

Colon Cancer

Jose Tamez

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```
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
#library(corrplot)
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)

data(cancer)
colon <- subset(colon, etype==1)
colon$etype <- NULL
rownames(colon) <- colon$id
colon$id <- NULL
```

```

colon <- colon[complete.cases(colon),]
time <- colon$time
status <- colon$status
data <- colon
data$time <- NULL
data$study <- NULL
table(data$status)

```

```
0 1 442 446
```

```

#dataColon <- as.data.frame(model.matrix(status~.*.,data))
dataColon <- as.data.frame(model.matrix(status~.*age,data))
dataColon$`(Intercept)` <- NULL
dataColon$time <- time/365
dataColon$status <- status
colnames(dataColon) <-str_replace_all(colnames(dataColon),":","_")
colnames(dataColon) <-str_replace_all(colnames(dataColon),"\\.","_")
colnames(dataColon) <-str_replace_all(colnames(dataColon),"\\+", "_")
data <- NULL

trainsamples <- sample(nrow(dataColon),0.7*nrow(dataColon))
dataColonTrain <- dataColon[trainsamples,]
dataColonTest <- dataColon[-trainsamples,]

```

```
pander::pander(table(dataColonTrain$status))
```

0	1
320	301

```
pander::pander(table(dataColonTest$status))
```

0	1
122	145

0.1 Modeling

```
ml <- BSWiMS.model(Surv(time,status)~1,data=dataColonTrain,NumberOfRepeats = 10)
```

```
[+++++++++++++++++++++++++++++++++++++++++++++++++++++++++++++++++++++]....
```

```

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

```

Table 3: Table continues below

	Estimate	lower	HR	upper	u.Accuracy
nodes	0.018702	1.013	1.019	1.025	0.602
node4	0.205428	1.142	1.228	1.321	0.612
age_node4	0.005120	1.003	1.005	1.007	0.612
age_nodes	0.000160	1.000	1.000	1.000	0.593

	Estimate	lower	HR	upper	u.Accuracy
age	-0.016078	0.978	0.984	0.991	0.517
age_extent	0.003637	1.002	1.004	1.006	0.517
extent	0.130698	1.062	1.140	1.223	0.543
rxLev_5FU_age	-0.001709	0.997	0.998	0.999	0.552
rxLev_5FU	-0.142480	0.782	0.867	0.962	0.552
surg	0.089631	1.016	1.094	1.177	0.546
age_surg	0.000302	1.000	1.000	1.001	0.546

Table 4: Table continues below

	r.Accuracy	full.Accuracy	u.AUC	r.AUC	full.AUC
nodes	0.512	0.607	0.597	0.518	0.603
node4	0.583	0.620	0.604	0.586	0.613
age_node4	0.590	0.628	0.604	0.594	0.627
age_nodes	0.498	0.594	0.588	0.508	0.589
age	0.619	0.628	0.515	0.614	0.627
age_extent	0.615	0.628	0.518	0.611	0.627
extent	0.612	0.620	0.554	0.604	0.613
rxLev_5FU_age	0.620	0.620	0.558	0.613	0.613
rxLev_5FU	0.632	0.628	0.558	0.629	0.627
surg	0.605	0.628	0.539	0.602	0.627
age_surg	0.603	0.628	0.539	0.601	0.626

	IDI	NRI	z.IDI	z.NRI	Delta.AUC	Frequency
nodes	0.02913	0.390	5.87	5.21	0.085718	1.0
node4	0.03800	0.418	5.51	6.12	0.027357	1.0
age_node4	0.03330	0.418	5.14	6.12	0.033102	1.0
age_nodes	0.02376	0.350	5.03	4.69	0.080486	1.0
age	0.02789	0.220	4.77	2.76	0.012935	1.0
age_extent	0.01884	0.215	3.86	3.92	0.015303	1.0
extent	0.01718	0.215	3.63	3.92	0.008799	1.0
rxLev_5FU_age	0.01209	0.231	3.18	3.10	-0.000197	1.0
rxLev_5FU	0.00879	0.231	2.68	3.10	-0.002800	1.0
surg	0.00576	0.156	2.37	2.18	0.024751	1.0
age_surg	0.00513	0.156	2.23	2.18	0.025497	0.5

0.2 Cox Model Performance

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

0.2.1 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

```
index <- predict(ml,dataColonTrain)
timeinterval <- 2*mean(subset(dataColonTrain,status==1)$time)

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

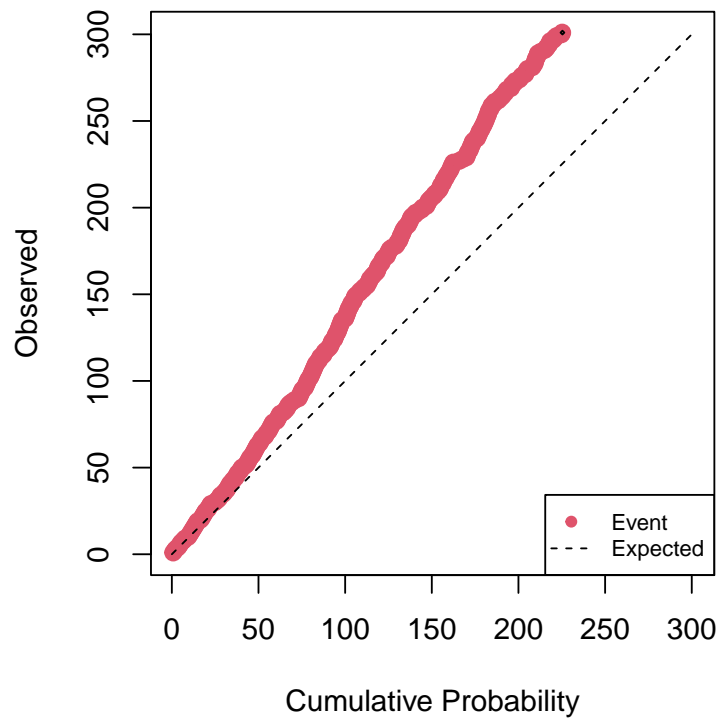
```

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

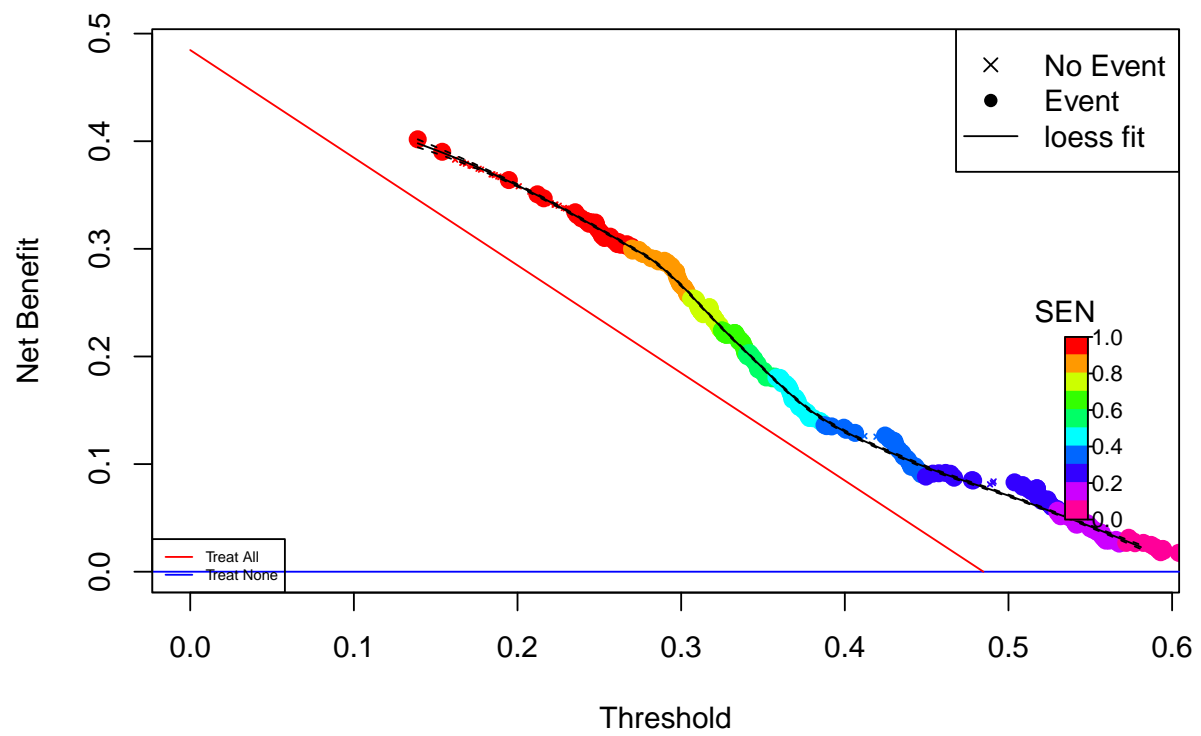
rrAnalysisTrain <- RRPlot(rdata, atProb=c(0.90),
  timetoEvent=dataColonTrain$time,
  title="Raw Train: Colon Cancer",
  ysurvlim=c(0.00, 1.0),
  riskTimeInterval=timeinterval)

```

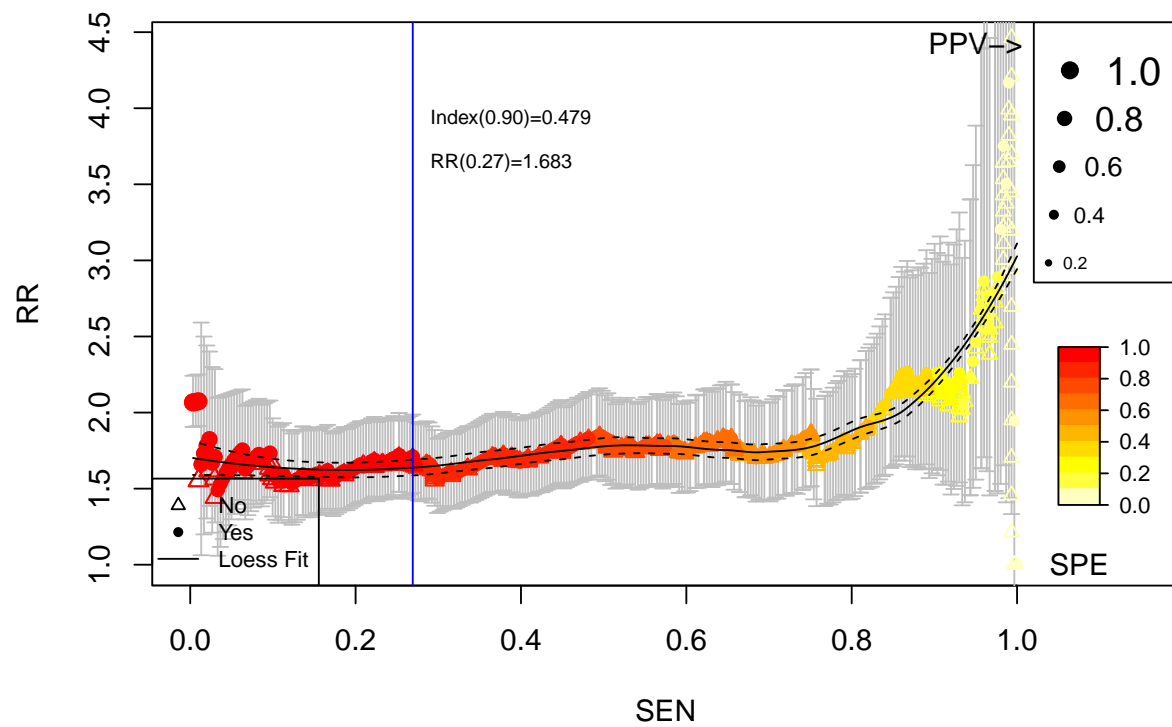
Cumulative vs. Observed: Raw Train: Colon Cancer

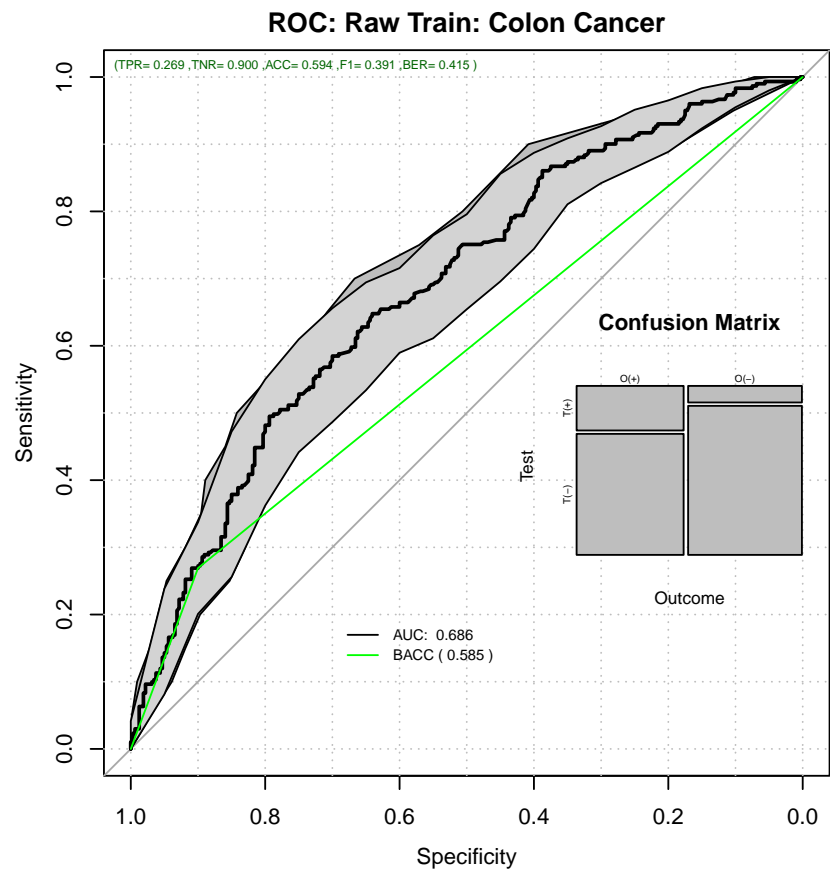


Decision Curve Analysis: Raw Train: Colon Cancer

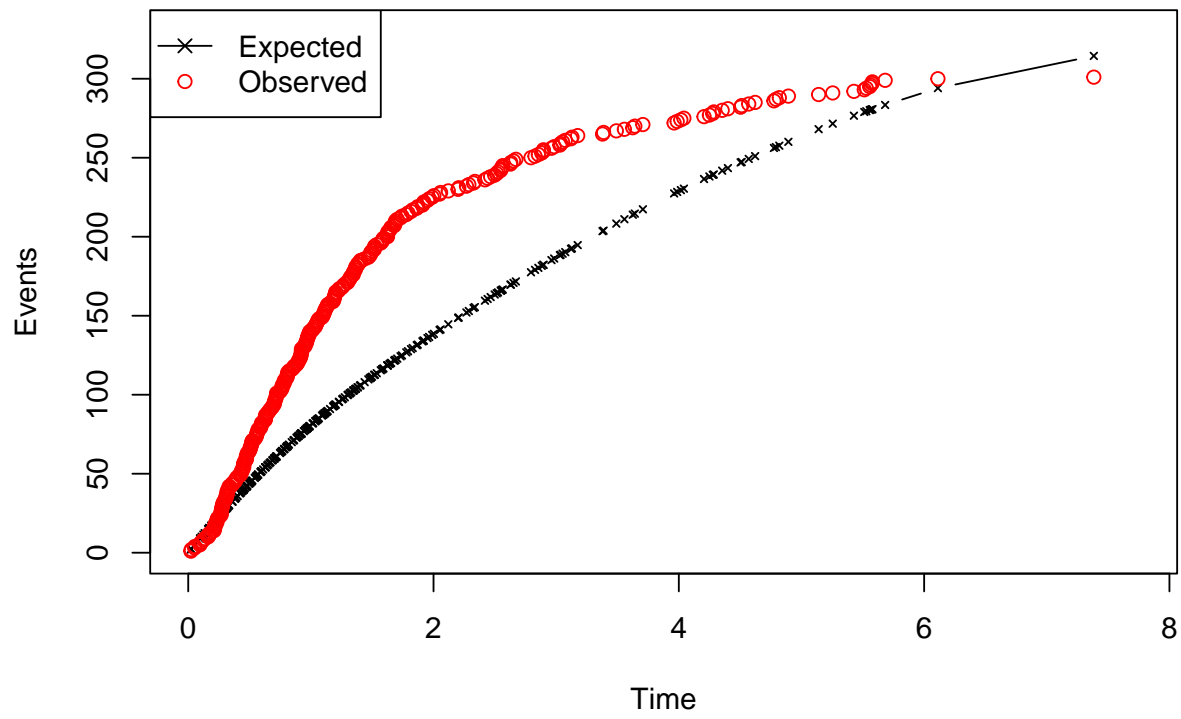


Relative Risk: Raw Train: Colon Cancer

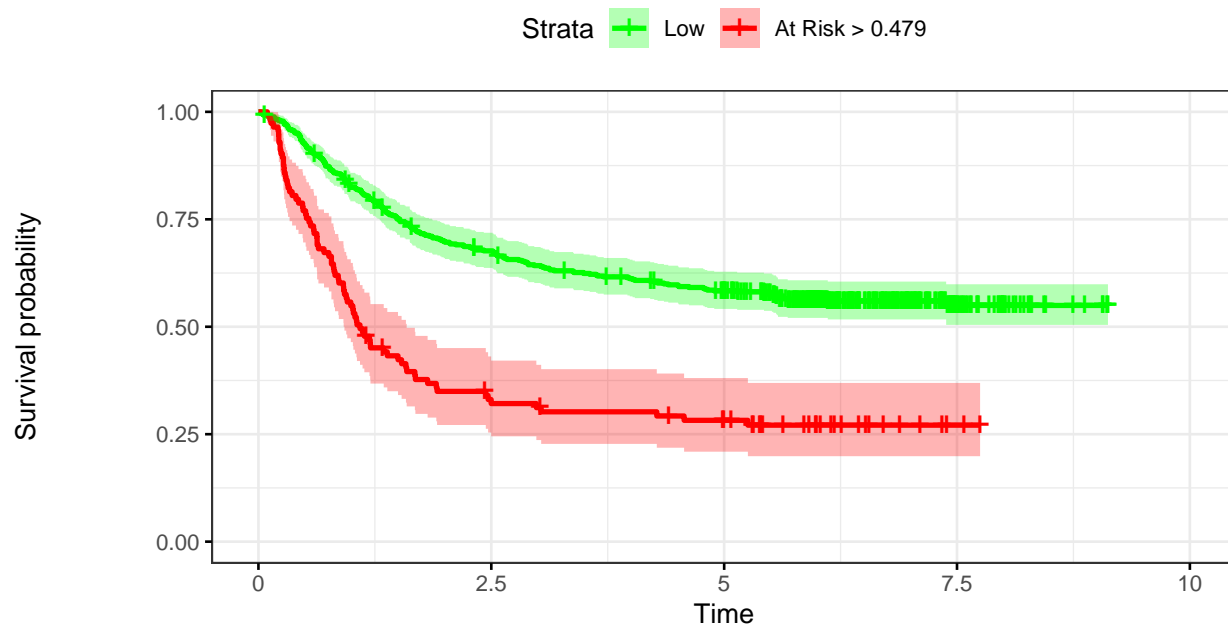




Time vs. Events: Raw Train: Colon Cancer



Kaplan–Meier: Raw Train: Colon Cancer



Number at risk

Low	508	336	279	43	0
At Risk > 0.479	113	34	26	2	0

0.2.2 Uncalibrated Performance Report

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

Table 6: O/E Ratio

est	lower	upper
0.957	0.852	1.07

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

Table 7: O/E Ratio

mean	50%	2.5%	97.5%
1.5	1.5	1.47	1.53

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

Table 8: 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
0.66	0.66	0.629	0.69

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

Table 10: ROC AUC

est	lower	upper
0.686	0.644	0.727

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

Table 11: Sensitivity

est	lower	upper
0.269	0.22	0.323

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

Table 12: Specificity

est	lower	upper
0.9	0.862	0.931

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

Table 13: Probability Thresholds

90%
0.479

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

Table 14: Risk Ratio

est	lower	upper
1.68	1.45	1.96

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

Table 15: Logrank test Chisq = 56.555705 on 1 degrees of freedom,
p = 0.000000

	N	Observed	Expected	(O-E)^2/E	(O-E)^2/V
class=0	508	220	263.1	7.06	56.6
class=1	113	81	37.9	49.00	56.6

0.2.3 Cox Calibration

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

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

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

h0	Gain	DeltaTime
0.646	1.49	2.98

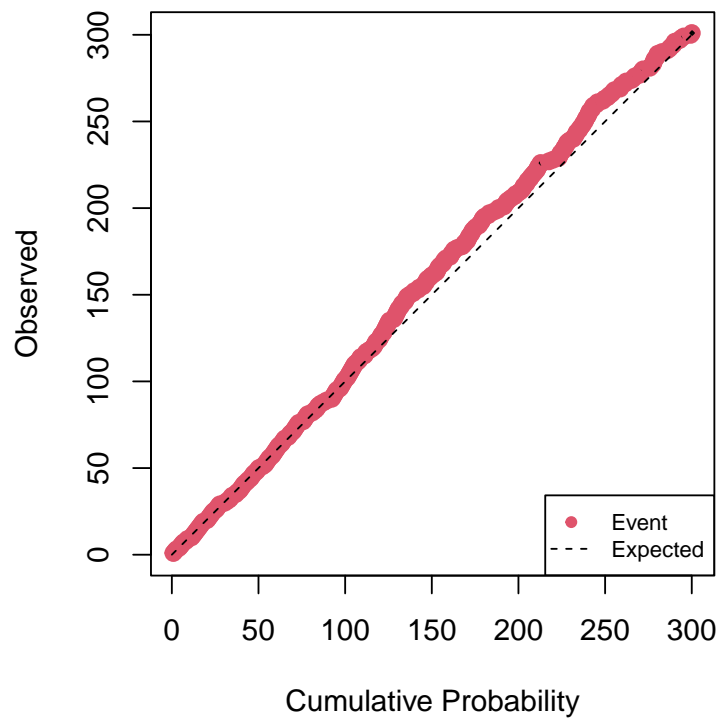
0.2.4 The RRplot() of the calibrated model

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

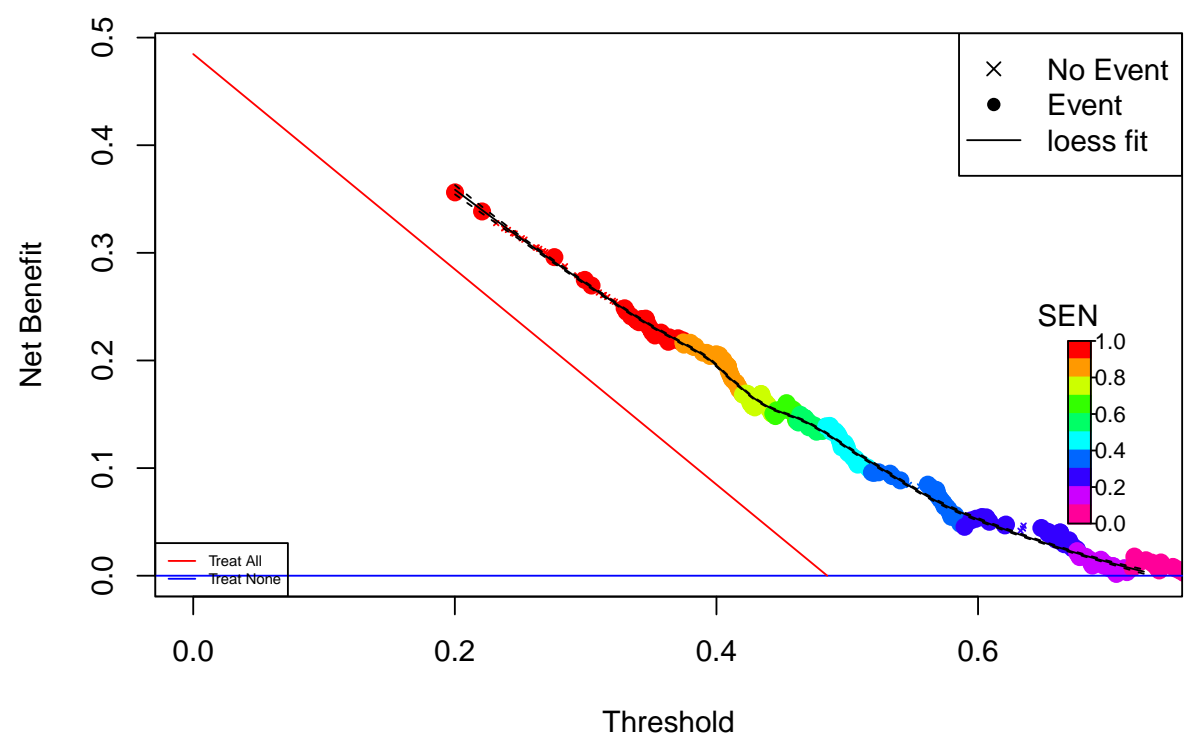
rdata <- cbind(dataColonTrain$status, calprob$prob)

rrAnalysisTrain <- RRPlot(rdata, atProb=c(0.90),
  timetoEvent=dataColonTrain$time,
  title="Calibrated Train: Colon",
  ysurvlim=c(0.00, 1.0),
  riskTimeInterval=timeinterval)
```

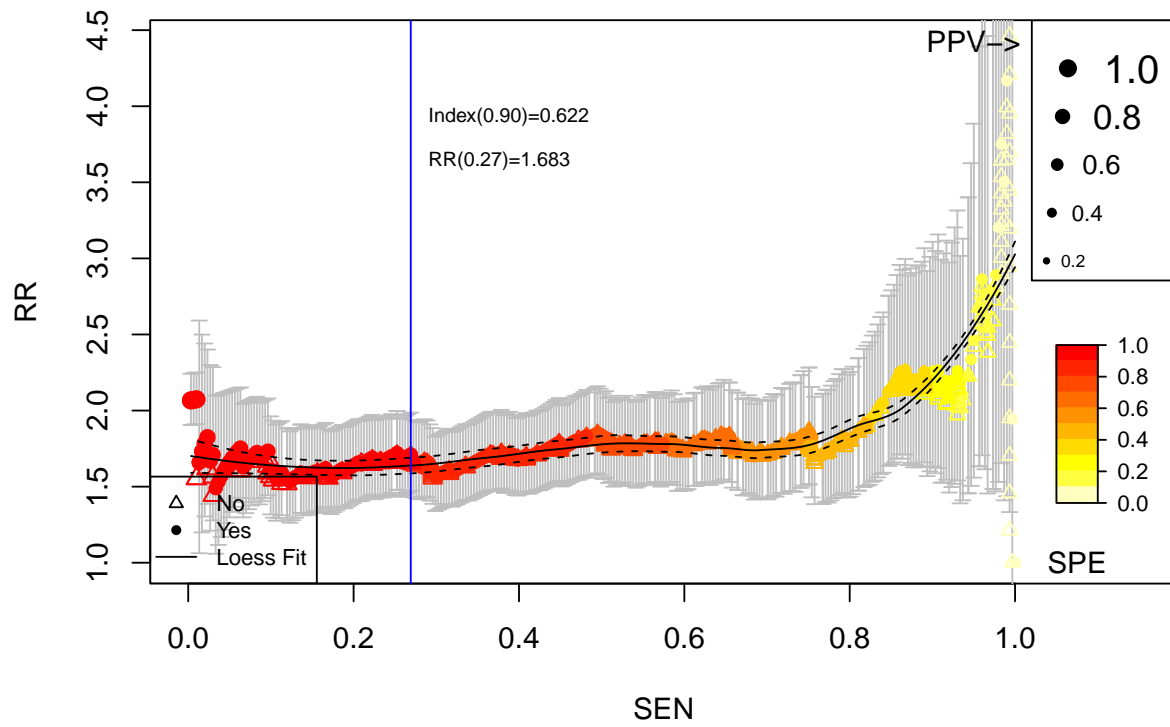
Cumulative vs. Observed: Calibrated Train: Colon

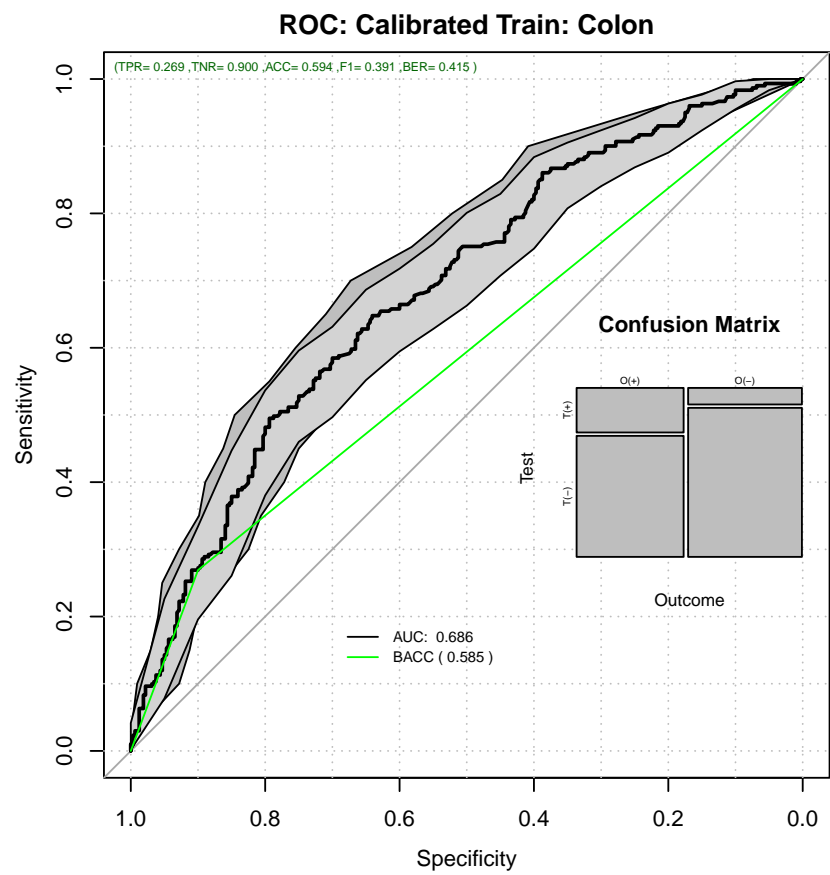


Decision Curve Analysis: Calibrated Train: Colon

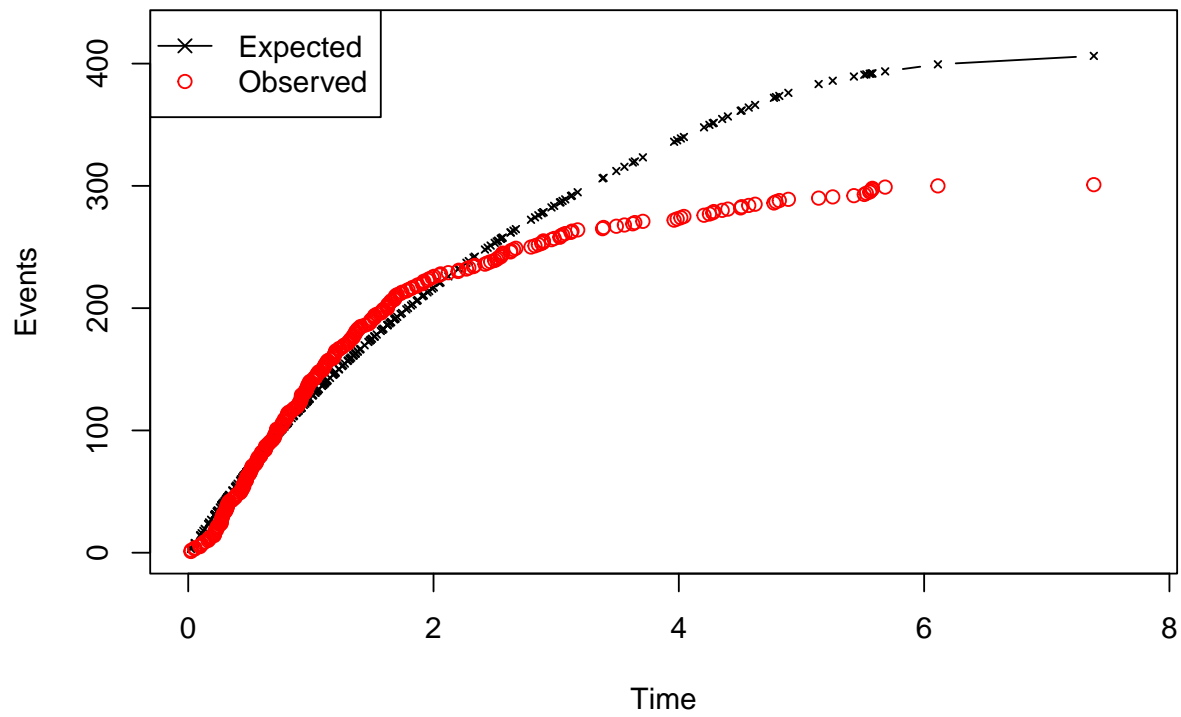


Relative Risk: Calibrated Train: Colon

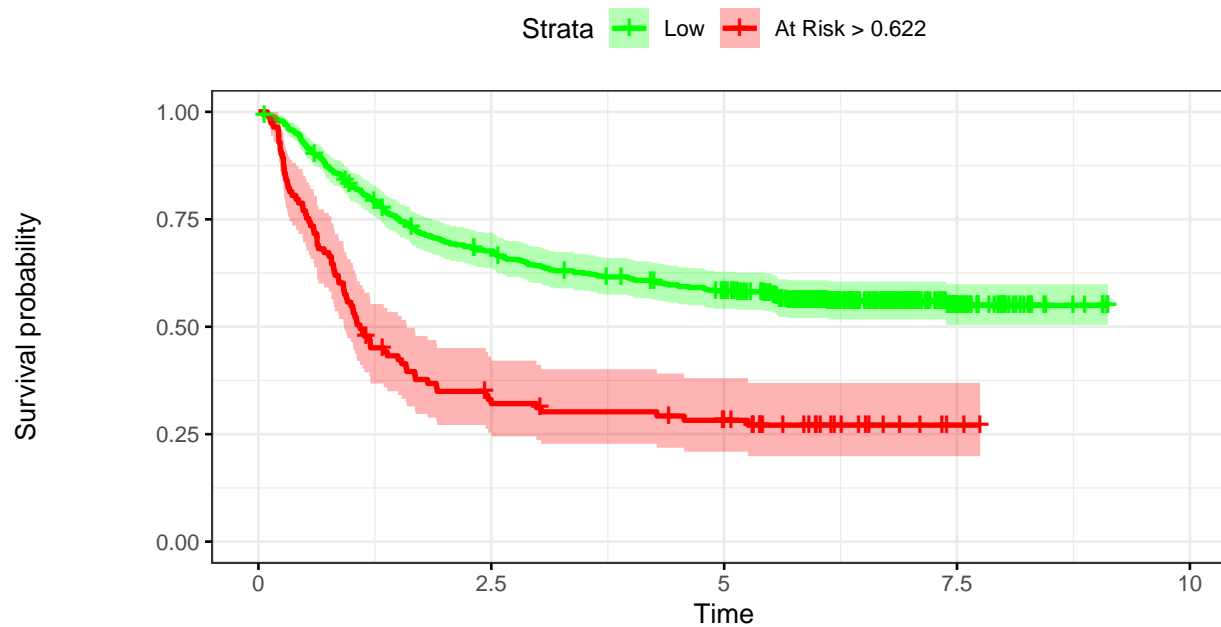




Time vs. Events: Calibrated Train: Colon



Kaplan–Meier: Calibrated Train: Colon



Number at risk

Low	508	336	279	43	0
At Risk > 0.622	113	34	26	2	0

0.2.5 Calibrated Train Performance

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

Table 17: O/E Ratio

est	lower	upper
0.741	0.659	0.829

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

Table 18: O/E Ratio

mean	50%	2.5%	97.5%
0.964	0.964	0.948	0.981

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

Table 19: O/Acum Ratio

mean	50%	2.5%	97.5%
1.04	1.04	1.03	1.04

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

mean.C Index	median	lower	upper
0.66	0.659	0.629	0.688

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

Table 21: ROC AUC

est	lower	upper
0.686	0.644	0.727

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

Table 22: Sensitivity

est	lower	upper
0.269	0.22	0.323

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

Table 23: Specificity

est	lower	upper
0.9	0.862	0.931

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

Table 24: Probability Thresholds

90%
0.622

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

Table 25: Risk Ratio

est	lower	upper
1.68	1.45	1.96

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

Table 26: Logrank test Chisq = 56.555705 on 1 degrees of freedom,
p = 0.000000

	N	Observed	Expected	(O-E)^2/E	(O-E)^2/V
class=0	508	220	263.1	7.06	56.6
class=1	113	81	37.9	49.00	56.6

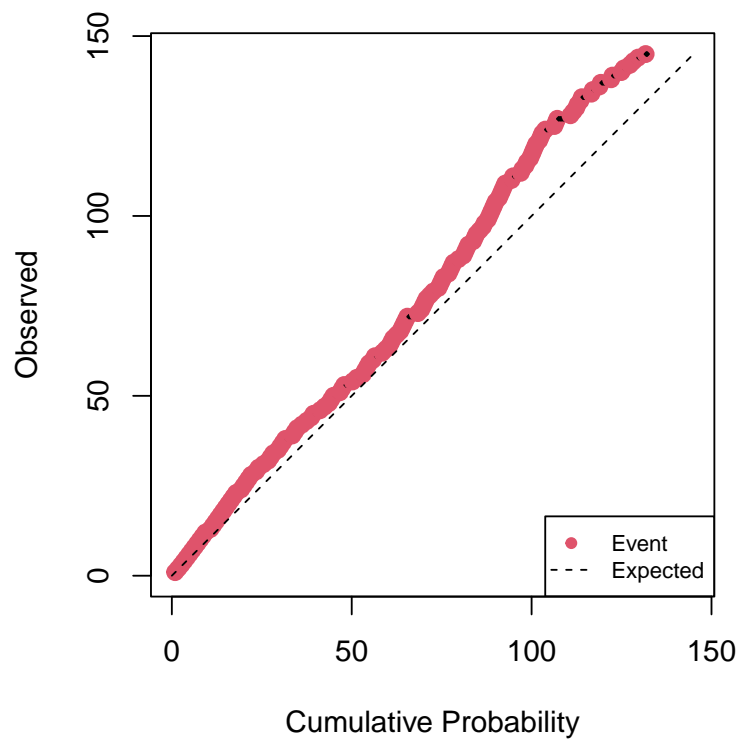
0.2.6 Evaluating on the test set

The calibrated h0 and timeinterval were estimated on the training set

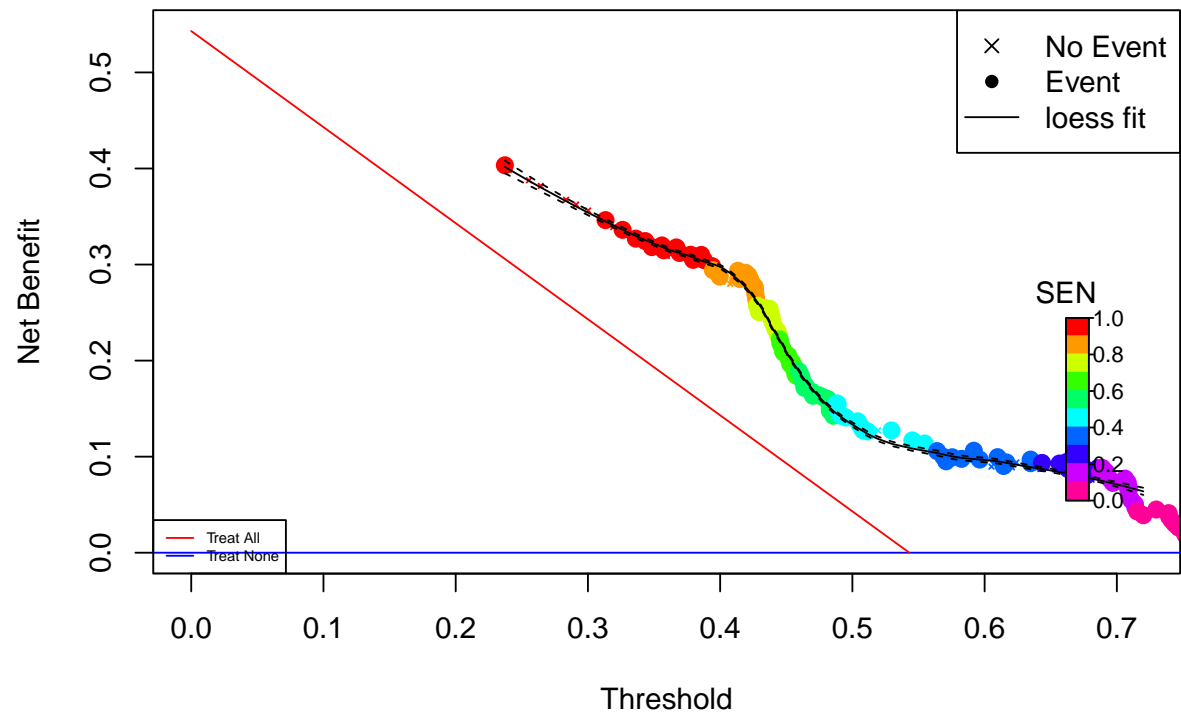
```
index <- predict(ml,dataColonTest)
rdata <- cbind(dataColonTest$status,ppoisGzero(index,h0))

rrAnalysisTest <- RRPlot(rdata,atThr = rrAnalysisTrain$thr_atP,
  timetoEvent=dataColonTest$time,
  title="Test: Colon Cancer",
  ysurvlim=c(0.00,1.0),
  riskTimeInterval=timeinterval)
```

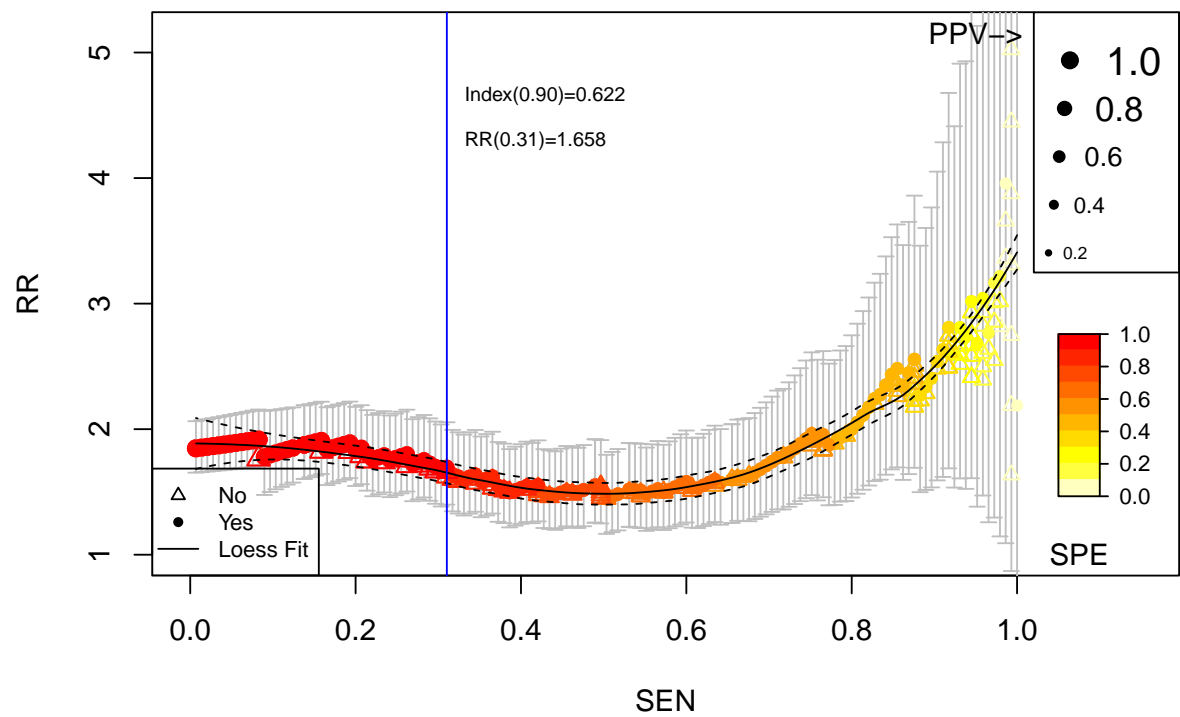
Cumulative vs. Observed: Test: Colon Cancer

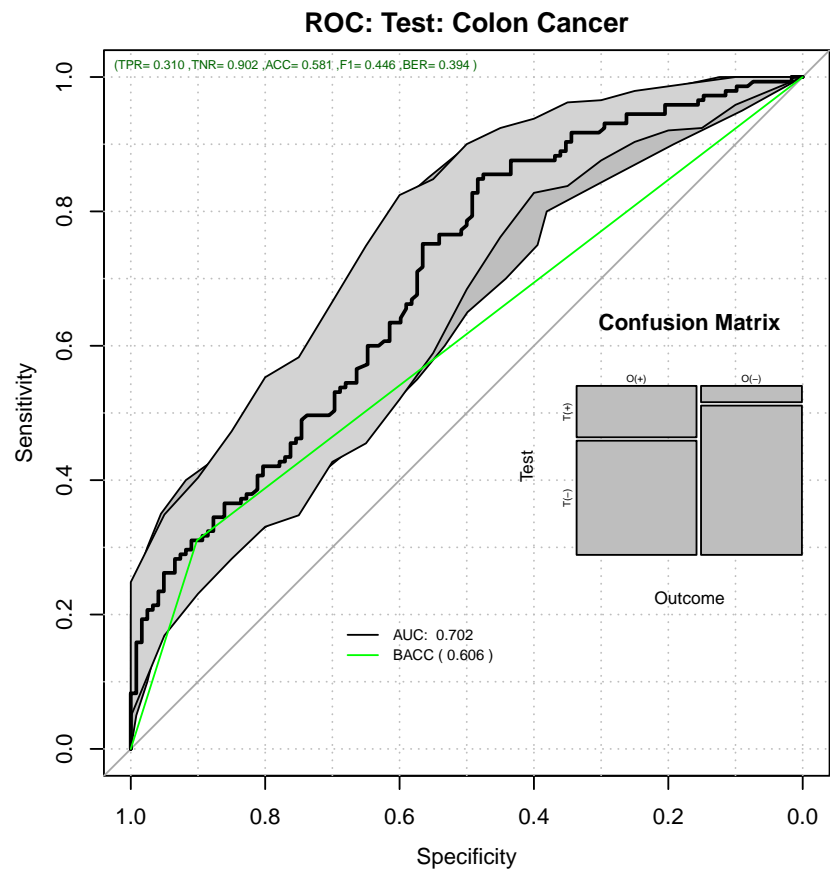


Decision Curve Analysis: Test: Colon Cancer

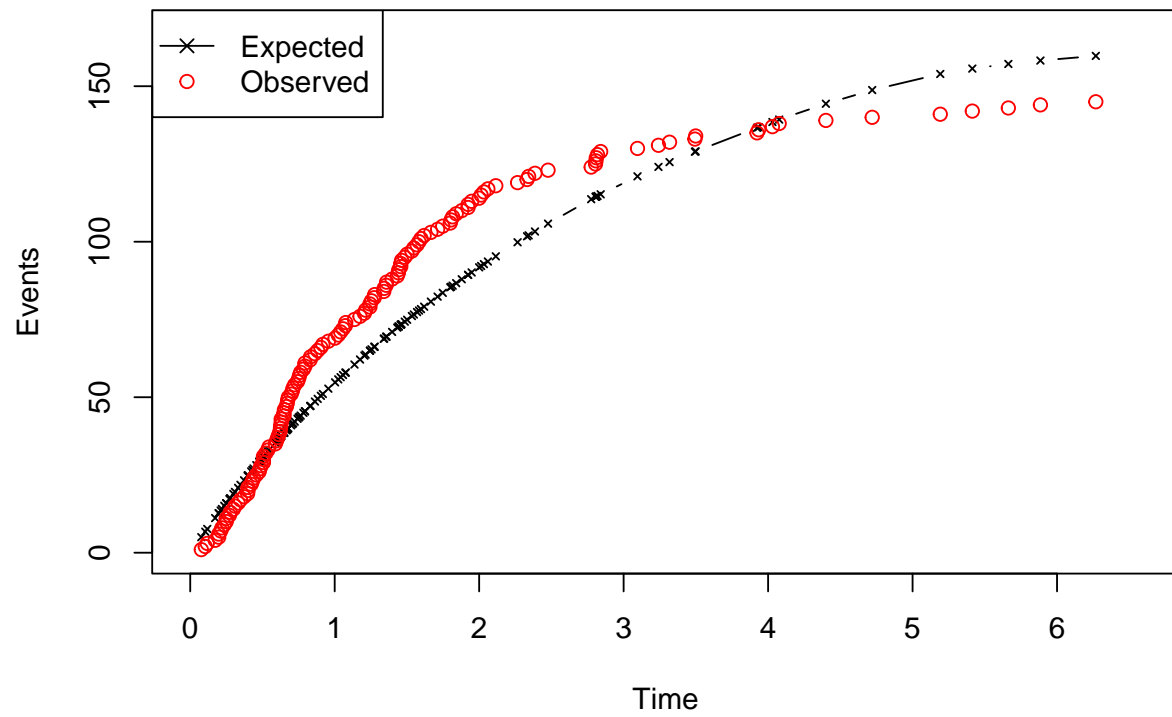


Relative Risk: Test: Colon Cancer

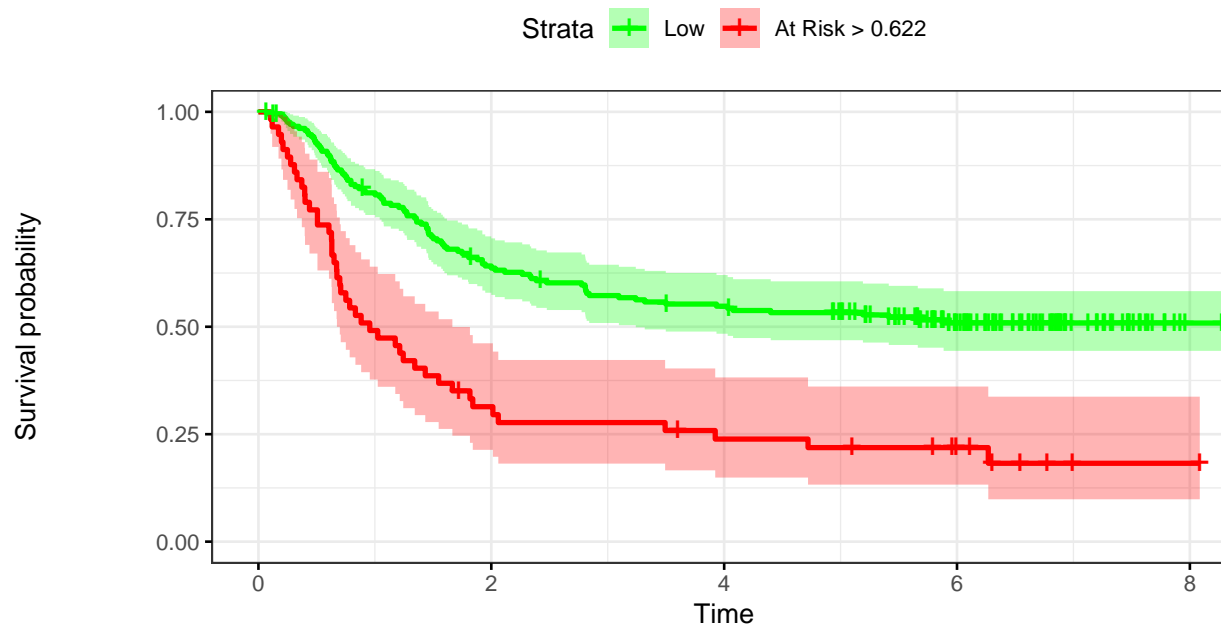




Time vs. Events: Test: Colon Cancer



Kaplan–Meier: Test: Colon Cancer



Number at risk

Low	210	131	110	68	4
At Risk > 0.622	57	17	12	7	1

0.2.7 Test Performance

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

Table 27: O/E Ratio

est	lower	upper
0.908	0.766	1.07

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

Table 28: O/E Ratio

mean	50%	2.5%	97.5%
1.12	1.12	1.09	1.15

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

Table 29: O/Acum Ratio

mean	50%	2.5%	97.5%
1.14	1.14	1.13	1.14

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

mean.C Index	median	lower	upper
0.653	0.654	0.609	0.698

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

Table 31: ROC AUC

est	lower	upper
0.702	0.64	0.764

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

Table 32: Sensitivity

est	lower	upper
0.31	0.236	0.392

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

Table 33: Specificity

est	lower	upper
0.902	0.834	0.948

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

Table 34: Probability Thresholds

90%
0.622

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

Table 35: Risk Ratio

est	lower	upper
1.66	1.36	2.02

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

Table 36: Logrank test Chisq = 30.415807 on 1 degrees of freedom,
p = 0.000000

	N	Observed	Expected	(O-E)^2/E	(O-E)^2/V
class=0	210	100	123.5	4.47	30.4
class=1	57	45	21.5	25.65	30.4

0.3 Cross-Validation

Here we will cross validate the training set and evaluate also on the testing set. The h0 and the timeinterval are the ones estimated on the calibration process

```
rcv <- randomCV(theData=dataColonTrain,
  theOutcome = Surv(time,status)~1,
  fittingFunction=BSWiMS.model,
  trainFraction = 0.75,
  repetitions=50,
  classSamplingType = "Pro",
  testingSet=dataColonTest
)
```

```
.[++++-].[+++++].[+++++].[++++-].[+++++].[+++++].[+++++].[+++++].[+++++].[+++++]10
Tested: 858 Avg. Selected: 8 Min Tests: 1 Max Tests: 10 Mean Tests: 4.93007 . MAD: 0.4773942
.[+++].[+++++].[+++++].[+++++].[+++++].[++++-].[+++++].[+++++].[++++-].[++-+ ]20 Tested:
887 Avg. Selected: 8.4 Min Tests: 1 Max Tests: 20 Mean Tests: 9.537768 . MAD: 0.4757449 .[+++-
].[+++++].[+++++].[+++++].[++++-].[+++++].[+++++].[+++++].[+++++].[+++++].[+++++]30
Tested: 888 Avg. Selected: 8.333333 Min Tests: 1 Max Tests: 30 Mean Tests: 14.29054 . MAD:
0.4755663 .[+++++].[+++++].[++++-].[+++++].[++++-].[+++++].[+++++].[+++++].[+++++].[+++++]40
Tested: 888 Avg. Selected: 8.25 Min Tests: 3 Max Tests: 40 Mean Tests: 19.05405 . MAD: 0.4757152
.[++++-].[+++++].[++++-].[+++++].[++++-].[+++++].[+++++].[+++++].[++++-].[+++++]50 Tested: 888 Avg.
Selected: 8.16 Min Tests: 4 Max Tests: 50 Mean Tests: 23.81757 . MAD: 0.4752228
```

```
stp <- rcv$urvTestPredictions
stp <- stp[!is.na(stp[,4]),]

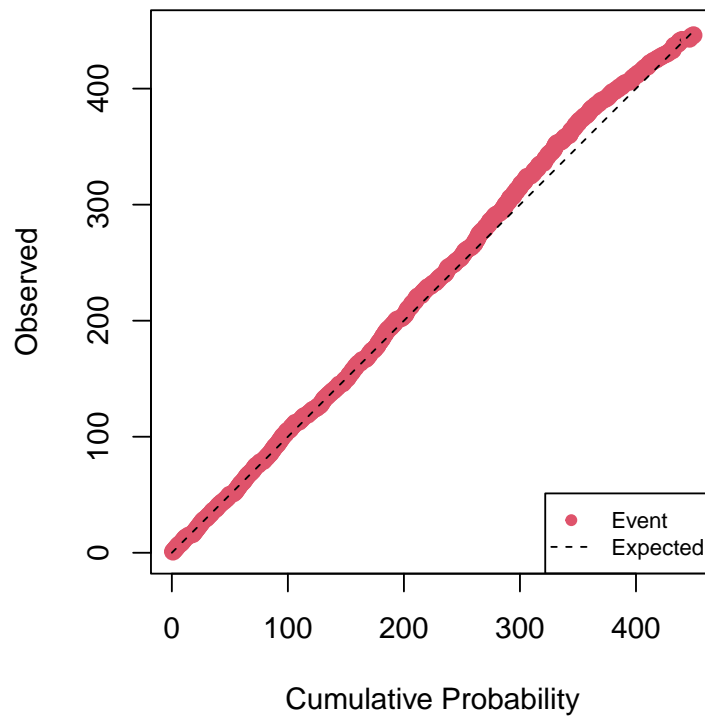
bbx <- boxplot(unlist(stp[,1])~rownames(stp), plot=FALSE)
times <- bbx$stats[3,]
status <- boxplot(unlist(stp[,2])~rownames(stp), plot=FALSE)$stats[3,]
prob <- ppoisGzero(boxplot(unlist(stp[,4])~rownames(stp), plot=FALSE)$stats[3,], h0)

rdatacv <- cbind(status, prob)
rownames(rdatacv) <- bbx$names
names(times) <- bbx$names

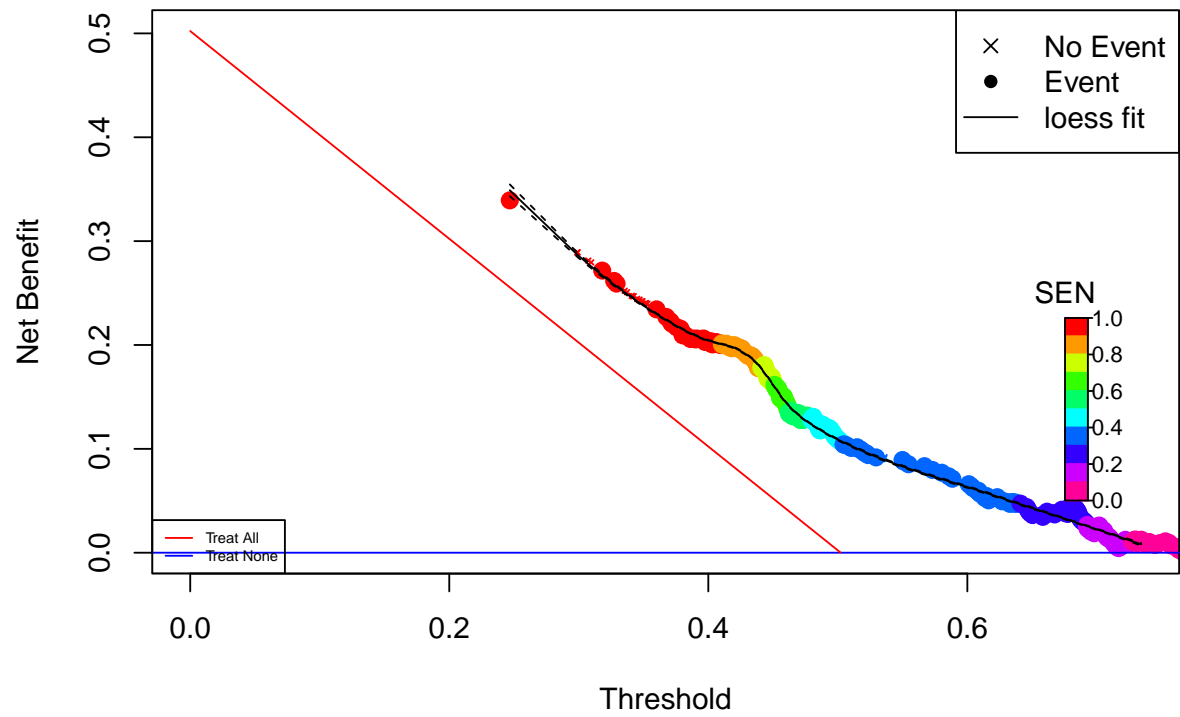
rrAnalysisCVTest <- RRPlot(rdatacv, atThr = rrAnalysisTrain$thr_atP,
  timetoEvent=times,
```

```
title="CV Test: Colon Cancer",  
ysurvlim=c(0.00,1.0),  
riskTimeInterval=timeinterval)
```

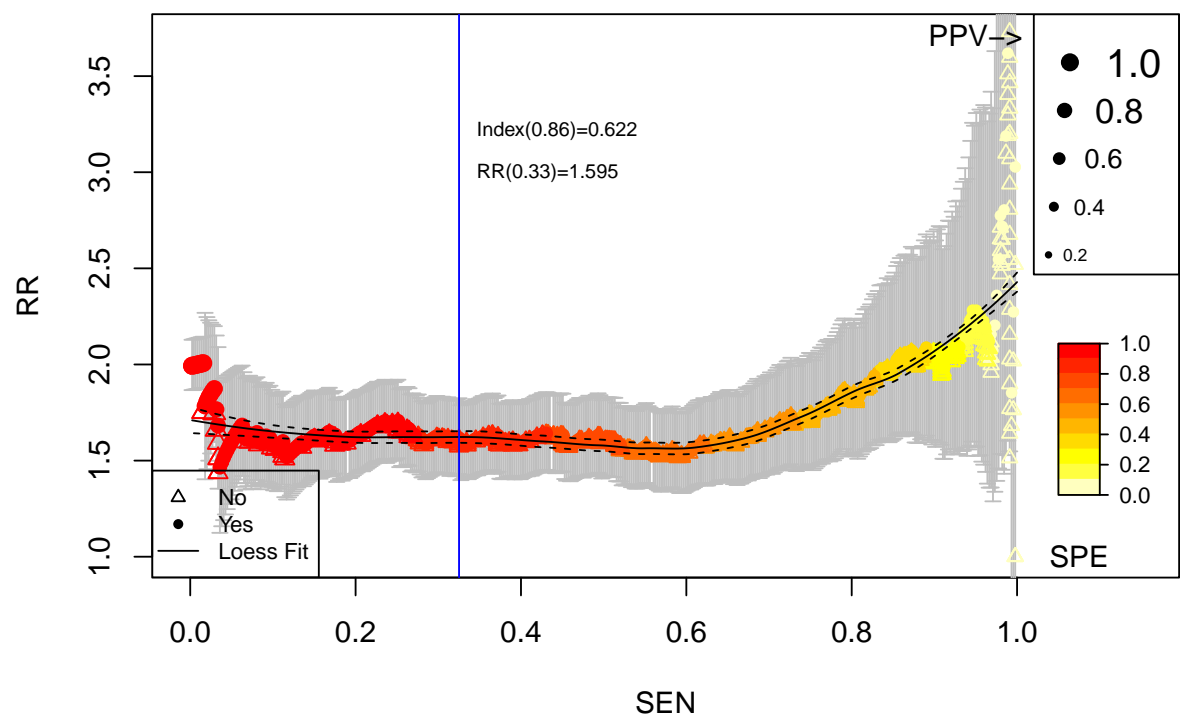
Cumulative vs. Observed: CV Test: Colon Cancer

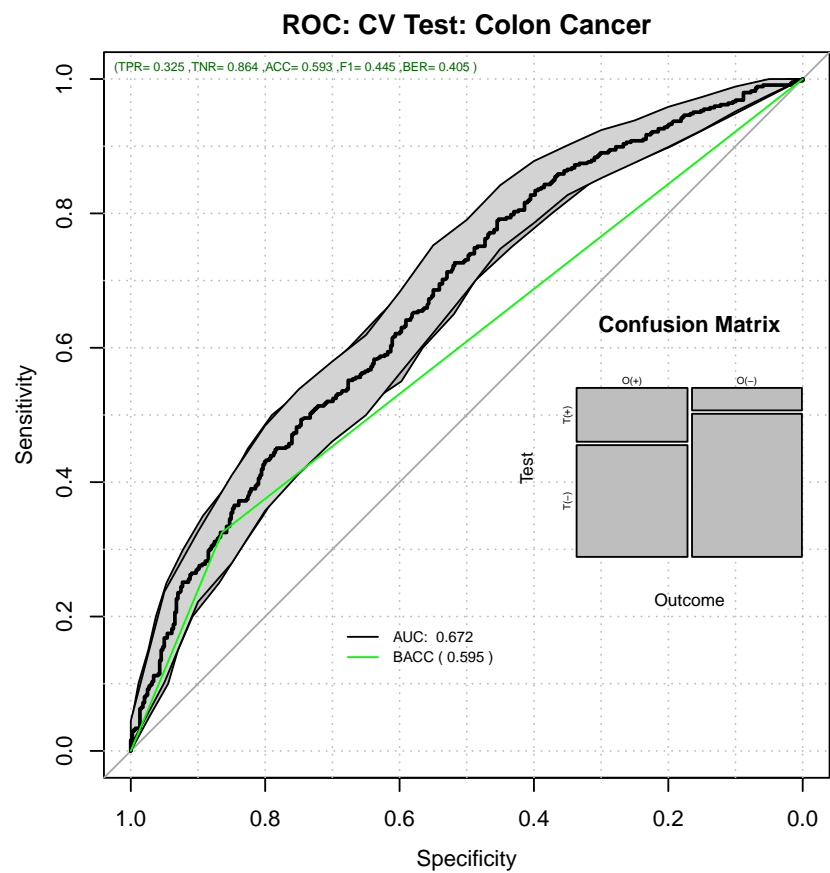


Decision Curve Analysis: CV Test: Colon Cancer

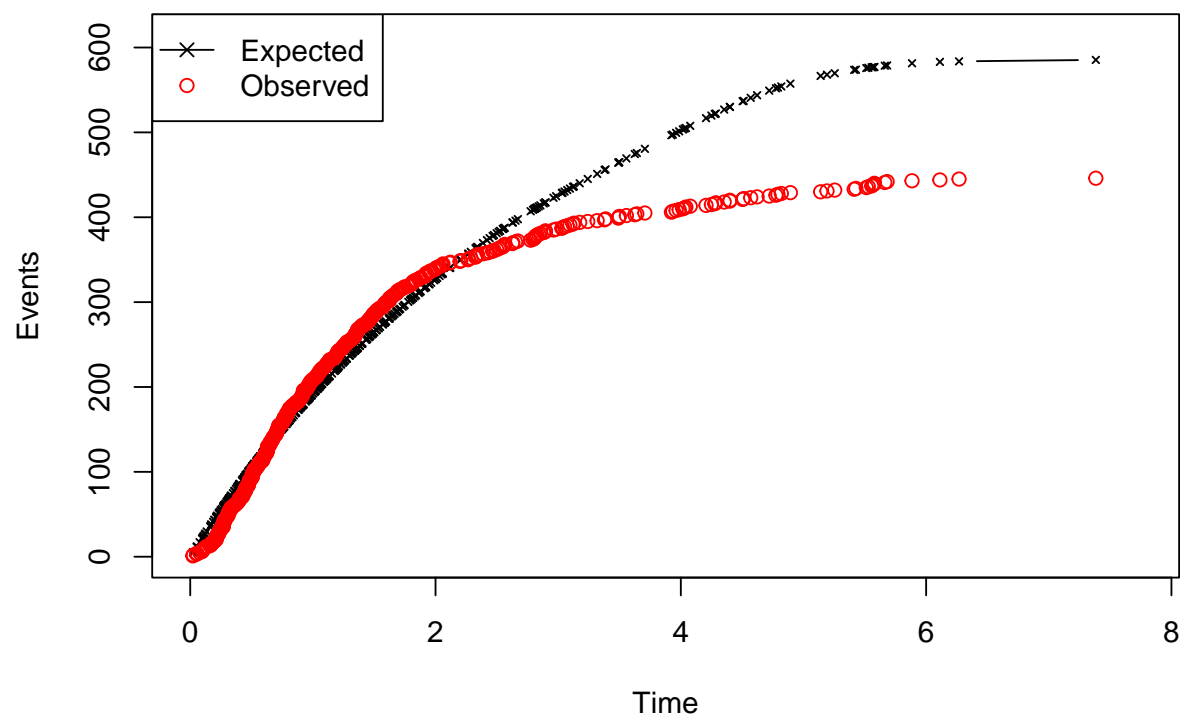


Relative Risk: CV Test: Colon Cancer

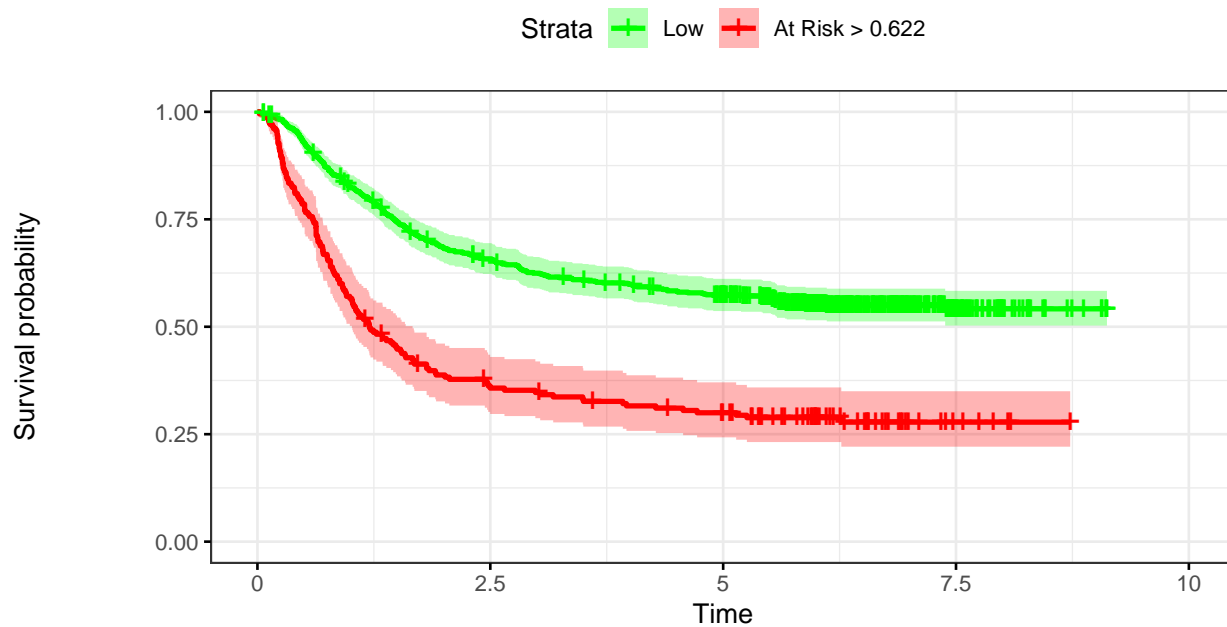




Time vs. Events: CV Test: Colon Cancer



Kaplan–Meier: CV Test: Colon Cancer



Number at risk

Low	683	437	365	56	0
At Risk > 0.622	205	70	54	6	0

0.3.1 CV Test Performance

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

Table 37: O/E Ratio

est	lower	upper
0.762	0.693	0.836

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

Table 38: O/E Ratio

mean	50%	2.5%	97.5%
0.954	0.954	0.942	0.967

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

Table 39: O/Acum Ratio

mean	50%	2.5%	97.5%
1.03	1.03	1.03	1.03

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

mean.C Index	median	lower	upper
0.645	0.646	0.62	0.67

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

Table 41: ROC AUC

est	lower	upper
0.672	0.637	0.707

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

Table 42: Sensitivity

est	lower	upper
0.325	0.282	0.371

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

Table 43: Specificity

est	lower	upper
0.864	0.829	0.895

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

Table 44: Probability Thresholds

90%
0.622

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

Table 45: Risk Ratio

est	lower	upper
1.59	1.41	1.8

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

Table 46: Logrank test Chisq = 74.685264 on 1 degrees of freedom,
p = 0.000000

	N	Observed	Expected	(O-E)^2/E	(O-E)^2/V
class=0	683	301	369.5	12.7	74.7
class=1	205	145	76.5	61.4	74.7