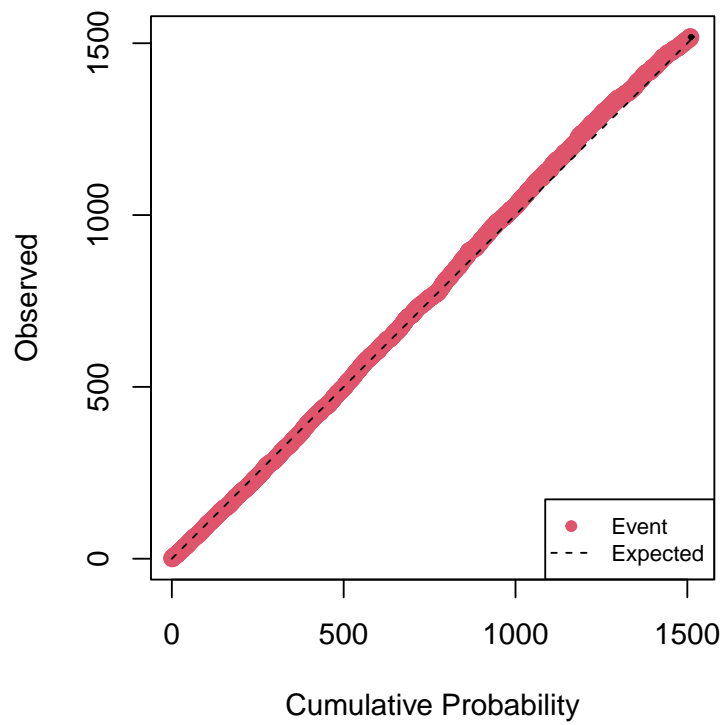
1.4.4 The RRplot() of the calibrated model

```
h0 <- calprob$h0
timeinterval <- calprob$timeInterval;

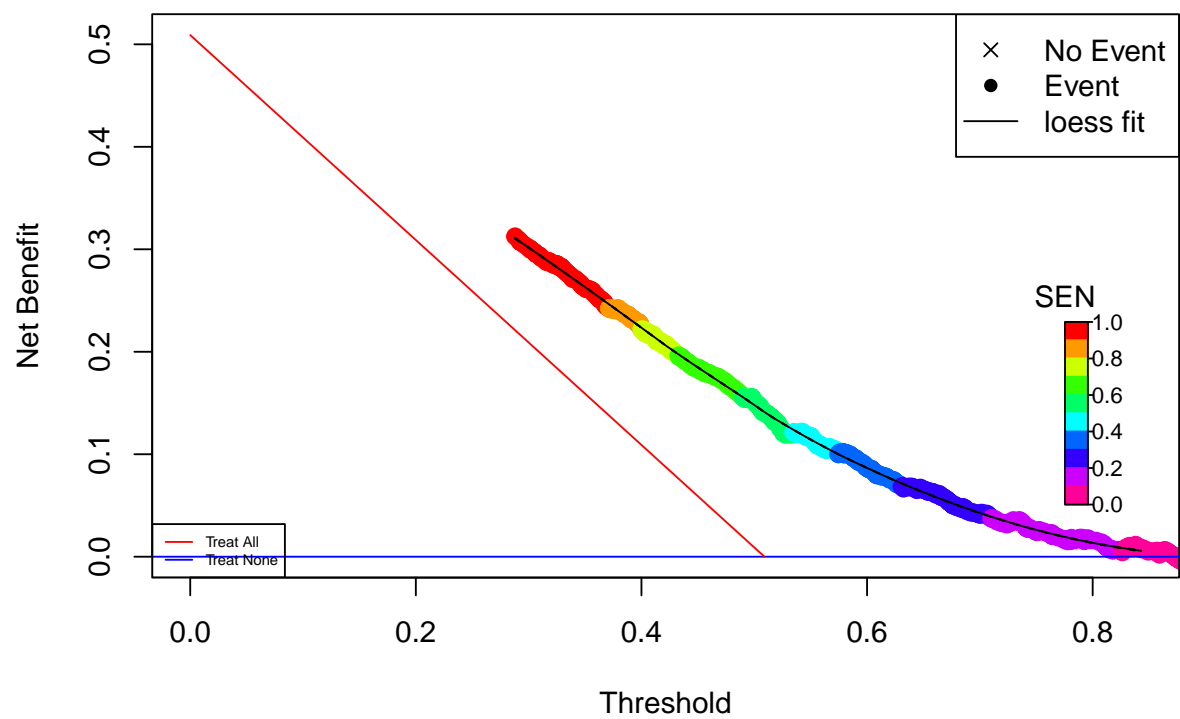
rdata <- cbind(dataBrestCancerTrain$status, calprob$prob)

rrAnalysisTrain <- RRPlot(rdata, atProb=c(0.90, 0.80),
  timetoEvent=dataBrestCancerTrain$time,
  title="Cal. Train: Breast Cancer",
  ysurvlim=c(0.00, 1.0),
  riskTimeInterval=timeinterval)
```

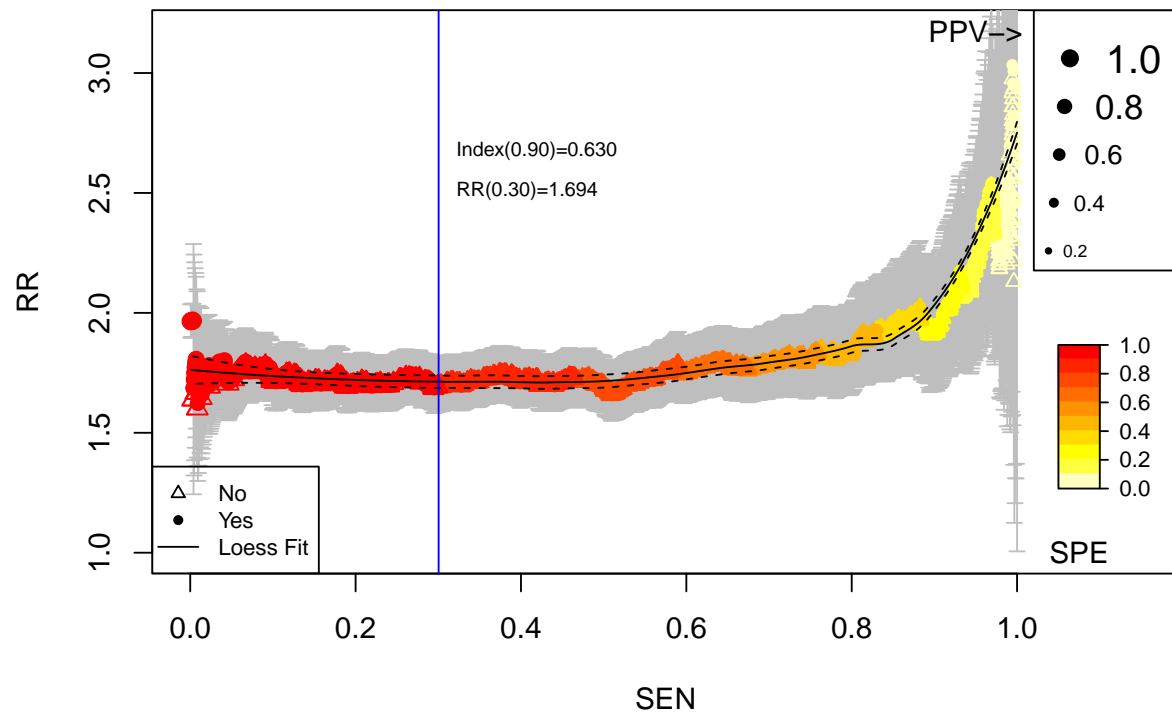
Cumulative vs. Observed: Cal. Train: Breast Cancer

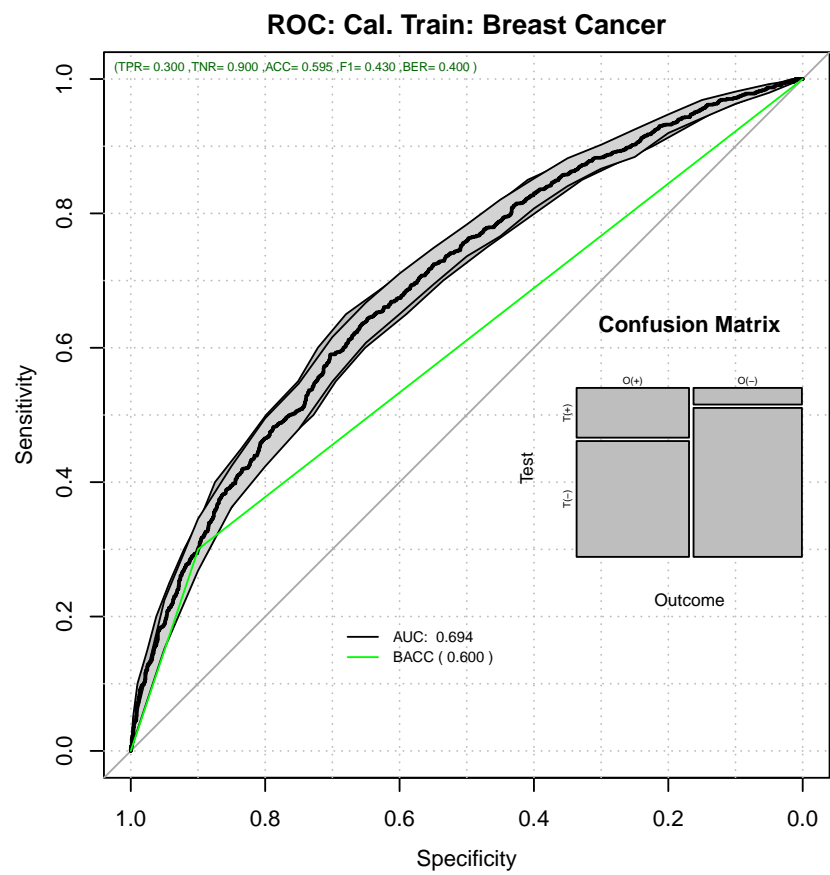


Decision Curve Analysis: Cal. Train: Breast Cancer

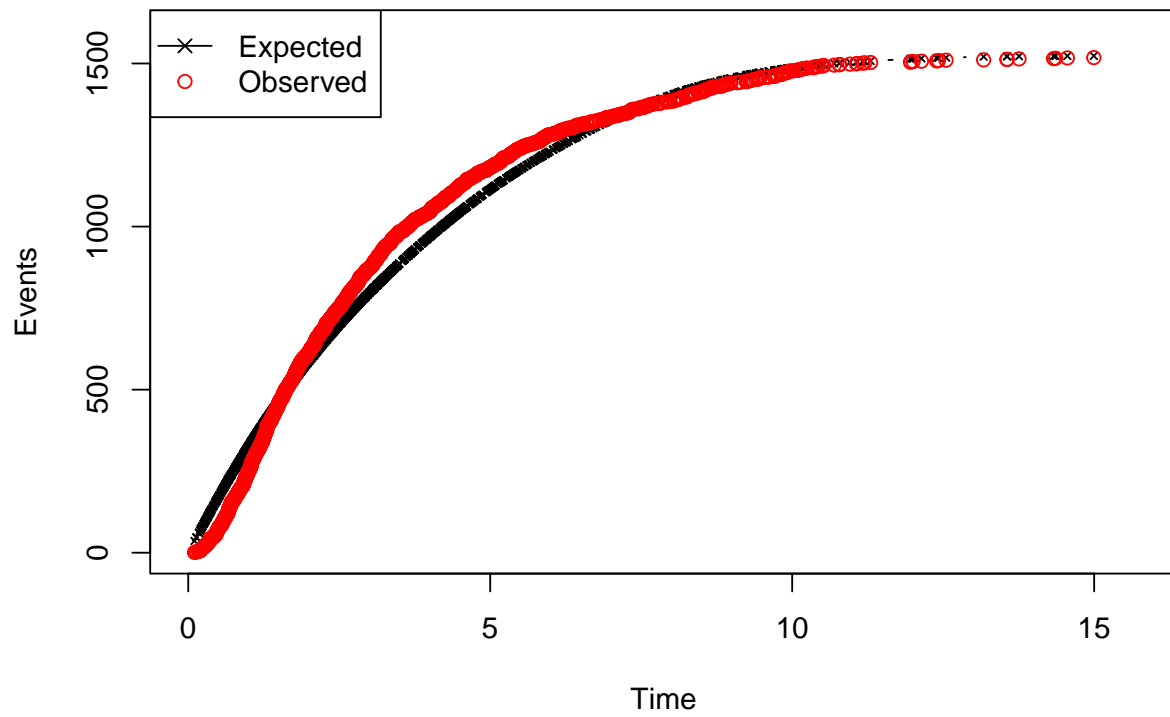


Relative Risk: Cal. Train: Breast Cancer

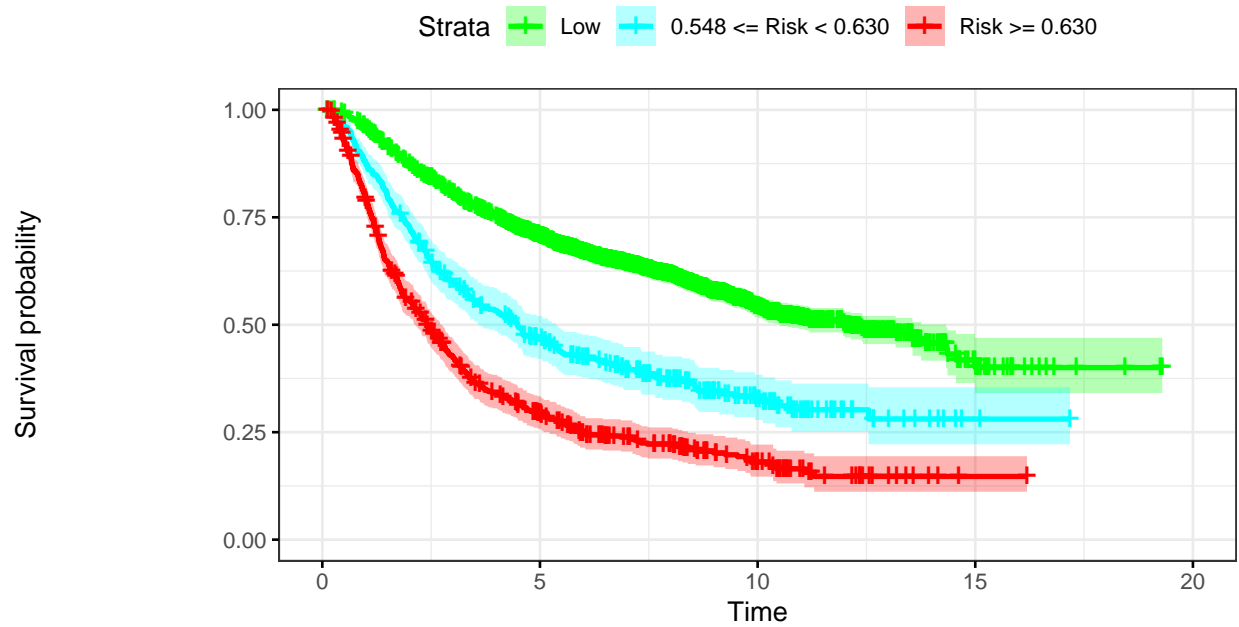




Time vs. Events: Cal. Train: Breast Cancer



Kaplan–Meier: Cal. Train: Breast Cancer



Number at risk

| | | | | | |
|----------------------------------|------|------|-----|----|---|
| Low | 1983 | 1263 | 396 | 23 | 0 |
| $0.548 \leq \text{Risk} < 0.630$ | 396 | 165 | 49 | 2 | 0 |
| $\text{Risk} \geq 0.630$ | 603 | 143 | 38 | 1 | 0 |

1.4.5 Calibrated Train Performance

```
pander::pander(t(rrAnalysisTrain$OERatio),caption="O/E Ratio")
```

Table 18: O/E Ratio

| est | lower | upper |
|-------|-------|-------|
| 0.997 | 0.947 | 1.05 |

```
pander::pander(t(rrAnalysisTrain$OE95ci),caption="O/E Ratio")
```

Table 19: O/E Ratio

| mean | 50% | 2.5% | 97.5% |
|-------|-------|-------|-------|
| 0.977 | 0.977 | 0.969 | 0.984 |

```
pander::pander(t(rrAnalysisTrain$OAcum95ci),caption="O/Acum Ratio")
```

Table 20: O/Acum Ratio

| mean | 50% | 2.5% | 97.5% |
|------|------|------|-------|
| 1.01 | 1.01 | 1.01 | 1.01 |

```
pander::pander(rrAnalysisTrain$c.index$cstatCI,caption="C. Index")
```

| mean.C Index | median | lower | upper |
|--------------|--------|-------|-------|
| 0.677 | 0.677 | 0.663 | 0.69 |

```
pander::pander(t(rrAnalysisTrain$ROCAalysis$aucs),caption="ROC AUC")
```

Table 22: ROC AUC

| est | lower | upper |
|-------|-------|-------|
| 0.694 | 0.676 | 0.713 |

```
pander::pander((rrAnalysisTrain$ROCAalysis$sensitivity),caption="Sensitivity")
```

Table 23: Sensitivity

| est | lower | upper |
|-----|-------|-------|
| 0.3 | 0.277 | 0.324 |

```
pander::pander((rrAnalysisTrain$ROCAalysis$specificity),caption="Specificity")
```

Table 24: Specificity

| est | lower | upper |
|-----|-------|-------|
| 0.9 | 0.883 | 0.915 |

```
pander::pander(t(rrAnalysisTrain$thr_atP),caption="Probability Thresholds")
```

Table 25: Probability Thresholds

| 90% | 80% |
|------|-------|
| 0.63 | 0.548 |

```
pander::pander(t(rrAnalysisTrain$RR_atP),caption="Risk Ratio")
```

Table 26: Risk Ratio

| est | lower | upper |
|------|-------|-------|
| 1.69 | 1.59 | 1.81 |

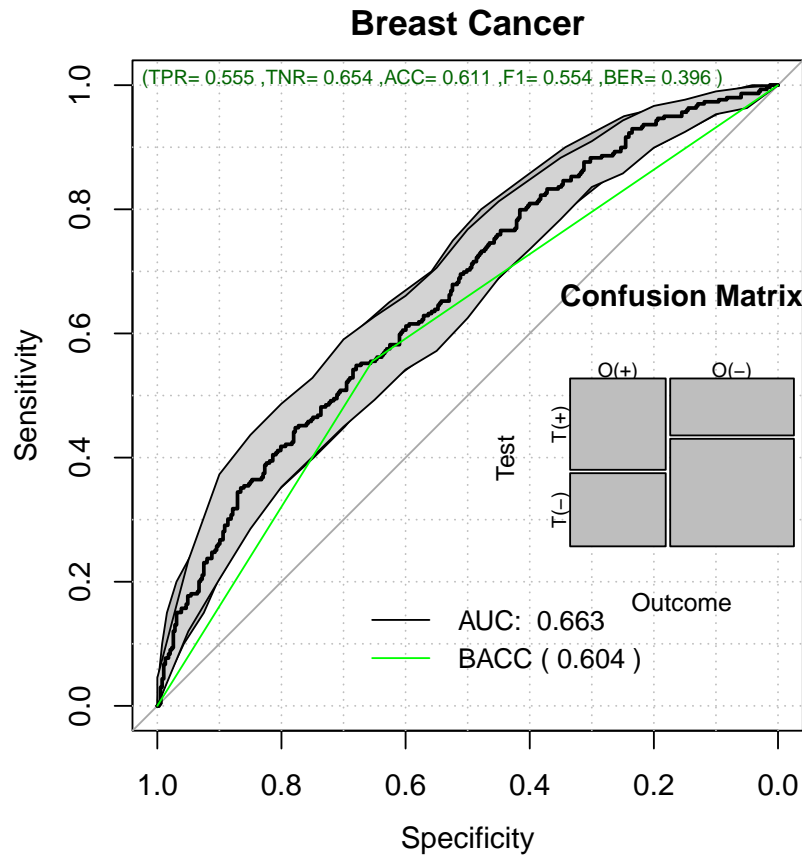
```
pander::pander(rrAnalysisTrain$surdif,caption="Logrank test")
```

Table 27: Logrank test Chisq = 479.123919 on 2 degrees of freedom,
p = 0.000000

| | N | Observed | Expected | (O-E)^2/E | (O-E)^2/V |
|---------|------|----------|----------|-----------|-----------|
| class=0 | 1983 | 812 | 1145 | 96.7 | 398.2 |
| class=1 | 396 | 250 | 177 | 29.6 | 33.6 |
| class=2 | 603 | 456 | 196 | 345.4 | 401.8 |

1.5 Performance on the external data set

```
index <- predict(ml,dataBreStCancerTest)
pp <- predictionStats_binary(cbind(dataBreStCancerTest$status,index),plotname="Breast Cancer")
```



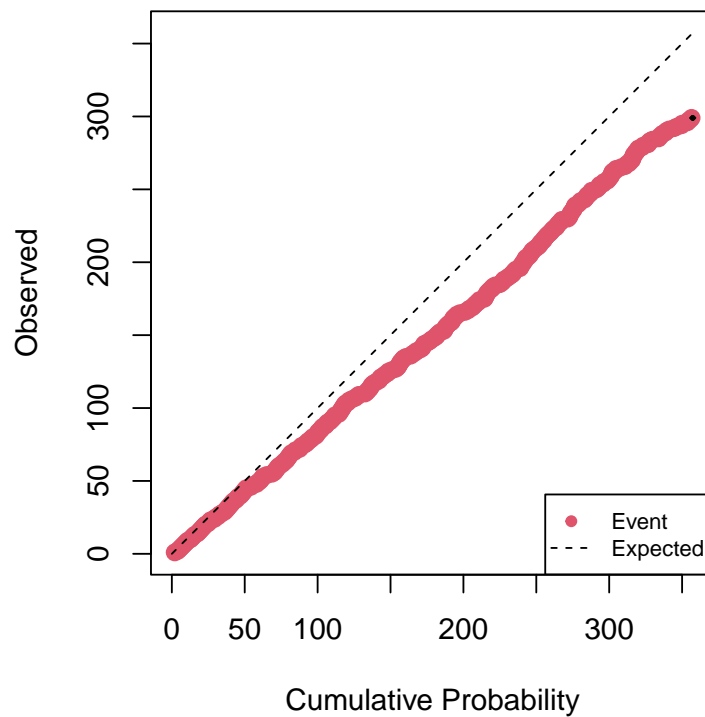
```
par(op)
```

```

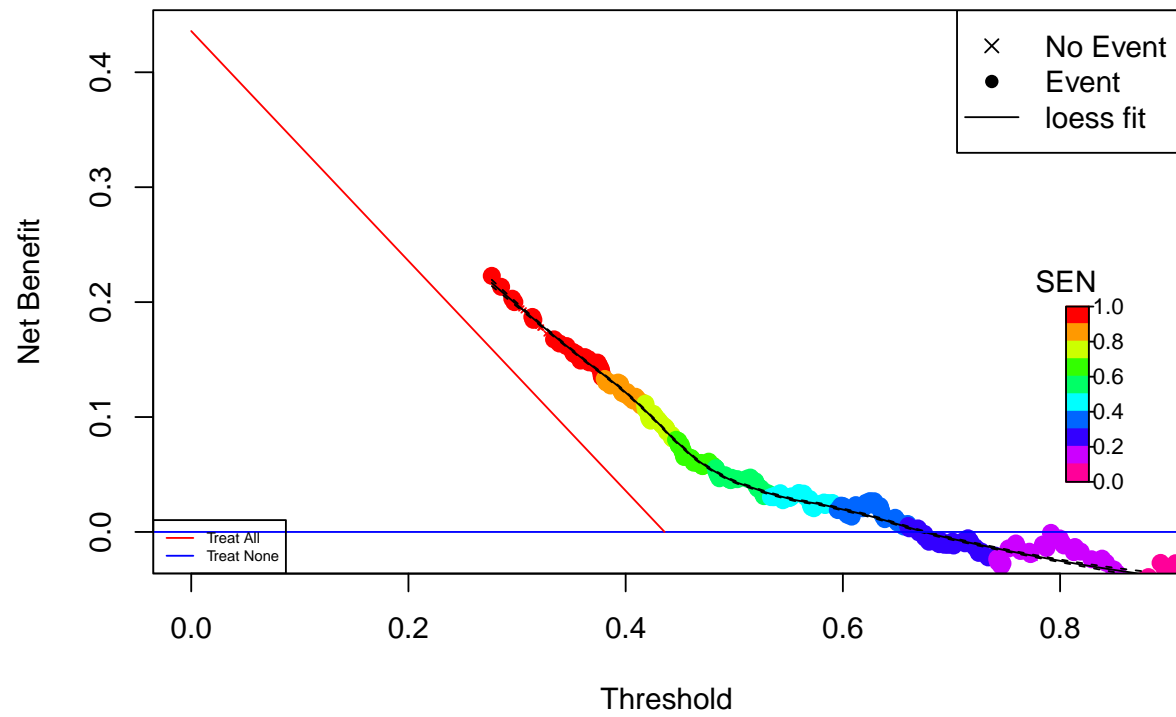
prob <- ppoisGzero(index,h0)
rdata <- cbind(dataBreastCancerTest$status,prob)
rrCoxTestAnalysis <- RRPlot(rdata,atThr=rrAnalysisTrain$thr_atP,
                             timetoEvent=dataBreastCancerTest$time,
                             title="Test: Breast Cancer",
                             ysurvlim=c(0.00,1.0),
                             riskTimeInterval=timeinterval)

```

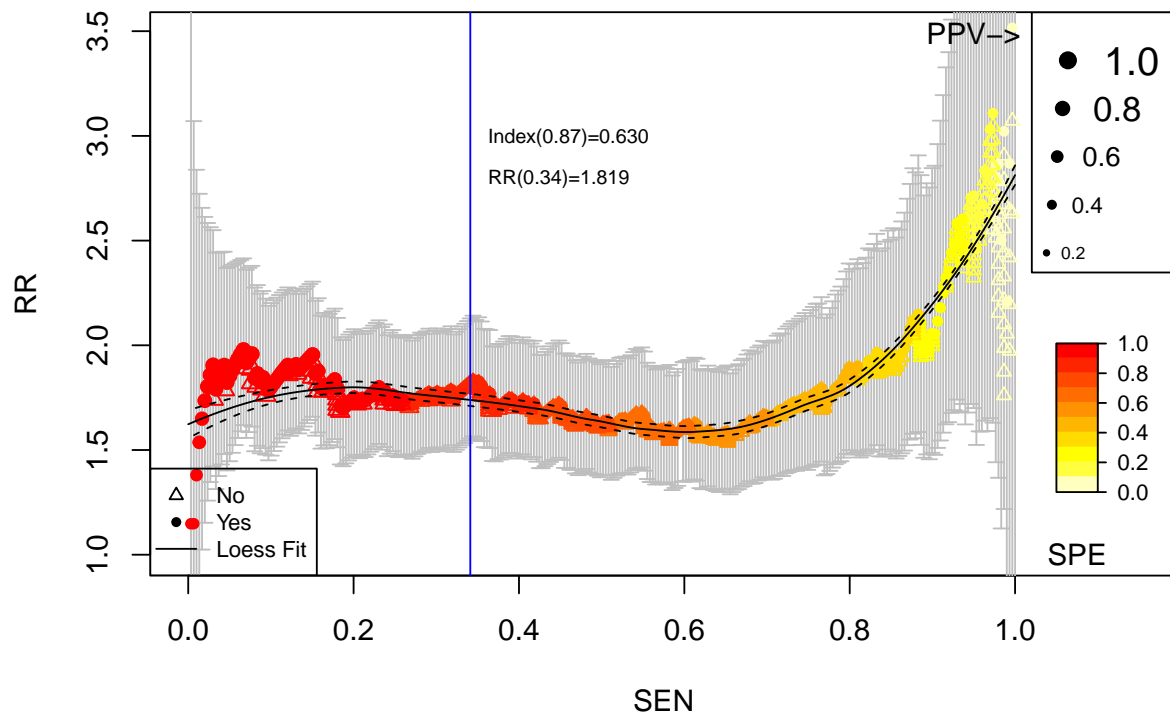
Cumulative vs. Observed: Test: Breast Cancer

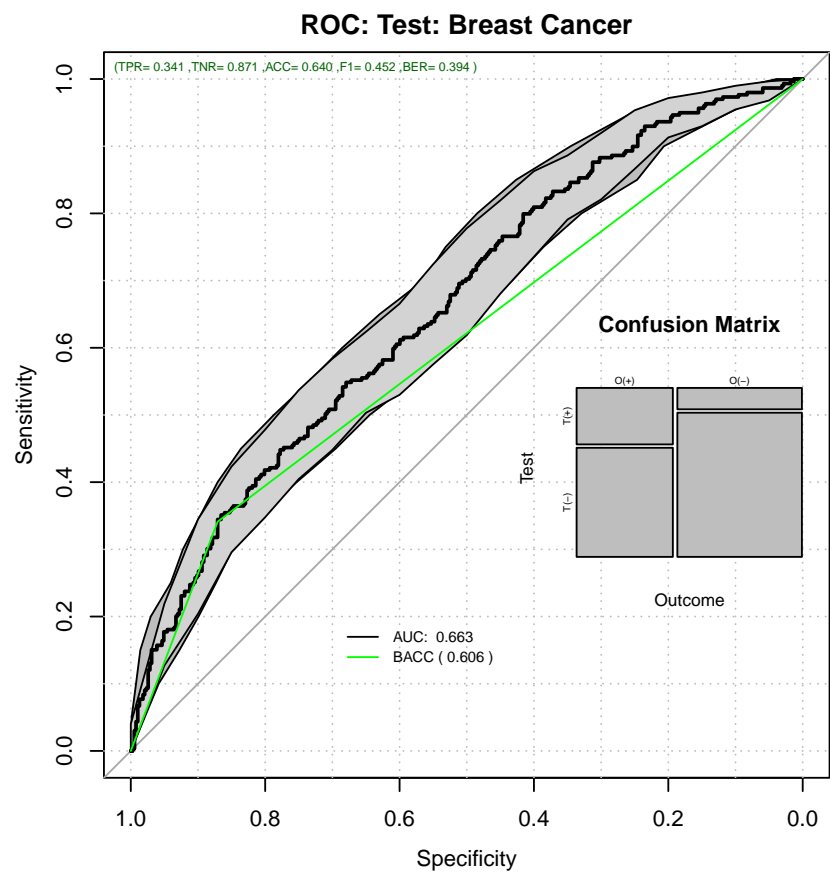


Decision Curve Analysis: Test: Breast Cancer

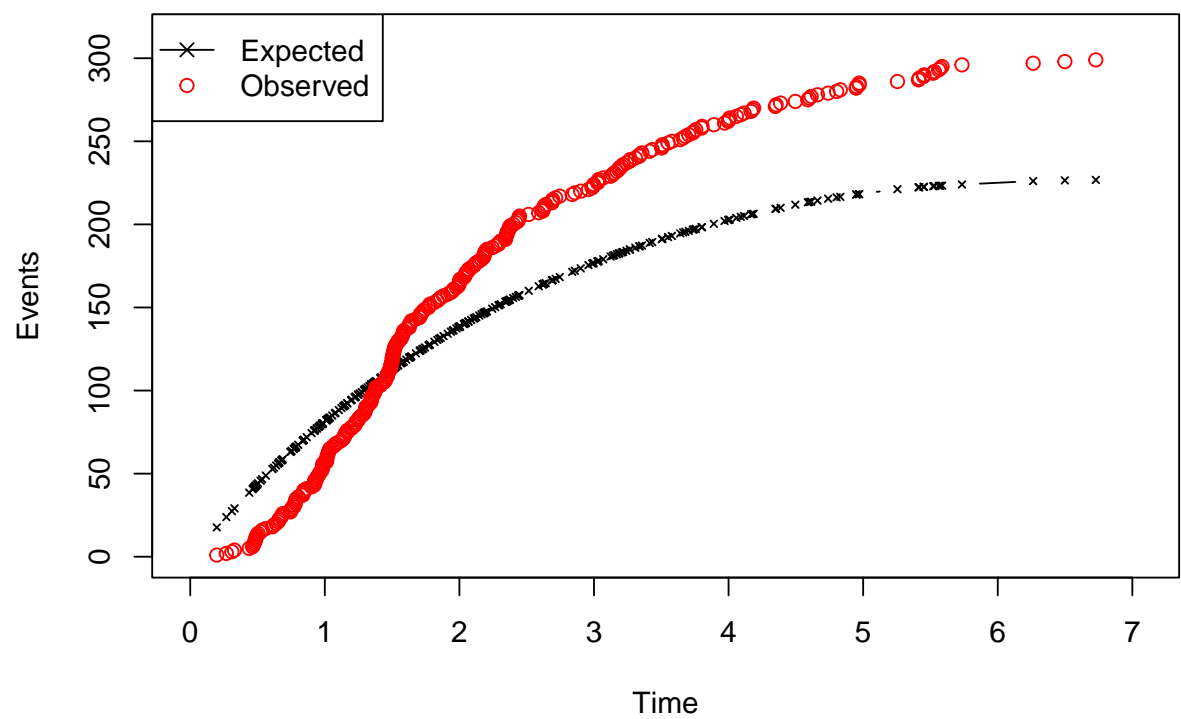


Relative Risk: Test: Breast Cancer

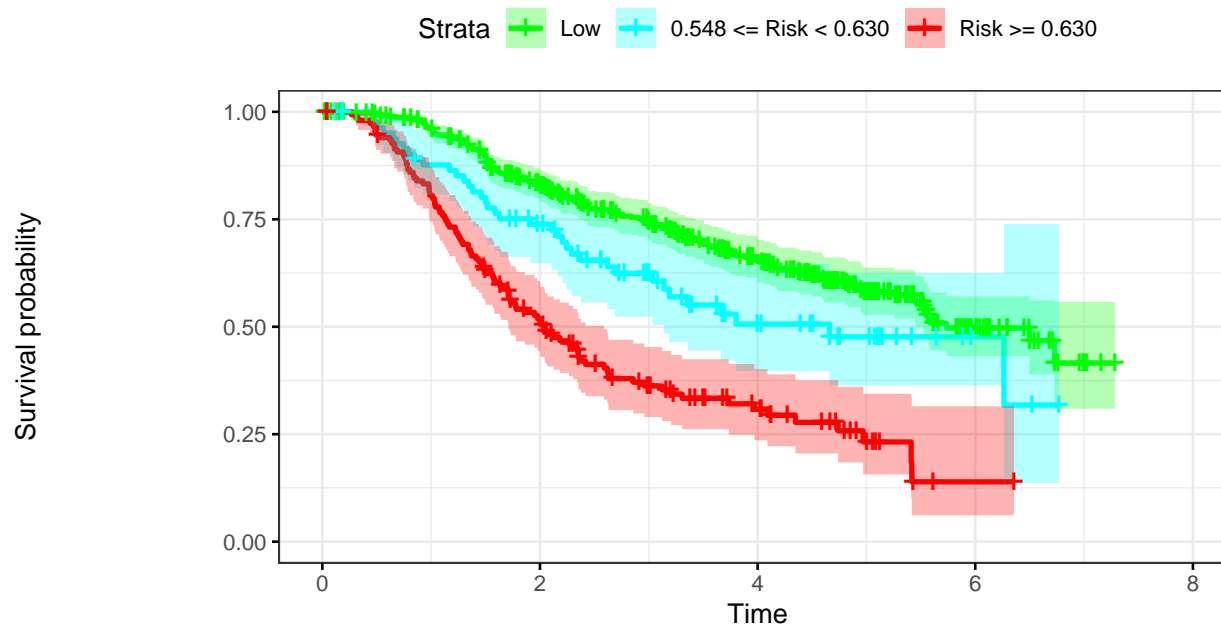




Time vs. Events: Test: Breast Cancer



Kaplan–Meier: Test: Breast Cancer



Number at risk

| | | | | | |
|-----------------------|-----|-----|-----|----|---|
| Low | 448 | 332 | 184 | 32 | 0 |
| 0.548 <= Risk < 0.630 | 86 | 55 | 21 | 3 | 0 |
| Risk >= 0.630 | 152 | 72 | 24 | 1 | 0 |

```
par(op)
```

1.5.1 External Data Report

```
pander::pander(t(rrCoxTestAnalysis$OERatio),caption="O/E Ratio")
```

Table 28: O/E Ratio

| est | lower | upper |
|------|-------|-------|
| 1.32 | 1.17 | 1.48 |

```
pander::pander(rrCoxTestAnalysis$c.index,caption="C. Index")
```

- C Index: 0.666
- Dxy: 0.331
- S.D.: 0.031
- n: 686
- missing: 0
- uncensored: 299
- Relevant Pairs: 266144

- **Concordant:** 177175
- **Uncertain:** 203702
- **cstatCI:**

| mean.C Index | median | lower | upper |
|--------------|--------|-------|-------|
| 0.666 | 0.665 | 0.637 | 0.696 |

```
pander::pander(t(rrCoxTestAnalysis$ROCAAnalysis$aucs),caption="ROC AUC")
```

Table 30: ROC AUC

| est | lower | upper |
|-------|-------|-------|
| 0.663 | 0.622 | 0.703 |

```
pander::pander((rrCoxTestAnalysis$ROCAAnalysis$sensitivity),caption="Sensitivity")
```

Table 31: Sensitivity

| est | lower | upper |
|-------|-------|-------|
| 0.341 | 0.288 | 0.398 |

```
pander::pander((rrCoxTestAnalysis$ROCAAnalysis$specificity),caption="Specificity")
```

Table 32: Specificity

| est | lower | upper |
|-------|-------|-------|
| 0.871 | 0.833 | 0.903 |

```
pander::pander(t(rrCoxTestAnalysis$thr_atP),caption="Probability Thresholds")
```

Table 33: Probability Thresholds

| 90% | 80% |
|------|-------|
| 0.63 | 0.548 |

```
pander::pander(t(rrCoxTestAnalysis$RR_atP),caption="Risk Ratio")
```

Table 34: Risk Ratio

| est | lower | upper |
|------|-------|-------|
| 1.82 | 1.55 | 2.13 |

```
pander::pander(rrCoxTestAnalysis$surdif,caption="Logrank test")
```

Table 35: Logrank test Chisq = 85.405348 on 2 degrees of freedom,
p = 0.000000

| | N | Observed | Expected | (O-E)^2/E | (O-E)^2/V |
|----------------|-----|----------|----------|-----------|-----------|
| class=0 | 448 | 160 | 219.0 | 15.89 | 60.334 |
| class=1 | 86 | 37 | 33.9 | 0.29 | 0.328 |
| class=2 | 152 | 102 | 46.1 | 67.59 | 81.340 |

1.5.2 Calibrating the index on the test data

```
calprob <- CoxRiskCalibration(ml,dataBrestCancerTest,"status","time")

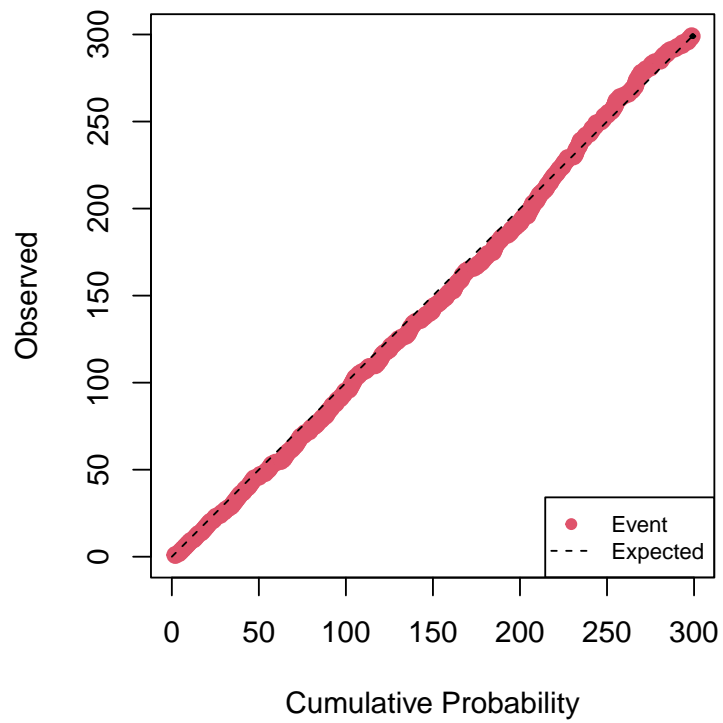
pander::pander(c(h0=calprob$h0,
                 Gain=calprob$hazardGain,
                 DeltaTime=calprob$timeInterval),
               caption="Cox Calibration Parameters")
```

| h0 | Gain | DeltaTime |
|-------|-------|-----------|
| 0.529 | 0.915 | 4.86 |

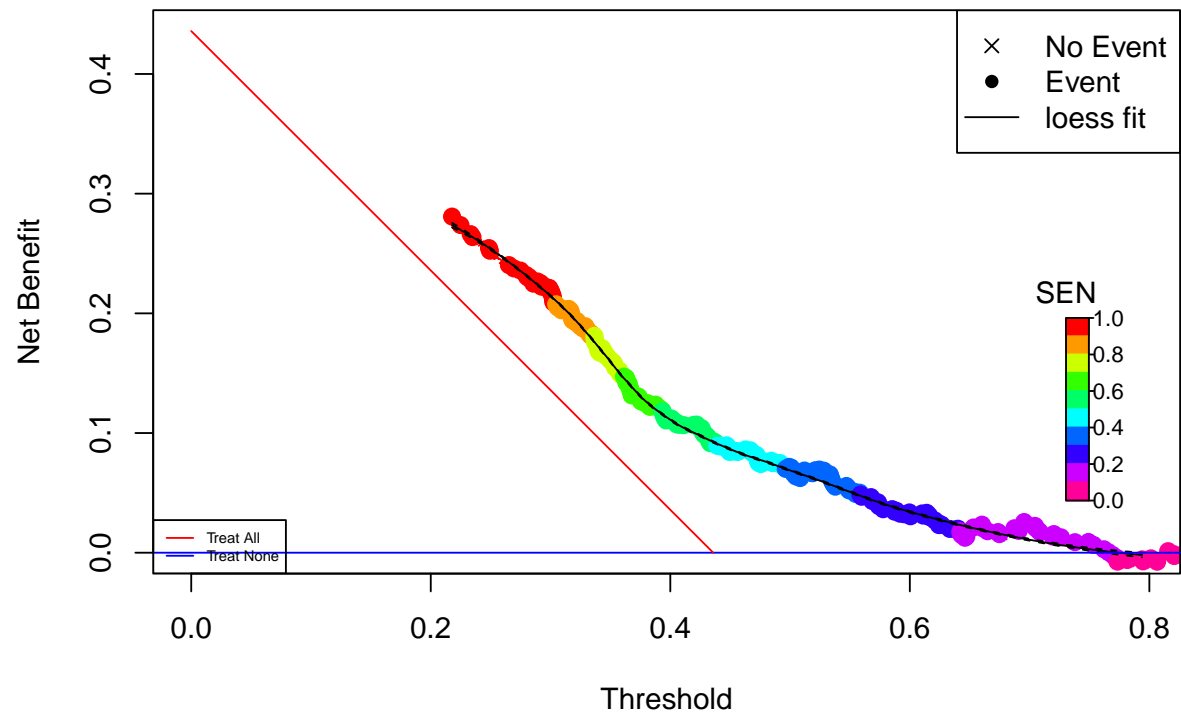
```
rdata <- cbind(dataBrestCancerTest$status,calprob$prob)

rrAnalysis <- RRPlot(rdata,atProb=c(0.90,0.80),
                    timetoEvent=dataBrestCancerTest$time,
                    title="Cal. Test: Breast Cancer",
                    ysurvlim=c(0.00,1.0),
                    riskTimeInterval=calprob$timeInterval)
```

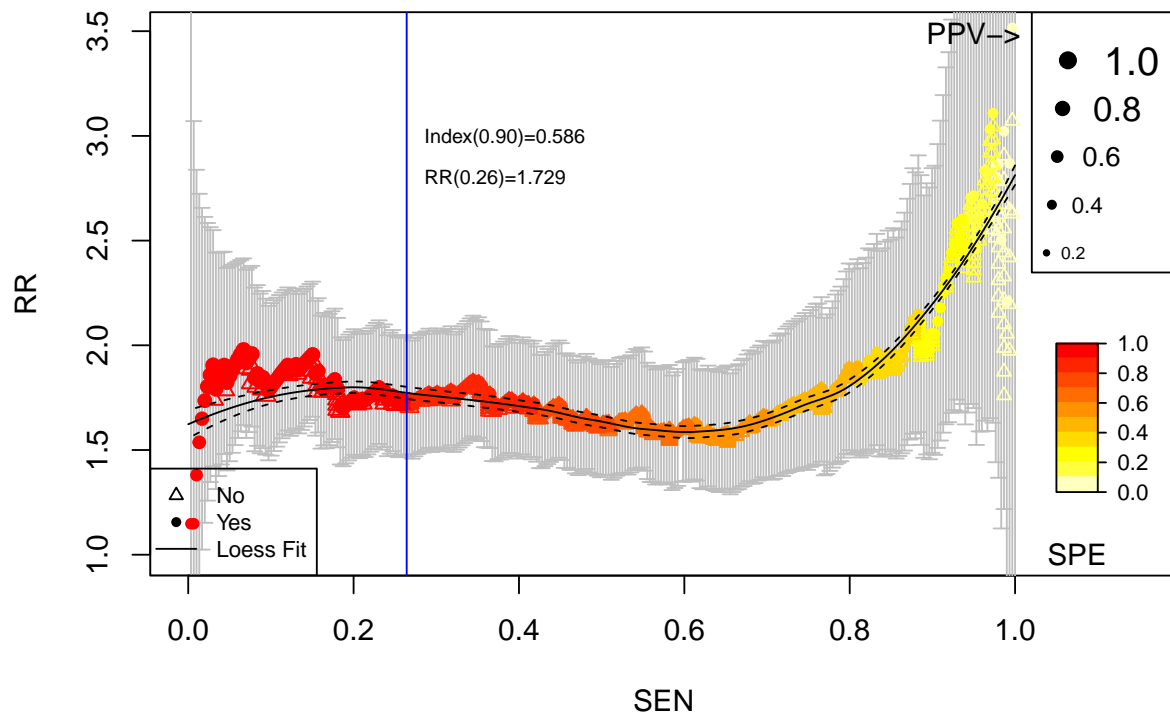
Cumulative vs. Observed: Cal. Test: Breast Cancer

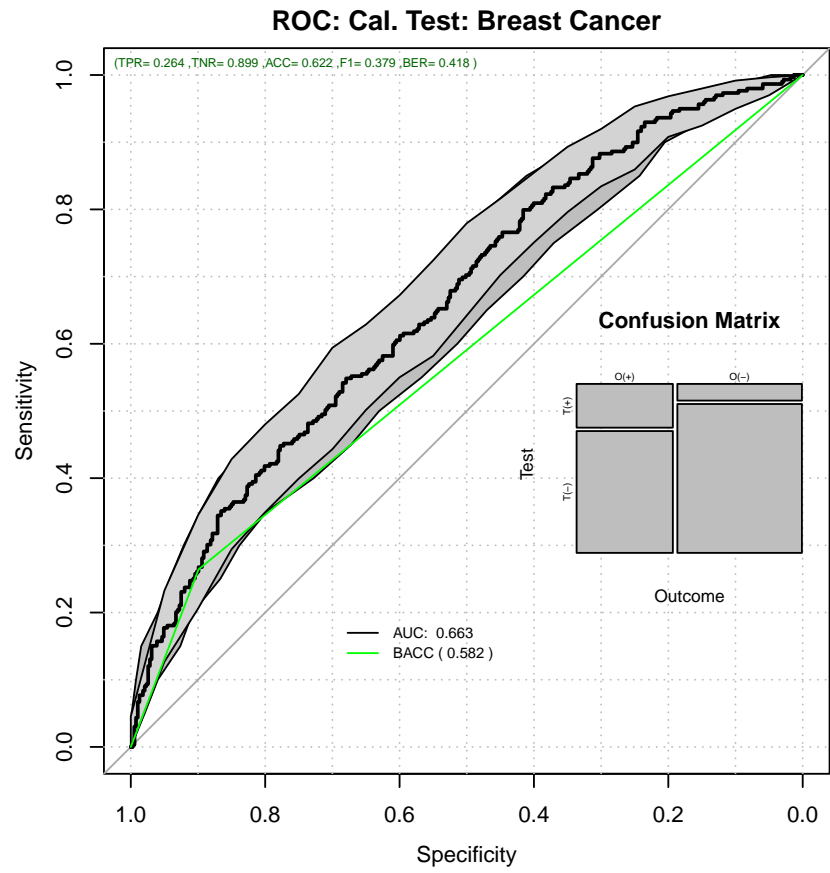


Decision Curve Analysis: Cal. Test: Breast Cancer

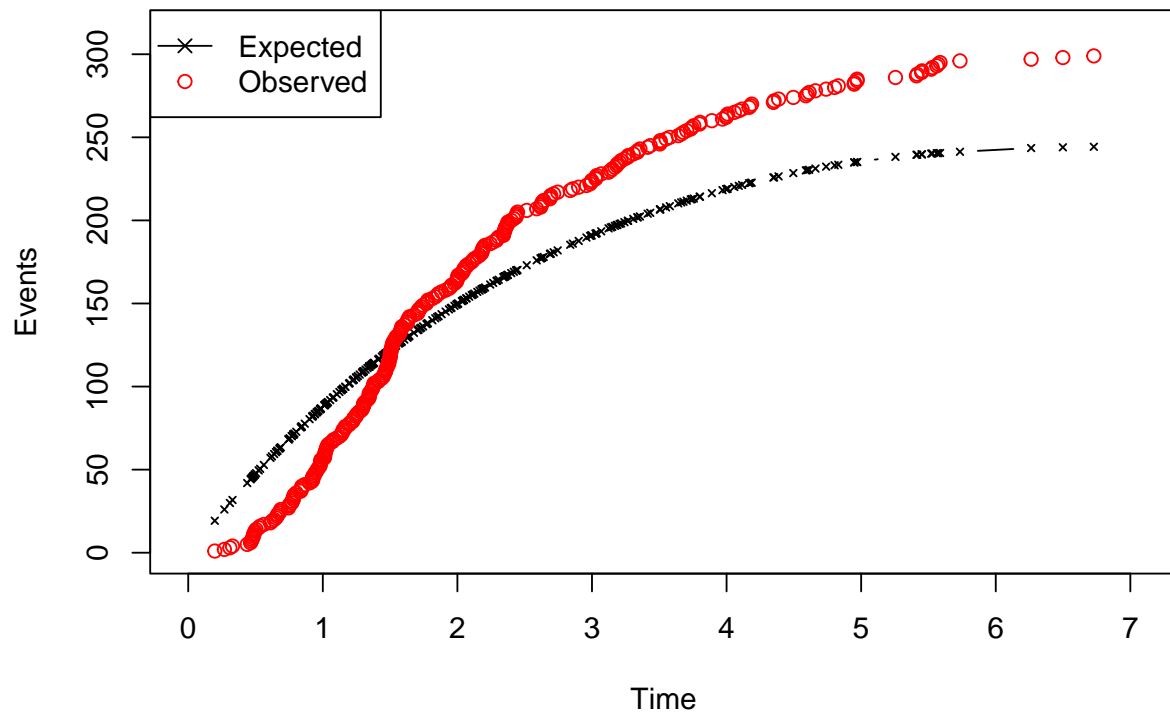


Relative Risk: Cal. Test: Breast Cancer

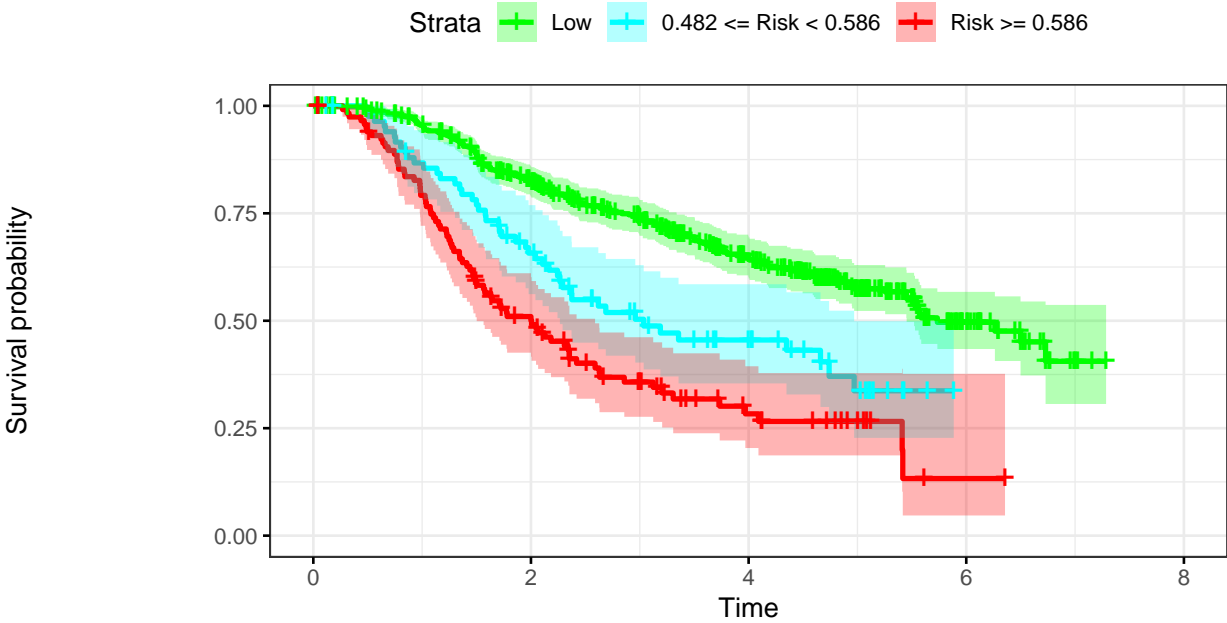




Time vs. Events: Cal. Test: Breast Cancer



Kaplan–Meier: Cal. Test: Breast Cancer



Number at risk

| | | | | | |
|-----------------------|-----|-----|-----|----|---|
| Low | 483 | 354 | 190 | 35 | 0 |
| 0.482 <= Risk < 0.586 | 85 | 51 | 23 | 0 | 0 |
| Risk >= 0.586 | 118 | 54 | 16 | 1 | 0 |

1.5.3 After Calibration Report

```
pander::pander(t(rrAnalysis$OERatio),caption="O/E Ratio")
```

Table 37: O/E Ratio

| est | lower | upper |
|------|-------|-------|
| 1.22 | 1.09 | 1.37 |

```
pander::pander(rrAnalysis$c.index,caption="C. Index")
```

- C Index: 0.666
- Dxy: 0.331
- S.D.: 0.031
- n: 686
- missing: 0
- uncensored: 299
- Relevant Pairs: 266144
- Concordant: 177175
- Uncertain: 203702

- `cstatCI`:

| mean.C Index | median | lower | upper |
|--------------|--------|-------|-------|
| 0.666 | 0.666 | 0.635 | 0.695 |

```
pander::pander(t(rrAnalysis$ROCAAnalysis$aucs),caption="ROC AUC")
```

Table 39: ROC AUC

| est | lower | upper |
|-------|-------|-------|
| 0.663 | 0.622 | 0.703 |

```
pander::pander((rrAnalysis$ROCAAnalysis$sensitivity),caption="Sensitivity")
```

Table 40: Sensitivity

| est | lower | upper |
|-------|-------|-------|
| 0.264 | 0.215 | 0.318 |

```
pander::pander((rrAnalysis$ROCAAnalysis$specificity),caption="Specificity")
```

Table 41: Specificity

| est | lower | upper |
|-------|-------|-------|
| 0.899 | 0.865 | 0.927 |

```
pander::pander(t(rrAnalysis$thr_atP),caption="Probability Thresholds")
```

Table 42: Probability Thresholds

| 90% | 80% |
|-------|-------|
| 0.586 | 0.482 |

```
pander::pander(t(rrAnalysis$RR_atP),caption="Risk Ratio")
```

Table 43: Risk Ratio

| est | lower | upper |
|------|-------|-------|
| 1.73 | 1.47 | 2.04 |

```
pander::pander(rrAnalysis$surdif,caption="Logrank test")
```

Table 44: Logrank test Chisq = 80.478762 on 2 degrees of freedom,
p = 0.000000

| | N | Observed | Expected | (O-E)^2/E | (O-E)^2/V |
|----------------|-----|----------|----------|-----------|-----------|
| class=0 | 483 | 174 | 232.7 | 14.80 | 67.96 |
| class=1 | 85 | 46 | 32.0 | 6.16 | 6.94 |
| class=2 | 118 | 79 | 34.3 | 58.05 | 66.45 |

1.6 Logistic Model

Here we train a logistic model on the same data set

```
## Only label subjects that present event withing five years

dataBrestCancerR <- subset(dataBrestCancerTrain, time>=5 | status==1)
dataBrestCancerR$status <- dataBrestCancerR$status * (dataBrestCancerR$time < 5)
dataBrestCancerR$time <- NULL

#ml <- BSWiMS.model(status~1,data=dataBrestCancerR,loops=20,NumberofRepeats = 5)
mlog <- BSWiMS.model(status~1,data=dataBrestCancerR,loops=1,NumberofRepeats = 5)

---..

sm <- summary(mlog)
pander::pander(sm$coefficients)
```

Table 45: Table continues below

| | Estimate | lower | OR | upper | u.Accuracy | r.Accuracy |
|---------------------|-----------|-------|-------|-------|------------|------------|
| size_nodes | 1.05e-03 | 1.001 | 1.001 | 1.001 | 0.669 | 0.571 |
| nodes | 4.33e-02 | 1.040 | 1.044 | 1.048 | 0.676 | 0.634 |
| grade_nodes | 1.50e-02 | 1.014 | 1.015 | 1.016 | 0.682 | 0.637 |
| age_nodes | 1.06e-03 | 1.001 | 1.001 | 1.001 | 0.678 | 0.653 |
| size_grade | 1.75e-03 | 1.001 | 1.002 | 1.002 | 0.632 | 0.682 |
| age_size | 8.73e-05 | 1.000 | 1.000 | 1.000 | 0.608 | 0.682 |
| grade | 2.27e-01 | 1.168 | 1.254 | 1.347 | 0.571 | 0.683 |
| age_meno | -6.04e-03 | 0.992 | 0.994 | 0.996 | 0.571 | 0.676 |
| age_pgr | -5.42e-06 | 1.000 | 1.000 | 1.000 | 0.571 | 0.686 |
| age_grade | -1.65e-03 | 0.997 | 0.998 | 0.999 | 0.574 | 0.690 |
| meno_grade | 1.02e-01 | 1.045 | 1.107 | 1.173 | 0.571 | 0.683 |
| nodes_hormon | -1.38e-02 | 0.979 | 0.986 | 0.994 | 0.587 | 0.688 |
| size | 3.94e-03 | 1.002 | 1.004 | 1.006 | 0.611 | 0.693 |
| meno_pgr | 3.19e-04 | 1.000 | 1.000 | 1.001 | 0.571 | 0.687 |
| pgr | -1.07e-04 | 1.000 | 1.000 | 1.000 | 0.571 | 0.689 |
| meno_nodes | -2.60e-02 | 0.955 | 0.974 | 0.994 | 0.640 | 0.686 |
| grade_pgr | -3.51e-05 | 1.000 | 1.000 | 1.000 | 0.571 | 0.669 |
| meno_size | 2.34e-03 | 1.000 | 1.002 | 1.004 | 0.604 | 0.691 |

Table 46: Table continues below

| | full.Accuracy | u.AUC | r.AUC | full.AUC | IDI |
|-------------------|---------------|-------|-------|----------|---------|
| size_nodes | 0.668 | 0.627 | 0.500 | 0.628 | 0.11233 |

| | full.Accuracy | u.AUC | r.AUC | full.AUC | IDI |
|--------------|---------------|-------|-------|----------|---------|
| nodes | 0.690 | 0.639 | 0.621 | 0.662 | 0.07110 |
| grade_nodes | 0.686 | 0.649 | 0.624 | 0.655 | 0.06580 |
| age_nodes | 0.686 | 0.642 | 0.621 | 0.657 | 0.03346 |
| size_grade | 0.686 | 0.626 | 0.646 | 0.655 | 0.01787 |
| age_size | 0.686 | 0.577 | 0.649 | 0.657 | 0.01534 |
| grade | 0.690 | 0.500 | 0.653 | 0.662 | 0.01340 |
| age_meno | 0.686 | 0.500 | 0.645 | 0.657 | 0.00782 |
| age_pgr | 0.686 | 0.500 | 0.656 | 0.657 | 0.00512 |
| age_grade | 0.690 | 0.507 | 0.661 | 0.662 | 0.00454 |
| meno_grade | 0.686 | 0.500 | 0.652 | 0.657 | 0.00425 |
| nodes_hormon | 0.686 | 0.526 | 0.658 | 0.655 | 0.00280 |
| size | 0.690 | 0.618 | 0.663 | 0.662 | 0.00507 |
| meno_pgr | 0.686 | 0.500 | 0.657 | 0.657 | 0.00316 |
| pgr | 0.686 | 0.500 | 0.659 | 0.655 | 0.00257 |
| meno_nodes | 0.686 | 0.595 | 0.656 | 0.657 | 0.00264 |
| grade_pgr | 0.668 | 0.500 | 0.627 | 0.628 | 0.00241 |
| meno_size | 0.690 | 0.578 | 0.663 | 0.662 | 0.00185 |

| | NRI | z.IDI | z.NRI | Delta.AUC | Frequency |
|--------------|----------|-------|--------|-----------|-----------|
| size_nodes | 0.63654 | 17.86 | 18.870 | 0.128490 | 1 |
| nodes | 0.57106 | 14.13 | 16.179 | 0.040494 | 1 |
| grade_nodes | 0.54866 | 13.66 | 15.650 | 0.031087 | 1 |
| age_nodes | 0.21312 | 9.39 | 5.710 | 0.035896 | 1 |
| size_grade | 0.29411 | 6.74 | 7.728 | 0.008648 | 1 |
| age_size | 0.29152 | 6.41 | 7.652 | 0.007600 | 1 |
| grade | 0.19036 | 6.20 | 4.983 | 0.008461 | 1 |
| age_meno | 0.08057 | 4.76 | 2.337 | 0.012065 | 1 |
| age_pgr | 0.00745 | 4.11 | 0.194 | 0.000417 | 1 |
| age_grade | 0.11372 | 3.60 | 2.960 | 0.000315 | 1 |
| meno_grade | 0.20428 | 3.47 | 5.343 | 0.004441 | 1 |
| nodes_hormon | 0.45522 | 3.44 | 12.150 | -0.002853 | 1 |
| size | 0.21050 | 3.42 | 5.600 | -0.001075 | 1 |
| meno_pgr | 0.05977 | 3.35 | 1.558 | -0.000429 | 1 |
| pgr | 0.19759 | 2.64 | 5.745 | -0.004123 | 1 |
| meno_nodes | -0.06329 | 2.59 | -1.645 | 0.000631 | 1 |
| grade_pgr | 0.17471 | 2.55 | 5.058 | 0.001252 | 1 |
| meno_size | 0.10227 | 2.43 | 2.662 | -0.001378 | 1 |

1.7 Logistic Model Performance

```

op <- par(no.readonly = TRUE)

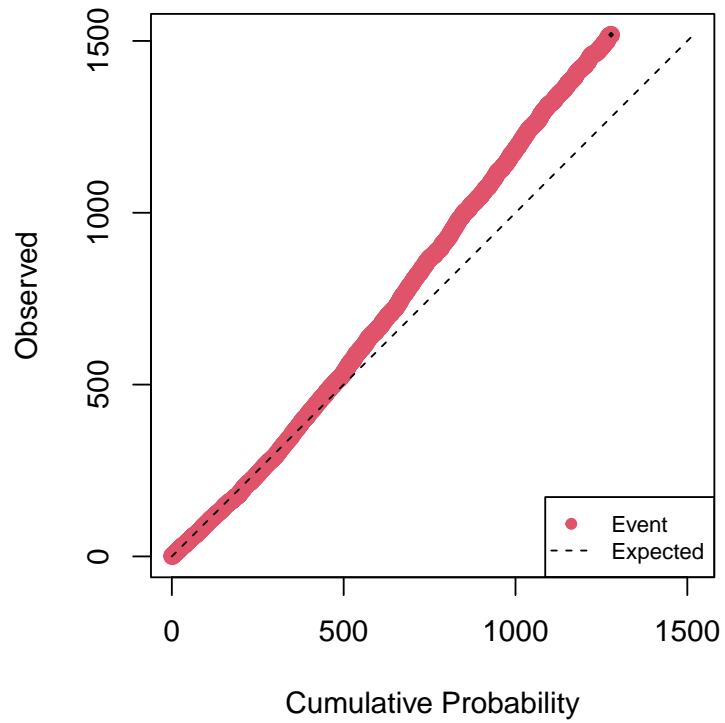
cprob <- predict(mlog,dataBrestCancerTrain)

rdata <- cbind(dataBrestCancerTrain$status,cprob)
rrAnalysisTrain <- RRPlot(rdata,atProb=c(0.90,0.80),
                           timetoEvent=dataBrestCancerTrain$time,
                           title="Logistic Train: Breast Cancer",

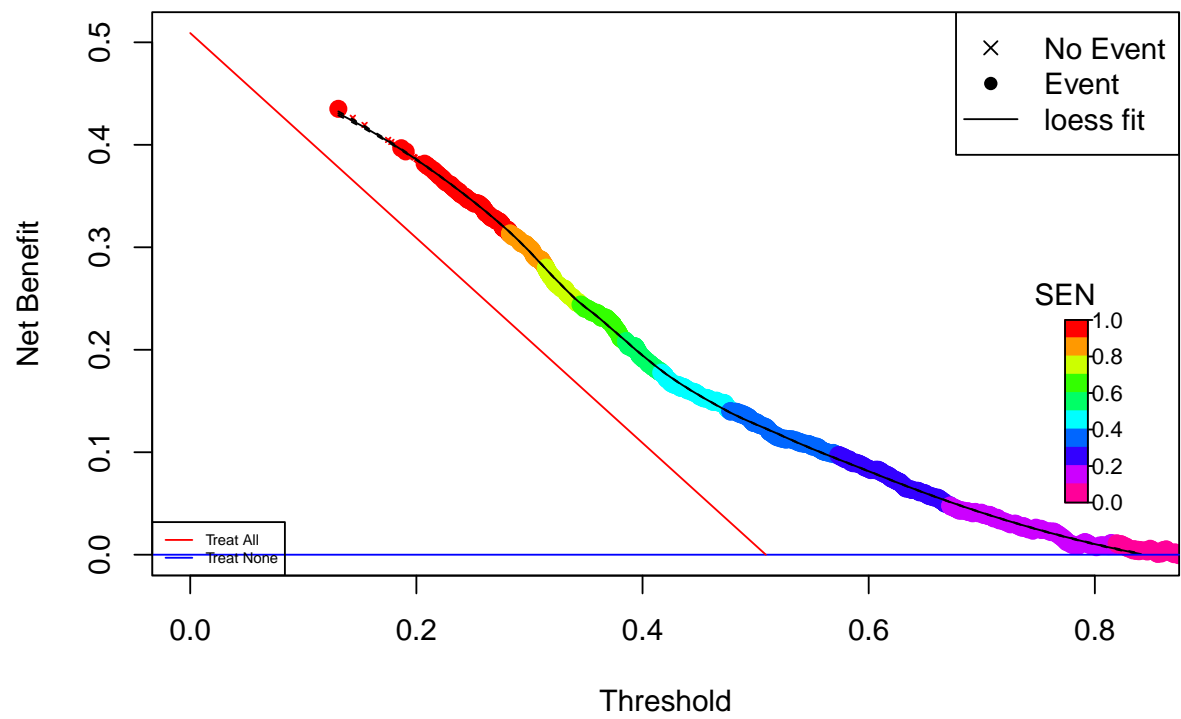
```

```
ysurvlim=c(0.00,1.0),  
riskTimeInterval=5.0)
```

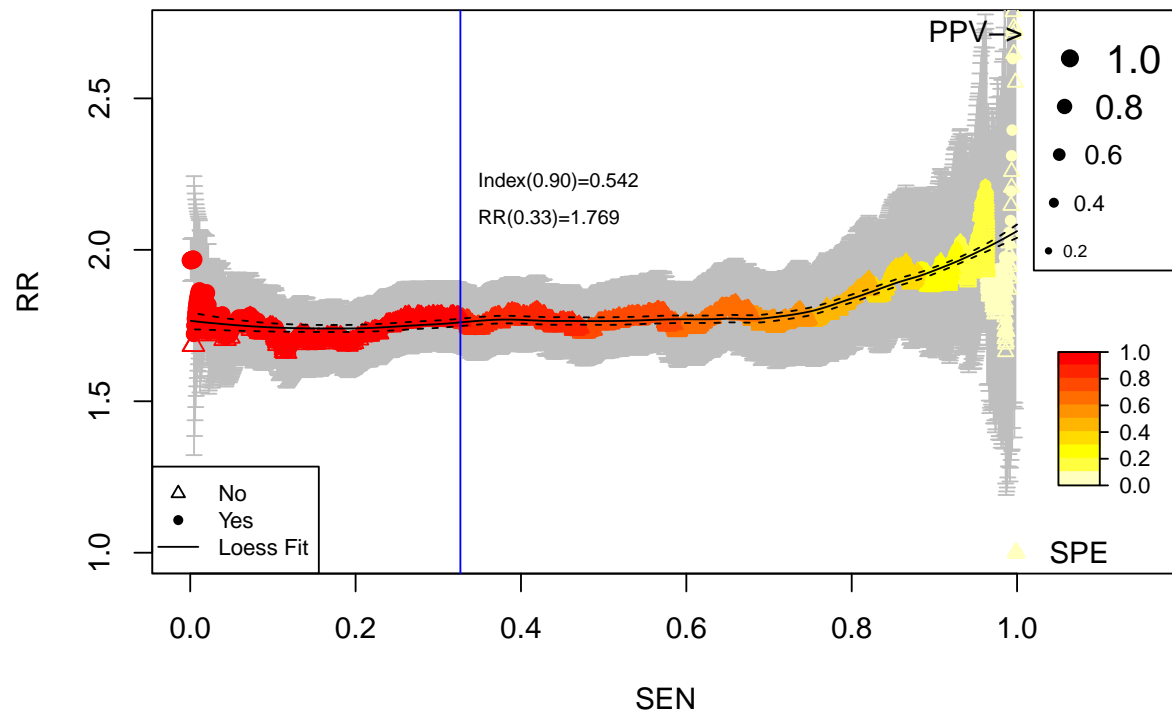
Cumulative vs. Observed: Logistic Train: Breast Cancer

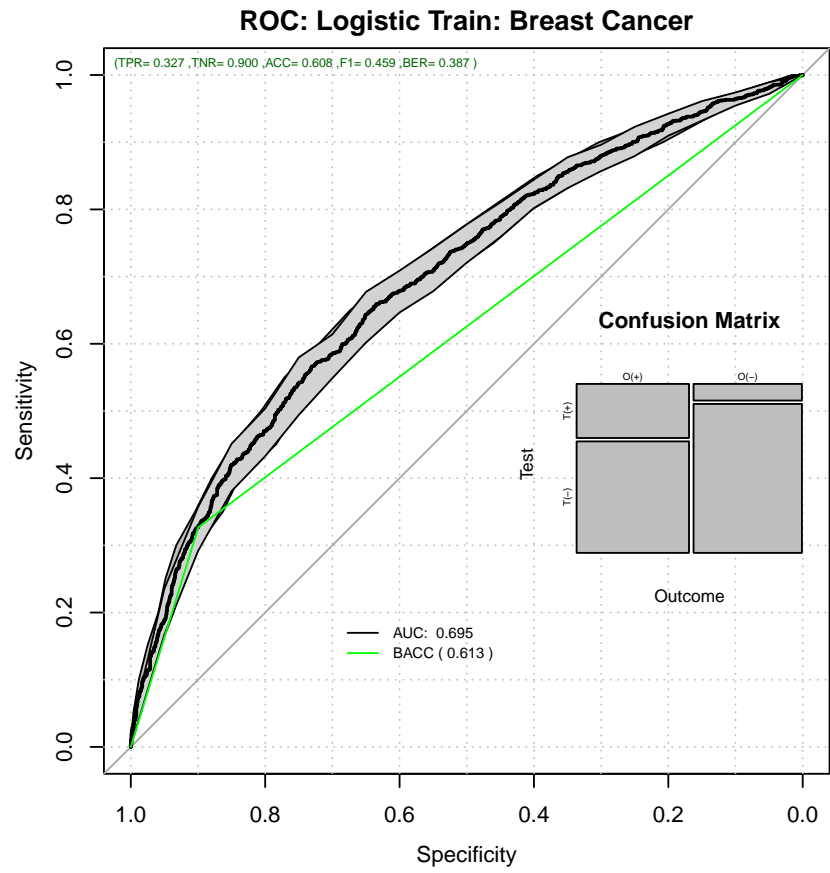


Decision Curve Analysis: Logistic Train: Breast Cancer

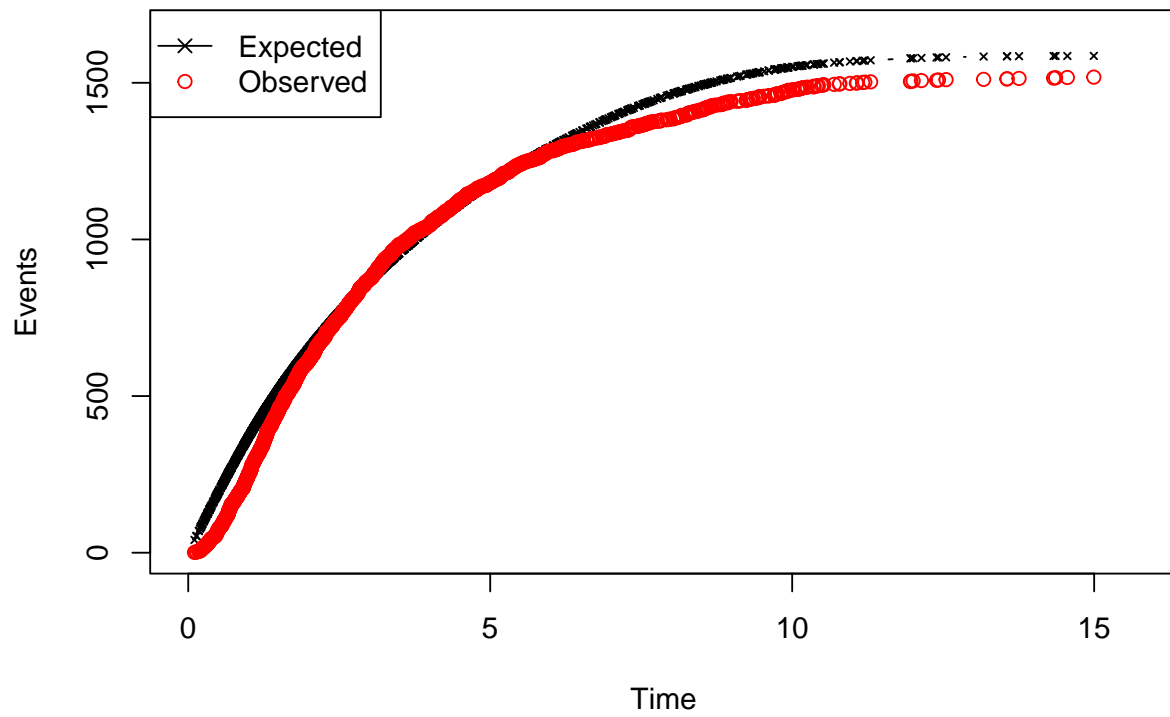


Relative Risk: Logistic Train: Breast Cancer

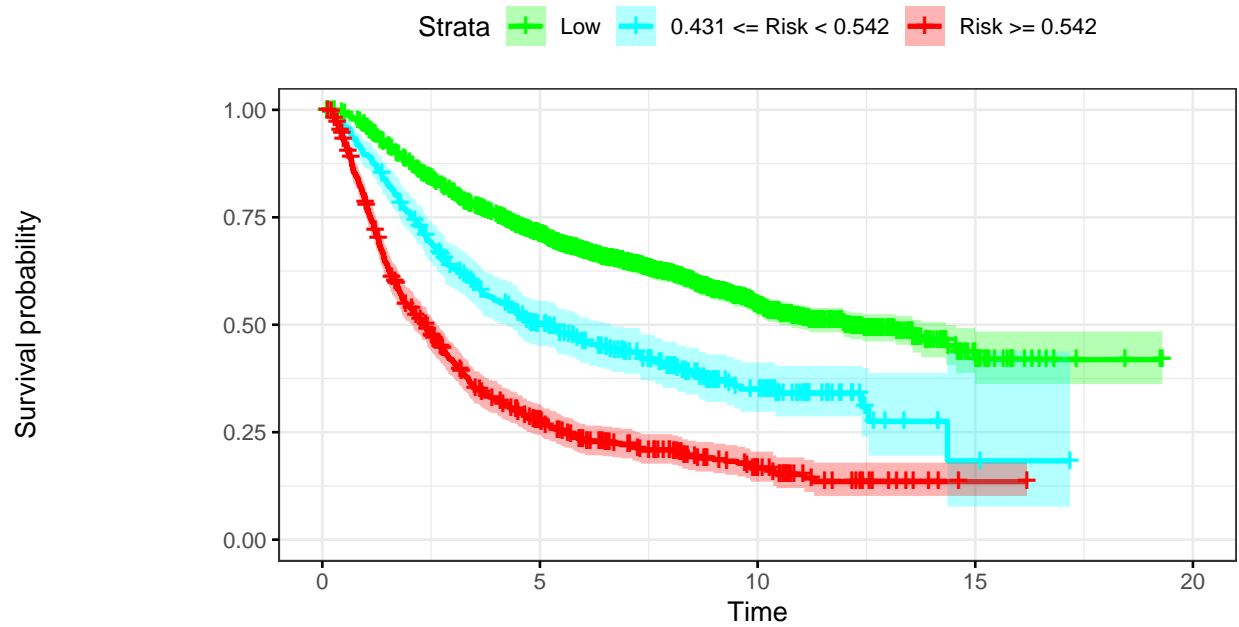




Time vs. Events: Logistic Train: Breast Cancer



Kaplan–Meier: Logistic Train: Breast Cancer



Number at risk

| | | | | | |
|-----------------------|------|------|-----|----|---|
| Low | 1975 | 1268 | 399 | 23 | 0 |
| 0.431 <= Risk < 0.542 | 364 | 160 | 47 | 2 | 0 |
| Risk >= 0.542 | 643 | 143 | 37 | 1 | 0 |

```
par(op)
```

1.7.1 Training Report

```
pander::pander(t(rrAnalysisTrain$OERatio),caption="O/E Ratio")
```

Table 48: O/E Ratio

| est | lower | upper |
|-------|-------|-------|
| 0.957 | 0.91 | 1.01 |

```
pander::pander(rrAnalysisTrain$c.index,caption="C. Index")
```

- C Index: 0.68
- Dxy: 0.36
- S.D.: 0.014
- n: 2982
- missing: 0
- uncensored: 1518
- Relevant Pairs: 6184528

- **Concordant:** *4206588*
- **Uncertain:** *2703838*
- **cstatCI:**

| mean.C Index | median | lower | upper |
|--------------|--------|-------|-------|
| 0.68 | 0.68 | 0.665 | 0.693 |

```
pander::pander(t(rrAnalysisTrain$ROCAalysis$aucs),caption="ROC AUC")
```

Table 50: ROC AUC

| est | lower | upper |
|-------|-------|-------|
| 0.695 | 0.677 | 0.714 |

```
pander::pander((rrAnalysisTrain$ROCAalysis$sensitivity),caption="Sensitivity")
```

Table 51: Sensitivity

| est | lower | upper |
|-------|-------|-------|
| 0.327 | 0.303 | 0.351 |

```
pander::pander((rrAnalysisTrain$ROCAalysis$specificity),caption="Specificity")
```

Table 52: Specificity

| est | lower | upper |
|-----|-------|-------|
| 0.9 | 0.883 | 0.915 |

```
pander::pander(t(rrAnalysisTrain$thr_atP),caption="Probability Thresholds")
```

Table 53: Probability Thresholds

| 90% | 80% |
|-------|-------|
| 0.542 | 0.431 |

```
pander::pander(t(rrAnalysisTrain$RR_atP),caption="Risk Ratio")
```

Table 54: Risk Ratio

| est | lower | upper |
|------|-------|-------|
| 1.77 | 1.66 | 1.88 |

```
pander::pander(rrAnalysisTrain$surdif,caption="Logrank test")
```

Table 55: Logrank test $\text{Chisq} = 543.347175$ on 2 degrees of freedom,
 $p = 0.000000$

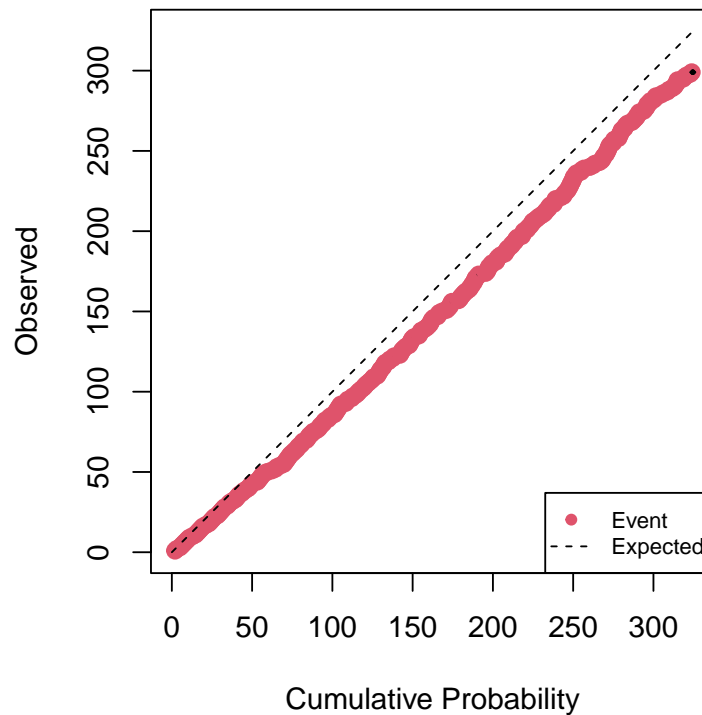
| | N | Observed | Expected | $(O-E)^2/E$ | $(O-E)^2/V$ |
|---------|------|----------|----------|-------------|-------------|
| class=0 | 1975 | 804 | 1145 | 101.5 | 418.9 |
| class=1 | 364 | 218 | 169 | 14.1 | 15.9 |
| class=2 | 643 | 496 | 204 | 418.2 | 490.7 |

1.7.2 Results on the validation set using Logistic model

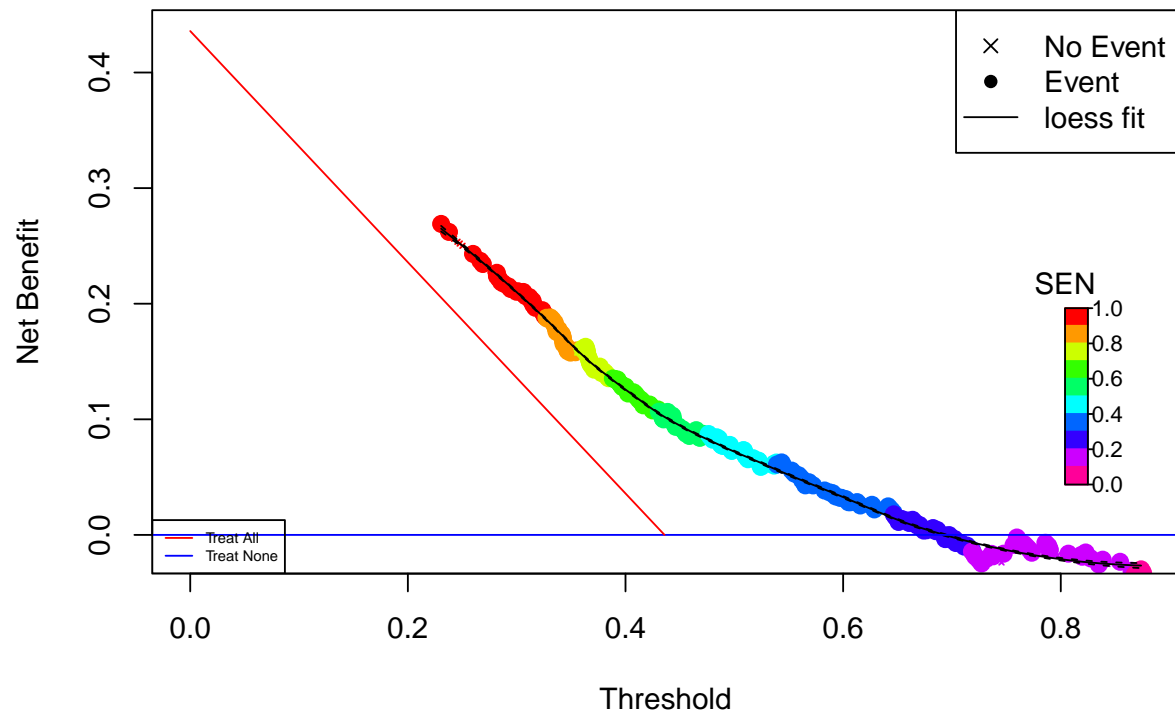
```
pre <- predict(mlog,dataBrestCancerTest)
rdata <- cbind(dataBrestCancerTest$status,pre)

rrAnalysis <- RRPlot(rdata,atThr=rrAnalysisTrain$thr_atP,
                     timetoEvent=dataBrestCancerTest$time,
                     title="Logistic Test: Breast Cancer",
                     ysurvlim=c(0.00,1.0),
                     riskTimeInterval=5)
```

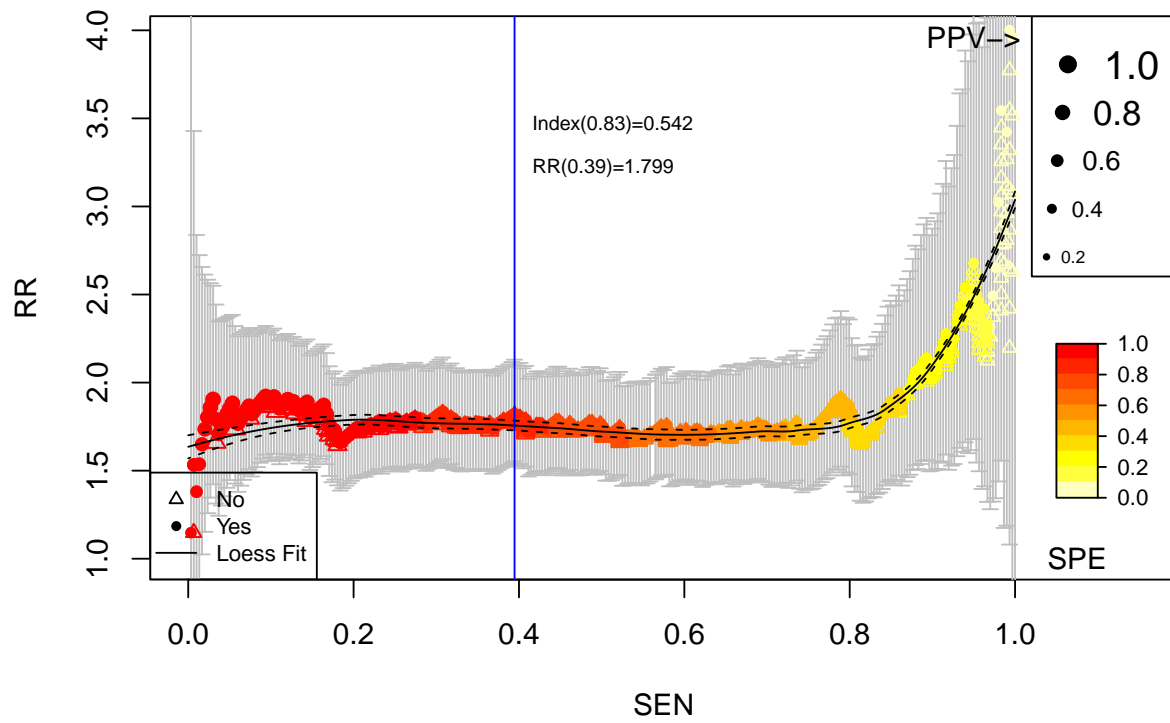
Cumulative vs. Observed: Logistic Test: Breast Cancer

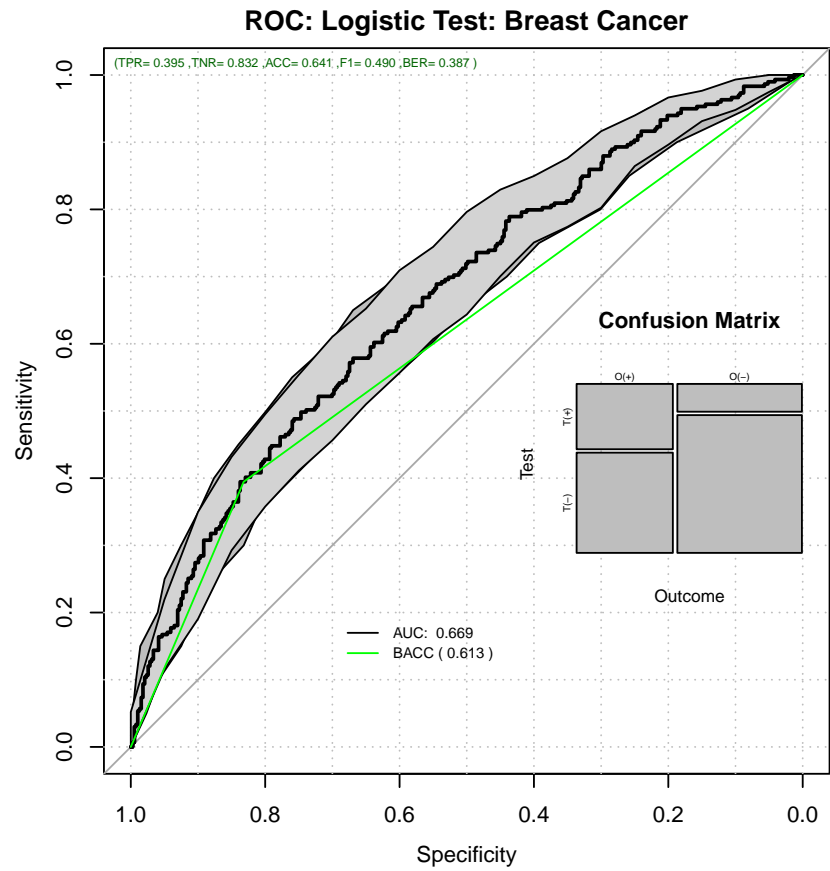


Decision Curve Analysis: Logistic Test: Breast Cancer

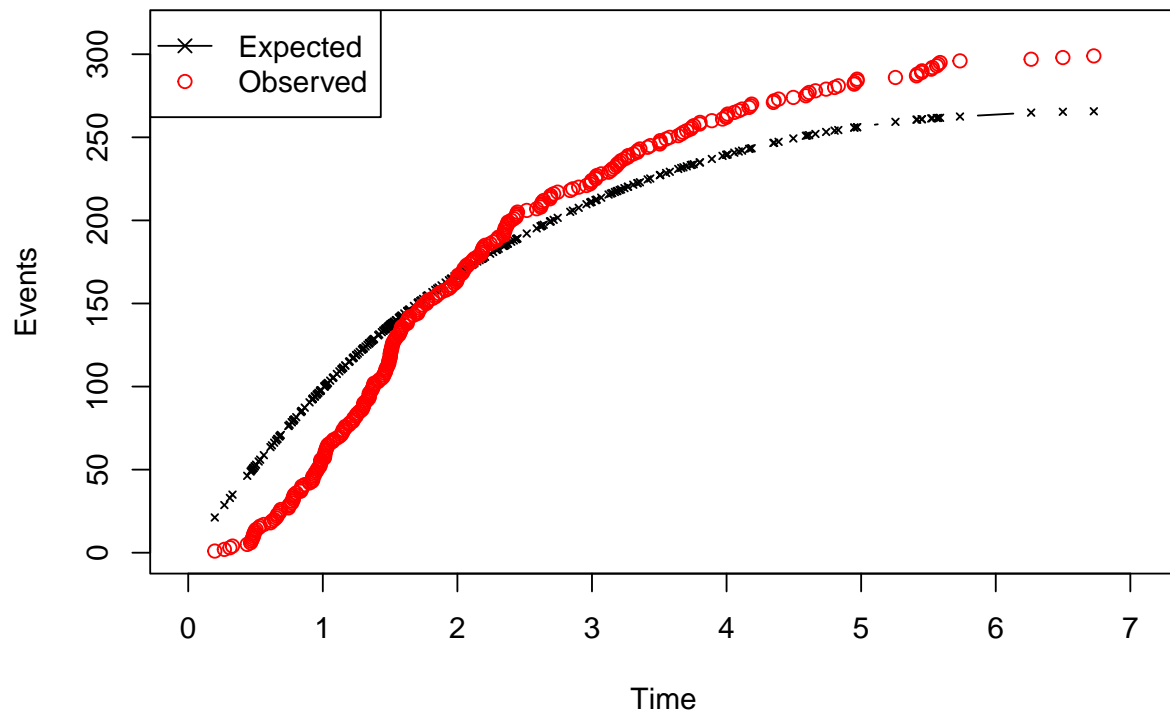


Relative Risk: Logistic Test: Breast Cancer

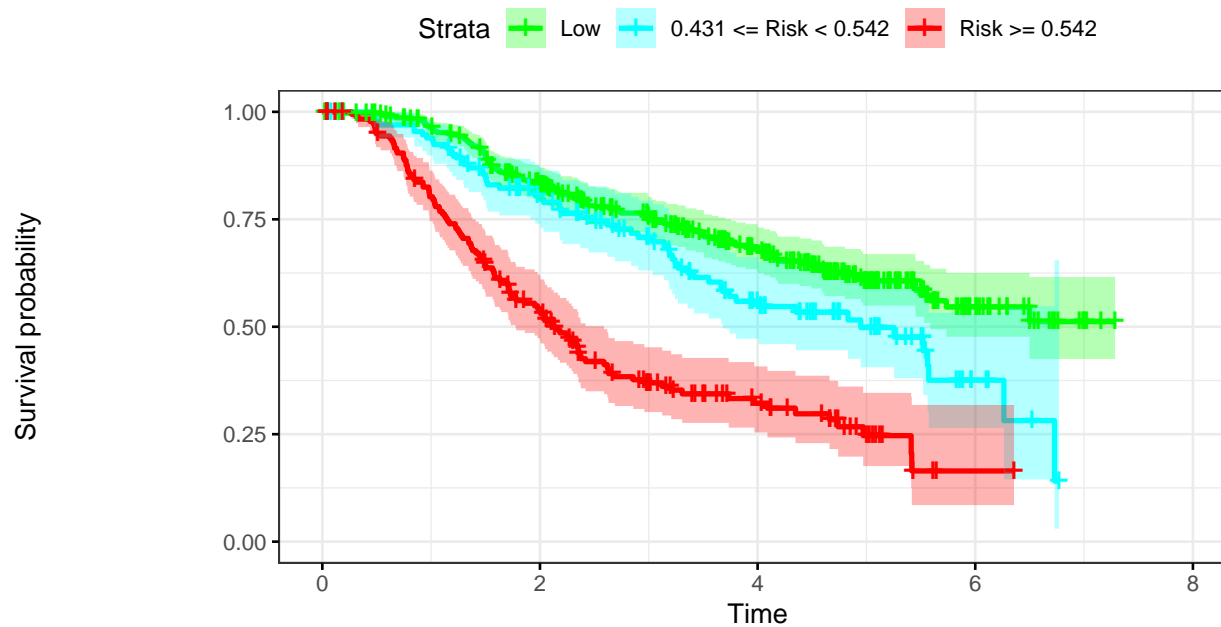




Time vs. Events: Logistic Test: Breast Cancer



Kaplan–Meier: Logistic Test: Breast Cancer



Number at risk

| | | | | | |
|-----------------------|-----|-----|-----|----|---|
| Low | 369 | 274 | 154 | 29 | 0 |
| 0.431 <= Risk < 0.542 | 134 | 96 | 46 | 6 | 0 |
| Risk >= 0.542 | 183 | 89 | 29 | 1 | 0 |

```
par(op)
```

1.7.3 Validation Report

```
pander::pander(t(rrAnalysis$OERatio),caption="O/E Ratio")
```

Table 56: O/E Ratio

| est | lower | upper |
|------|-------|-------|
| 1.13 | 1 | 1.26 |

```
pander::pander(rrAnalysis$c.index,caption="C. Index")
```

- C Index: 0.669
- Dxy: 0.338
- S.D.: 0.0309
- n: 686
- missing: 0
- uncensored: 299
- Relevant Pairs: 266144

- **Concordant:** 178115
- **Uncertain:** 203702
- **cstatCI:**

| mean.C Index | median | lower | upper |
|--------------|--------|-------|-------|
| 0.669 | 0.669 | 0.637 | 0.699 |

```
pander::pander(t(rrAnalysis$ROCAAnalysis$aucs),caption="ROC AUC")
```

Table 58: ROC AUC

| est | lower | upper |
|-------|-------|-------|
| 0.669 | 0.628 | 0.709 |

```
pander::pander((rrAnalysis$ROCAAnalysis$sensitivity),caption="Sensitivity")
```

Table 59: Sensitivity

| est | lower | upper |
|-------|-------|-------|
| 0.395 | 0.339 | 0.453 |

```
pander::pander((rrAnalysis$ROCAAnalysis$specificity),caption="Specificity")
```

Table 60: Specificity

| est | lower | upper |
|-------|-------|-------|
| 0.832 | 0.791 | 0.868 |

```
pander::pander(t(rrAnalysis$thr_atP),caption="Probability Thresholds")
```

Table 61: Probability Thresholds

| 90% | 80% |
|-------|-------|
| 0.542 | 0.431 |

```
pander::pander(t(rrAnalysis$RR_atP),caption="Risk Ratio")
```

Table 62: Risk Ratio

| est | lower | upper |
|-----|-------|-------|
| 1.8 | 1.54 | 2.11 |

```
pander::pander(rrAnalysis$surdif,caption="Logrank test")
```

Table 63: Logrank test $\text{Chisq} = 92.507991$ on 2 degrees of freedom,
 $p = 0.000000$

| | N | Observed | Expected | (O-E)^2/E | (O-E)^2/V |
|----------------|-----|----------|----------|-----------|-----------|
| class=0 | 369 | 121 | 181.7 | 20.2997 | 52.3868 |
| class=1 | 134 | 60 | 61.7 | 0.0479 | 0.0604 |
| class=2 | 183 | 118 | 55.5 | 70.2342 | 88.0195 |

1.8 Logistic Model Poisson Calibration

```
riskdata <- cbind(dataBrestCancerTrain$status, predict(mlog, dataBrestCancerTrain, type="prob"), dataBrestC
calprob <- CalibrationProbPoissonRisk(riskdata)
```

```
pander::pander(c(h0=calprob$h0,
                  Gain=calprob$hazardGain,
                  DeltaTime=calprob$timeInterval),
                caption="Logistic Calibration Parameters")
```

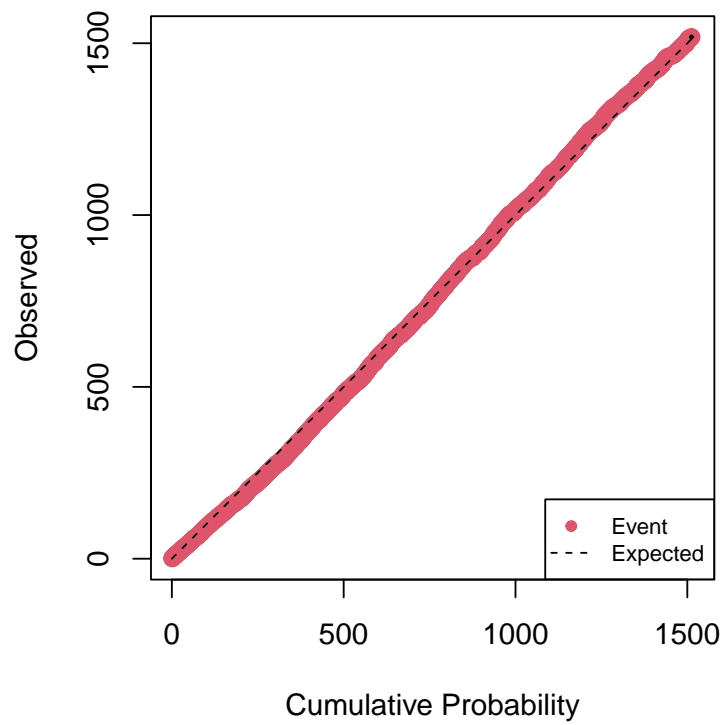
| h0 | Gain | DeltaTime |
|-------|------|-----------|
| 0.676 | 1.31 | 7.14 |

```
timeinterval <- calprob$timeInterval;
gain <- calprob$hazardGain

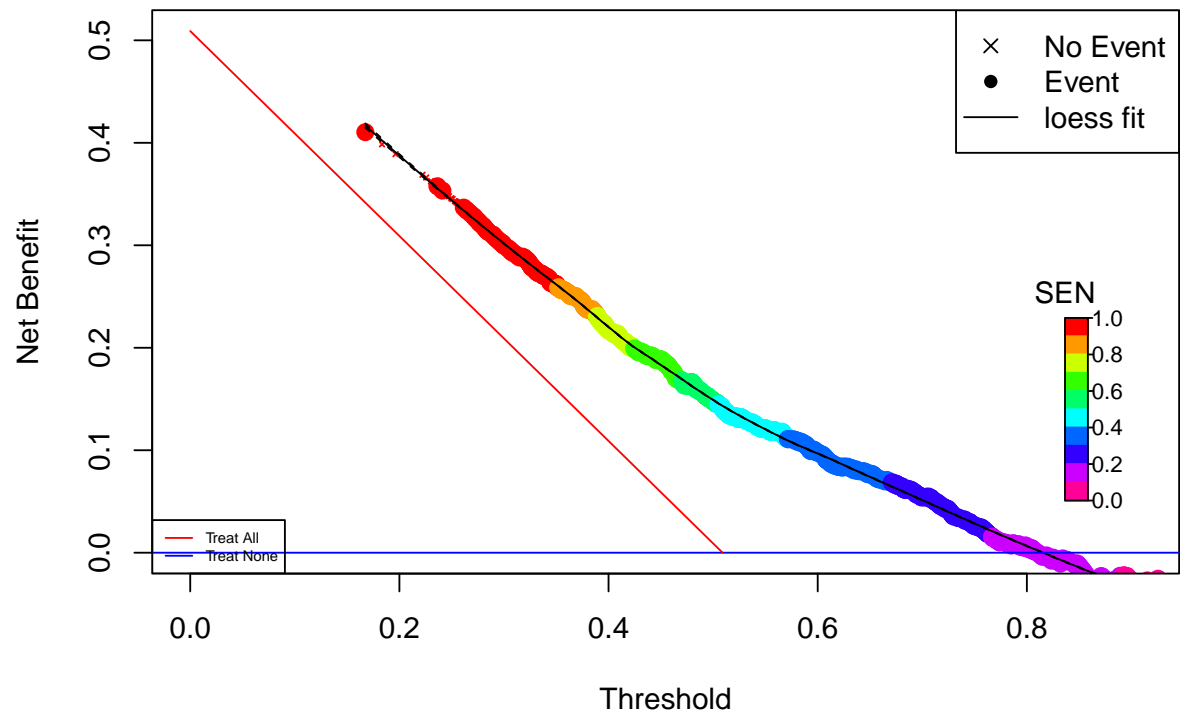
rdata <- cbind(dataBrestCancerTrain$status, calprob$prob)

rrAnalysisTrain <- RRPlot(rdata, atProb=c(0.90, 0.80),
                          timetoEvent=dataBrestCancerTrain$time,
                          title="Cal. Logistic Train: Breast Cancer",
                          ysurvlim=c(0.00, 1.0),
                          riskTimeInterval=timeinterval)
```

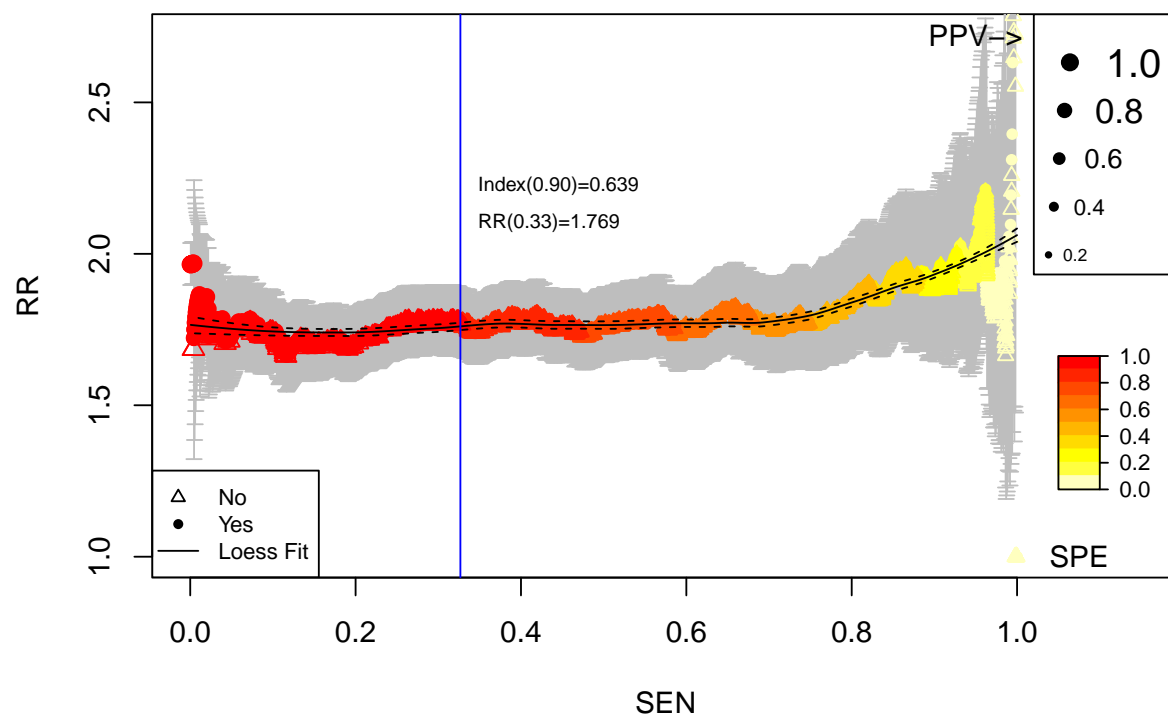
Cumulative vs. Observed: Cal. Logistic Train: Breast Cancer

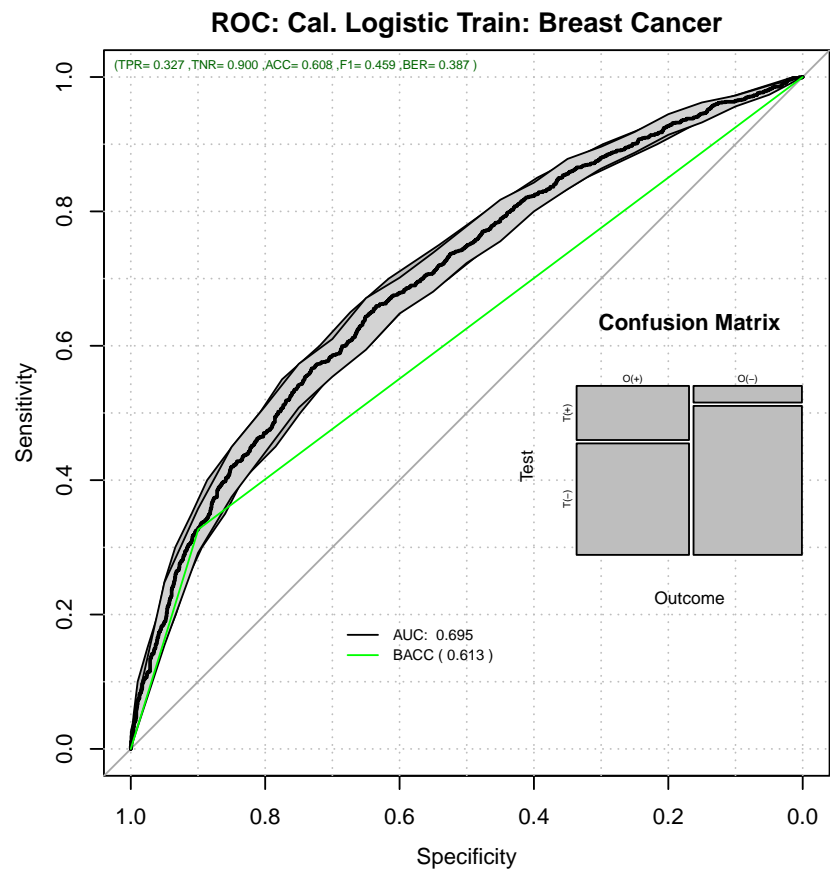


Decision Curve Analysis: Cal. Logistic Train: Breast Cancer

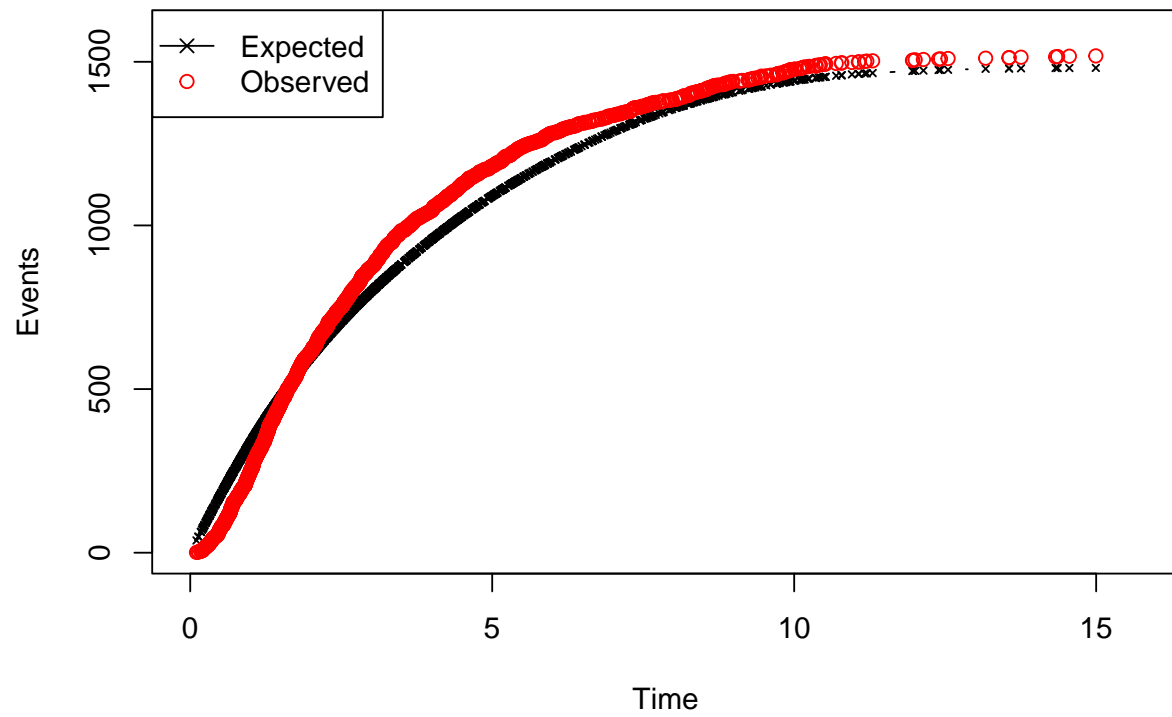


Relative Risk: Cal. Logistic Train: Breast Cancer

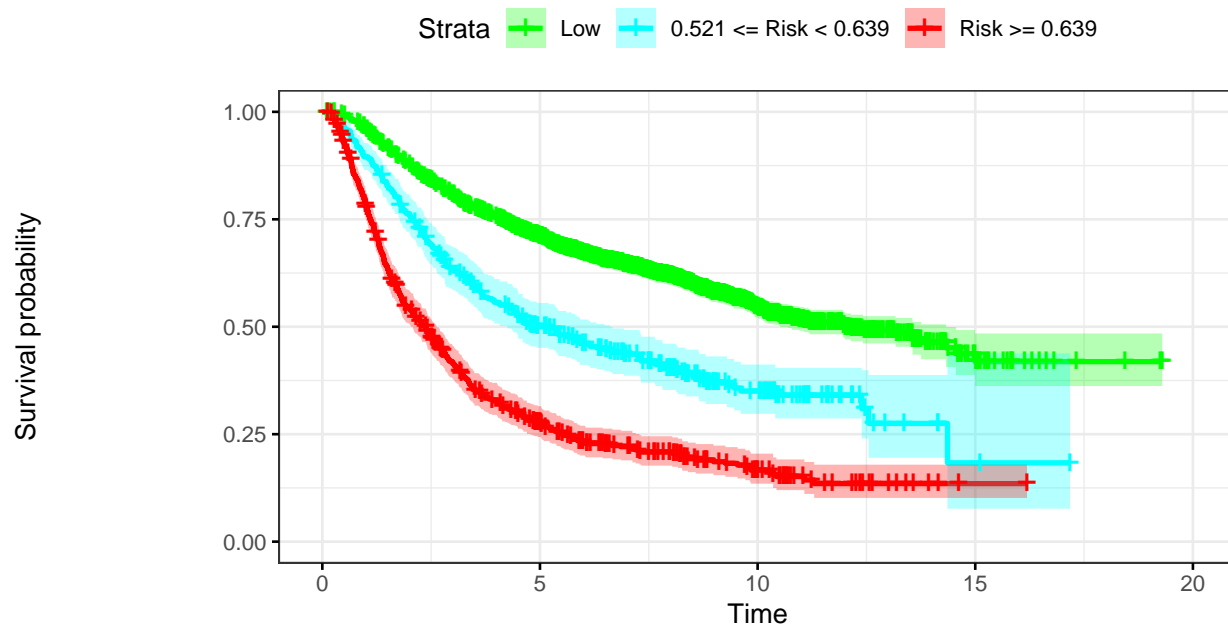




Time vs. Events: Cal. Logistic Train: Breast Cancer



Kaplan–Meier: Cal. Logistic Train: Breast Cancer



Number at risk

| | | | | | |
|-----------------------|------|------|-----|----|---|
| Low | 1975 | 1268 | 399 | 23 | 0 |
| 0.521 <= Risk < 0.639 | 364 | 160 | 47 | 2 | 0 |
| Risk >= 0.639 | 643 | 143 | 37 | 1 | 0 |

```
par(op)
```

1.8.1 Report of the calibrated logistic: training

```
pander::pander(t(rrAnalysisTrain$OERatio),caption="O/E Ratio")
```

Table 65: O/E Ratio

| est | lower | upper |
|------|-------|-------|
| 1.02 | 0.974 | 1.08 |

```
pander::pander(rrAnalysisTrain$c.index,caption="C. Index")
```

- C Index: 0.68
- Dxy: 0.36
- S.D.: 0.014
- n: 2982
- missing: 0
- uncensored: 1518
- Relevant Pairs: 6184528

- **Concordant:** 4206588
- **Uncertain:** 2703838
- **cstatCI:**

| mean.C Index | median | lower | upper |
|--------------|--------|-------|-------|
| 0.68 | 0.68 | 0.667 | 0.693 |

```
pander::pander(t(rrAnalysisTrain$ROCAalysis$aucs),caption="ROC AUC")
```

Table 67: ROC AUC

| est | lower | upper |
|-------|-------|-------|
| 0.695 | 0.677 | 0.714 |

```
pander::pander((rrAnalysisTrain$ROCAalysis$sensitivity),caption="Sensitivity")
```

Table 68: Sensitivity

| est | lower | upper |
|-------|-------|-------|
| 0.327 | 0.303 | 0.351 |

```
pander::pander((rrAnalysisTrain$ROCAalysis$specificity),caption="Specificity")
```

Table 69: Specificity

| est | lower | upper |
|-----|-------|-------|
| 0.9 | 0.883 | 0.915 |

```
pander::pander(t(rrAnalysisTrain$thr_atP),caption="Probability Thresholds")
```

Table 70: Probability Thresholds

| 90% | 80% |
|-------|-------|
| 0.639 | 0.521 |

```
pander::pander(t(rrAnalysisTrain$RR_atP),caption="Risk Ratio")
```

Table 71: Risk Ratio

| est | lower | upper |
|------|-------|-------|
| 1.77 | 1.66 | 1.88 |

```
pander::pander(rrAnalysisTrain$surdif,caption="Logrank test")
```

Table 72: Logrank test Chisq = 543.347175 on 2 degrees of freedom,
p = 0.000000

| | N | Observed | Expected | (O-E)^2/E | (O-E)^2/V |
|---------|------|----------|----------|-----------|-----------|
| class=0 | 1975 | 804 | 1145 | 101.5 | 418.9 |
| class=1 | 364 | 218 | 169 | 14.1 | 15.9 |
| class=2 | 643 | 496 | 204 | 418.2 | 490.7 |

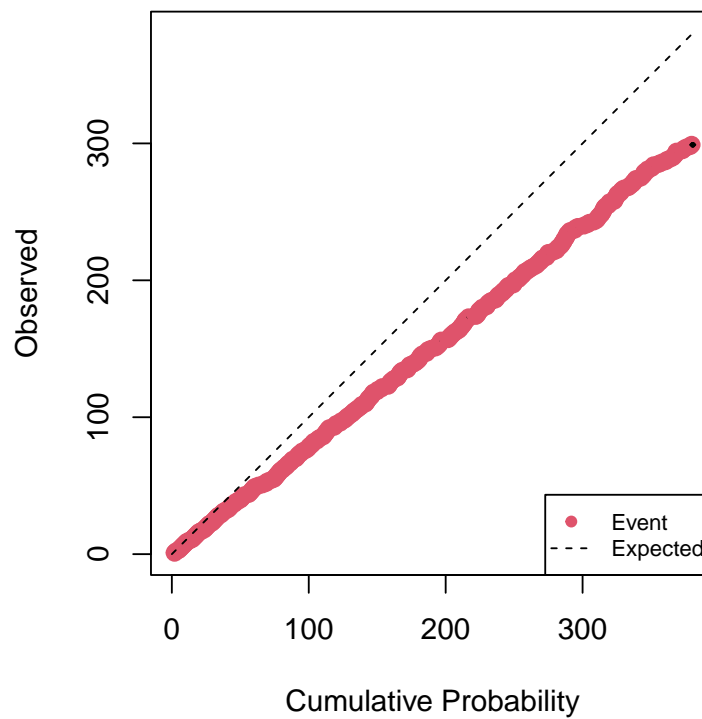
```

probLog <- predict(mlog,dataBreStCancerTest)
aprob <- adjustProb(probLog,gain)

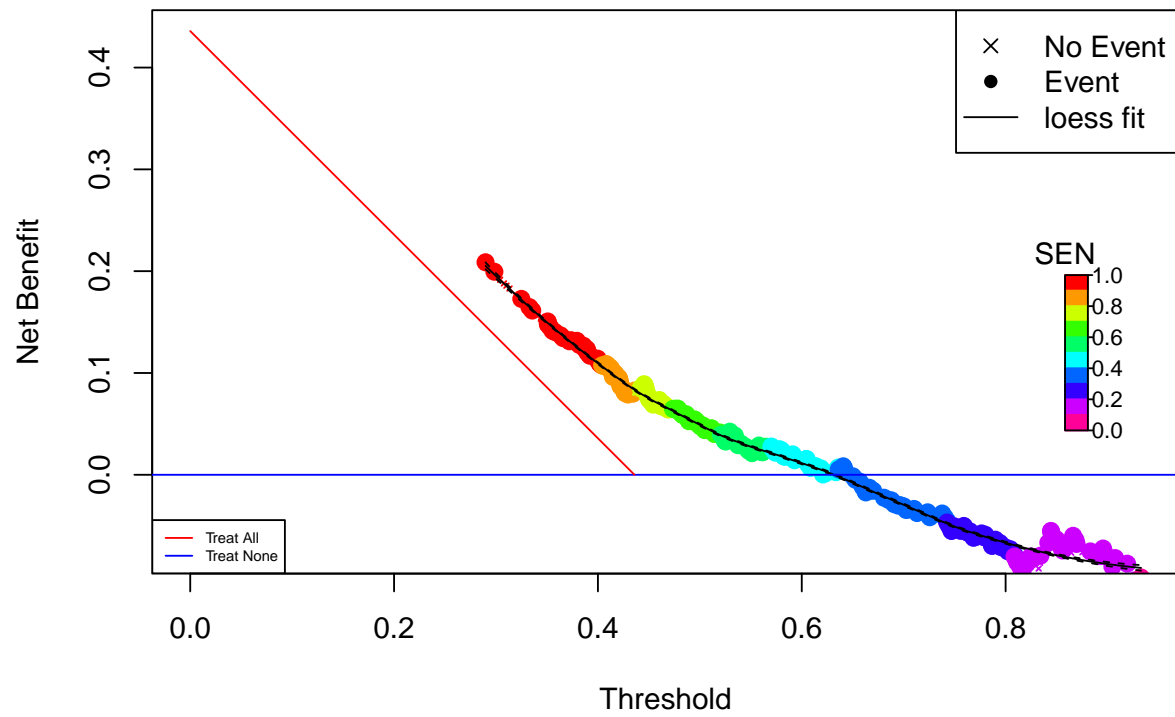
rdata <- cbind(dataBreStCancerTest$status,aprob)
rrAnalysisTestLogistic <- RRPlot(rdata,atThr=rrAnalysisTrain$thr_atP,
                                timetoEvent=dataBreStCancerTest$time,
                                title="Cal. Logistic Test: Breast Cancer",
                                ysurvlim=c(0.00,1.0),
                                riskTimeInterval=timeinterval)

```

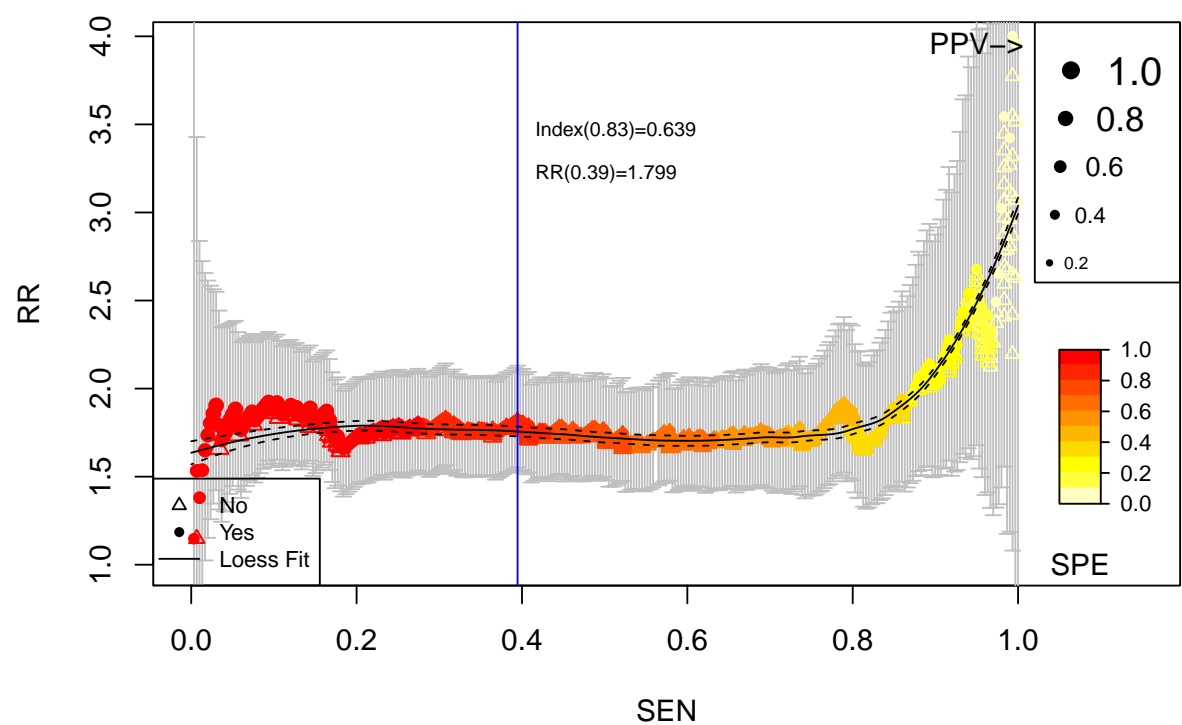
Cumulative vs. Observed: Cal. Logistic Test: Breast Cancer

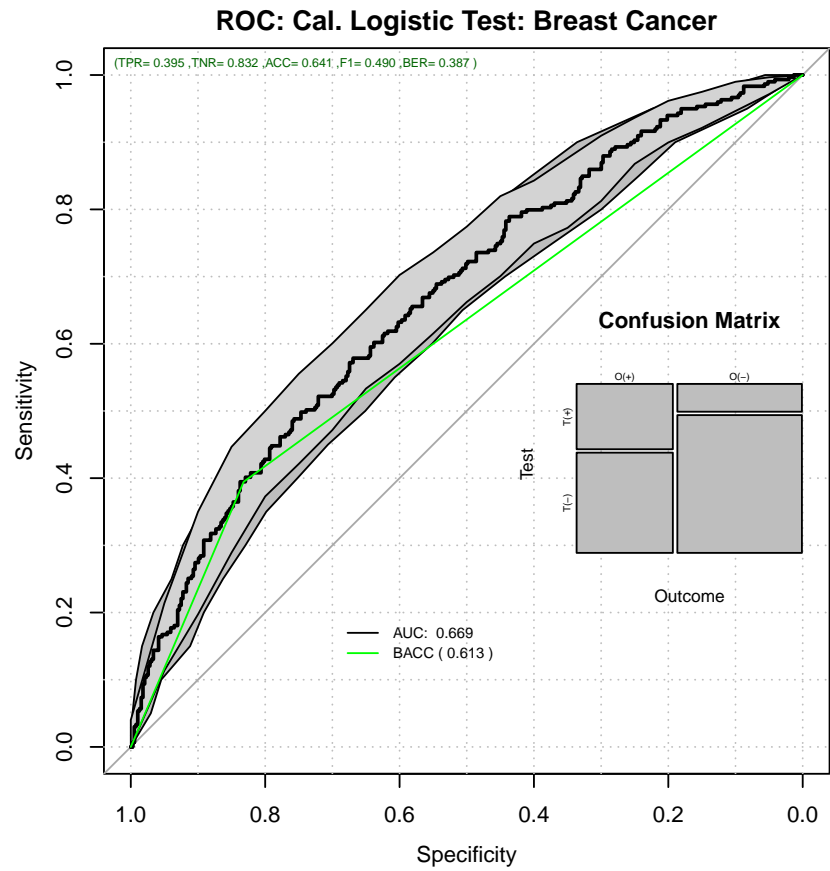


Decision Curve Analysis: Cal. Logistic Test: Breast Cancer

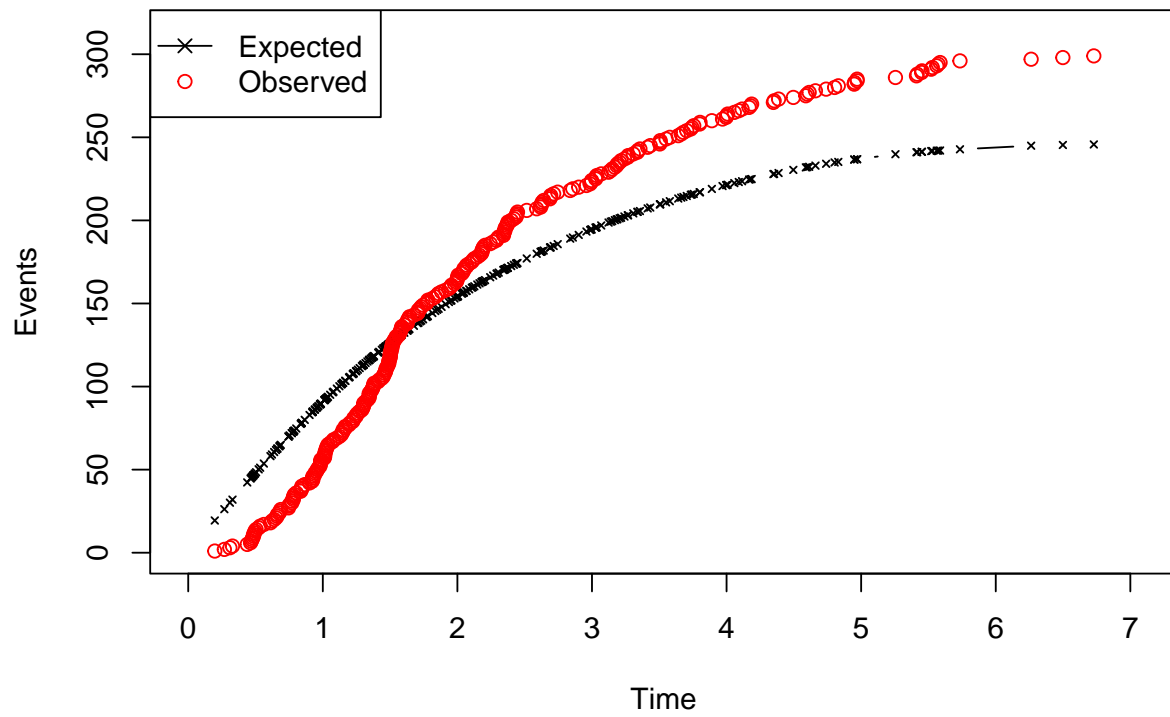


Relative Risk: Cal. Logistic Test: Breast Cancer

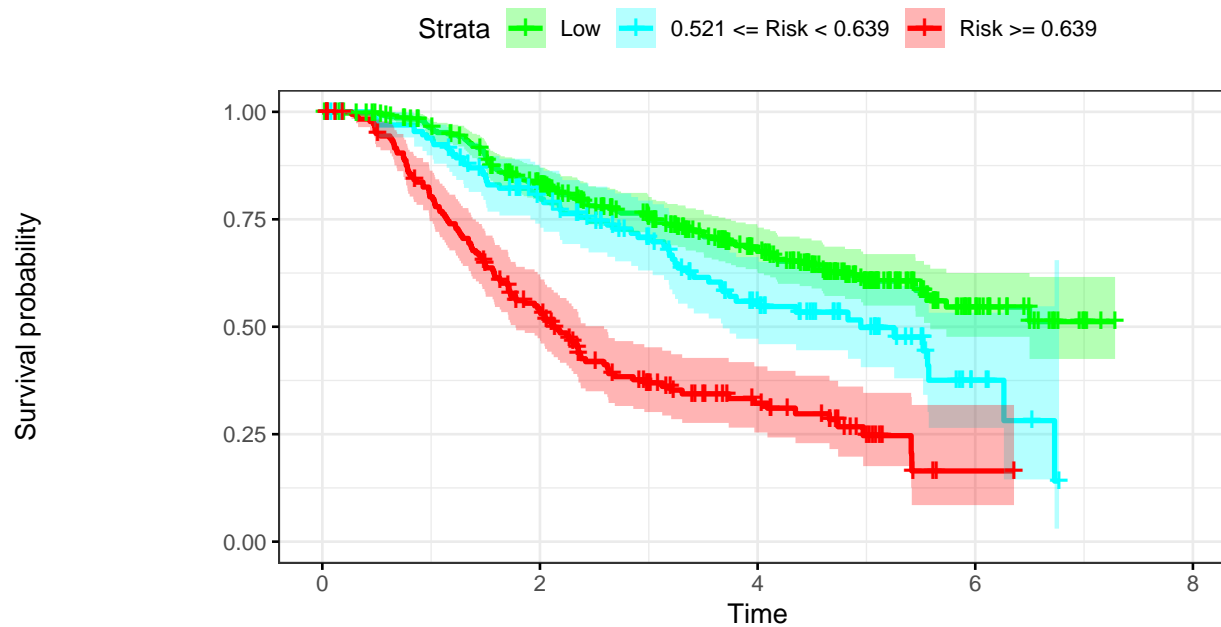




Time vs. Events: Cal. Logistic Test: Breast Cancer



Kaplan–Meier: Cal. Logistic Test: Breast Cancer



Number at risk

| | | | | | |
|-----------------------|-----|-----|-----|----|---|
| Low | 369 | 274 | 154 | 29 | 0 |
| 0.521 <= Risk < 0.639 | 134 | 96 | 46 | 6 | 0 |
| Risk >= 0.639 | 183 | 89 | 29 | 1 | 0 |

```
par(op)
```

1.8.2 Report of the calibrated validation

```
pander::pander(t(rrAnalysisTestLogistic$OERatio),caption="O/E Ratio")
```

Table 73: O/E Ratio

| est | lower | upper |
|------|-------|-------|
| 1.22 | 1.08 | 1.36 |

```
pander::pander(rrAnalysisTestLogistic$c.index,caption="C. Index")
```

- C Index: 0.669
- Dxy: 0.338
- S.D.: 0.0309
- n: 686
- missing: 0
- uncensored: 299
- Relevant Pairs: 266144

- **Concordant:** 178115
- **Uncertain:** 203702
- **cstatCI:**

| mean.C Index | median | lower | upper |
|--------------|--------|-------|-------|
| 0.669 | 0.67 | 0.64 | 0.698 |

```
pander::pander(t(rrAnalysisTestLogistic$ROCAalysis$aucs),caption="ROC AUC")
```

Table 75: ROC AUC

| est | lower | upper |
|-------|-------|-------|
| 0.669 | 0.628 | 0.709 |

```
pander::pander((rrAnalysisTestLogistic$ROCAalysis$sensitivity),caption="Sensitivity")
```

Table 76: Sensitivity

| est | lower | upper |
|-------|-------|-------|
| 0.395 | 0.339 | 0.453 |

```
pander::pander((rrAnalysisTestLogistic$ROCAalysis$specificity),caption="Specificity")
```

Table 77: Specificity

| est | lower | upper |
|-------|-------|-------|
| 0.832 | 0.791 | 0.868 |

```
pander::pander(t(rrAnalysisTestLogistic$thr_atP),caption="Probability Thresholds")
```

Table 78: Probability Thresholds

| 90% | 80% |
|-------|-------|
| 0.639 | 0.521 |

```
pander::pander(t(rrAnalysisTestLogistic$RR_atP),caption="Risk Ratio")
```

Table 79: Risk Ratio

| est | lower | upper |
|-----|-------|-------|
| 1.8 | 1.54 | 2.11 |

```
pander::pander(rrAnalysisTestLogistic$surdif,caption="Logrank test")
```

Table 80: Logrank test Chisq = 92.507991 on 2 degrees of freedom,
p = 0.000000

| | N | Observed | Expected | (O-E)^2/E | (O-E)^2/V |
|----------------|-----|----------|----------|-----------|-----------|
| class=0 | 369 | 121 | 181.7 | 20.2997 | 52.3868 |
| class=1 | 134 | 60 | 61.7 | 0.0479 | 0.0604 |
| class=2 | 183 | 118 | 55.5 | 70.2342 | 88.0195 |

1.9 Comparing the COX and Logistic Models on the Independent Data

```
pander::pander(t(rrCoxTestAnalysis$OAcum95ci))
```

| mean | 50% | 2.5% | 97.5% |
|------|------|-------|-------|
| 0.84 | 0.84 | 0.839 | 0.841 |

```
pander::pander(t(rrAnalysisTestLogistic$OAcum95ci))
```

| mean | 50% | 2.5% | 97.5% |
|-------|-------|-------|-------|
| 0.791 | 0.791 | 0.791 | 0.792 |

```
pander::pander(t(rrCoxTestAnalysis$OE95ci))
```

| mean | 50% | 2.5% | 97.5% |
|------|------|------|-------|
| 1.06 | 1.06 | 1.03 | 1.09 |

```
pander::pander(t(rrAnalysisTestLogistic$OE95ci))
```

| mean | 50% | 2.5% | 97.5% |
|-------|-------|-------|-------|
| 0.955 | 0.954 | 0.925 | 0.985 |

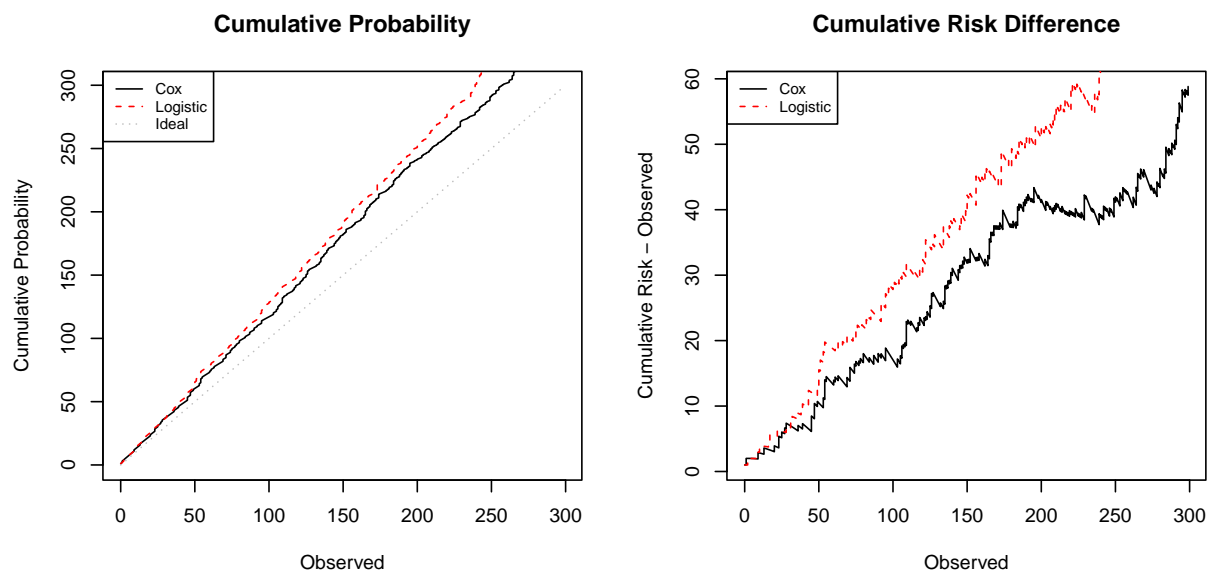
```
maxobs <- sum(dataBrestCancerTest$status)

par(mfrow=c(1,2),cex=0.75)

plot(rrCoxTestAnalysis$CumulativeOvs,type="l",lty=1,
     main="Cumulative Probability",
     xlab="Observed",
     ylab="Cumulative Probability",
     ylim=c(0,maxobs),
     xlim=c(0,maxobs))
lines(rrAnalysisTestLogistic$CumulativeOvs,lty=2,col="red")
lines(x=c(0,maxobs),y=c(0,maxobs),lty=3,col="gray")
legend("topleft",legend = c("Cox","Logistic","Ideal"),
     col=c("black","red","gray"),
     lty=c(1,2,3),
     cex=0.75)
```

```
)

plot(rrCoxTestAnalysis$CumulativeOvs$Observed,
     rrCoxTestAnalysis$CumulativeOvs$Cumulative-
     rrCoxTestAnalysis$CumulativeOvs$Observed,
     main="Cumulative Risk Difference",
     xlab="Observed",
     ylab="Cumulative Risk - Observed",
     type="l",
     lty=1)
lines(rrAnalysisTestLogistic$CumulativeOvs$Observed,
      rrAnalysisTestLogistic$CumulativeOvs$Cumulative-
      rrAnalysisTestLogistic$CumulativeOvs$Observed,
      lty=2,
      col="red")
legend("topleft", legend = c("Cox", "Logistic"),
      col=c("black", "red"),
      lty=c(1, 2),
      cex=0.75
)
```



```
plot(rrCoxTestAnalysis$OEData[,2:3], type="l", lty=1,
     main="Expected over Time",
     xlab="Observed",
     ylab="Expected",
     ylim=c(0, maxobs),
     xlim=c(0, maxobs))
lines(rrAnalysisTestLogistic$OEData[,2:3], lty=2, col="red")
lines(x=c(0, maxobs), y=c(0, maxobs), lty=3, col="gray")
legend("topleft", legend = c("Cox", "Logistic", "Ideal"),
      col=c("black", "red", "gray"),
      lty=c(1, 2, 3),
```

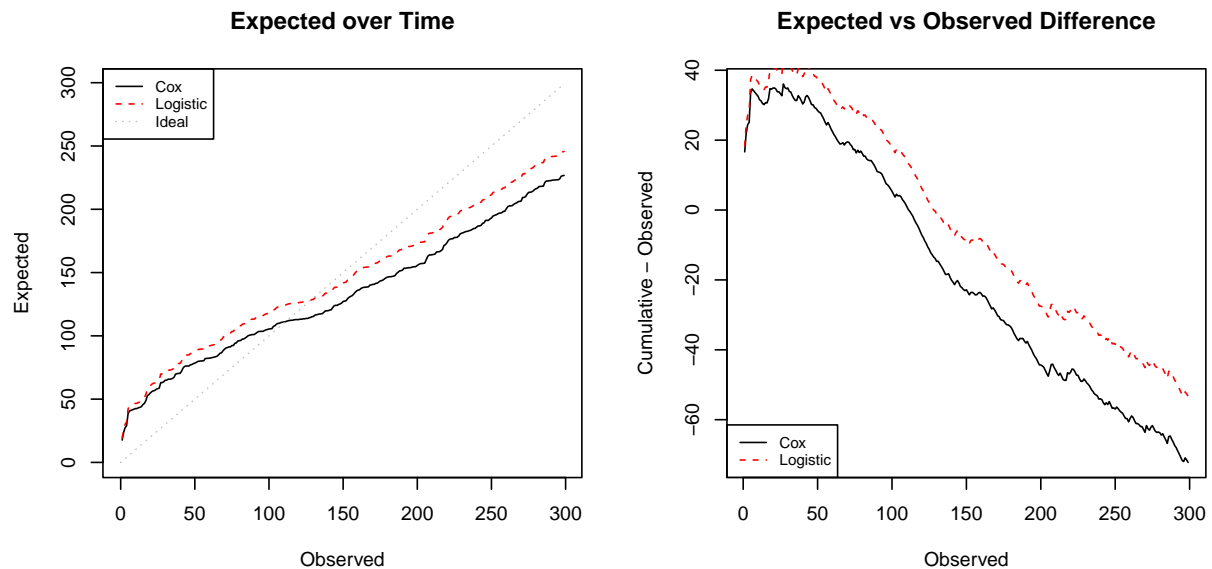
```

    cex=0.75
)

plot(rrCoxTestAnalysis$OEData$Observed,
     rrCoxTestAnalysis$OEData$Expected-
     rrCoxTestAnalysis$OEData$Observed,
     main="Expected vs Observed Difference",
     xlab="Observed",
     ylab="Cumulative - Observed",
     type="l",
     lty=1)
lines(rrAnalysisTestLogistic$OEData$Observed,
      rrAnalysisTestLogistic$OEData$Expected-
      rrAnalysisTestLogistic$OEData$Observed,
      lty=2,col="red")

legend("bottomleft",legend = c("Cox","Logistic"),
      col=c("black","red"),
      lty=c(1,2),
      cex=0.75
)

```



```

par(op)

```