

R markdown Survival Analysis

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Introduction

The following code was used to perform the various survival analyses for this research. Please see for the code to download and process the Forest Service data. The UMN-PALS data (Fleischman et al., 2020) can be downloaded from here: <https://conservancy.umn.edu/handle/11299/211669>

Load libraries

```
library(tidyverse)

## -- Attaching packages ----- tidyverse 1.3.0 --
## v ggplot2 3.3.3      v purrr   0.3.4
## v tibble  3.1.0      v dplyr  1.0.5
## v tidyr   1.1.3      v stringr 1.4.0
## v readr   1.4.0      v forcats 0.5.1

## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()

library(forcats)
library(survminer)

## Loading required package: ggpubr

library(survival)
library(lubridate)

##
## Attaching package: 'lubridate'

## The following objects are masked from 'package:base':
##
##     date, intersect, setdiff, union

library(reshape2)

##
## Attaching package: 'reshape2'

## The following object is masked from 'package:tidyr':
##
##     smiths
```

```

library(ggplot2)
library(ggpubr)

# Set your working directory to wherever you placed your processed FS-PALS dataset.

setwd("/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/Oct_presentation/")

# Set up to see more rows of output if desired

options(max.print = 10000)

```

Load the data

Data from post-processing. See *S3.1* and *S3.2*.

```

## Load the data for survival analysis

df_fin <- read.csv("df_c20201020_v02.csv")
df_fin <- df_fin %>%
  select(-X)

## Replace "Inf" in the project delay column with NA
df_fin$proj.delay[is.infinite(df_fin$proj.delay)] <- NA
## Bring in binomial data as a factor
df_fin$REGION <- as.factor(df_fin$REGION)
df_fin$LITIGATED <- as.factor(df_fin$LITIGATED)
df_fin$APPEALED <- as.factor(df_fin$APPEALED)
df_fin$size <- as.factor(df_fin$size)

## Relevel the size data in order from small to extra-large
df_fin <- df_fin %>%
  mutate(size = fct_relevel(size,
    "small", "medium", "large", "x-large"))

## Create different data frames with Regions relevelled.
## This is required to test the impact of using different
## Regions as a baseline or reference in the Cox proportional
## analysis.
df_fin2 <- df_fin %>%
  mutate(REGION = fct_relevel(REGION,
    "2", "1", "3", "4", "5", "6"))

df_fin3 <- df_fin %>%
  mutate(REGION = fct_relevel(REGION,
    "3", "1", "2", "4", "5", "6"))

df_fin4 <- df_fin %>%
  mutate(REGION = fct_relevel(REGION,
    "4", "1", "2", "3", "5", "6"))

df_fin5 <- df_fin %>%
  mutate(REGION = fct_relevel(REGION,
    "5", "1", "2", "3", "4", "6"))

```

```
df_fin6 <- df_fin %>%
  mutate(REGION = fct_relevel(REGION,
    "6", "1", "2", "3", "4", "5"))
```

Check for data correlations

Why care?

When it comes to USFS project planning and inititaion there is some anecdotal evidence for NEPA causing delays... but very few quantitative studies.

Fuel, weather, and topography comprise the three legs of the Wildland Fire Behavior Triangle. Of those components, fuel is the only factor we have the ability to manage. Yet year after year fuels accumulate in our forests as management projects go neglected, delayed or obstructed. There is a clear correlation between the decline in timber harvests experienced on our National Forests and the increase in intensity and size of wildfires over the past three decades—both of which have had lasting impacts on the economic vitality of Western, rural communities. While our forests burn, our economic, recreational, and aesthetic capital burns with it.

While we recognize the ecological role wildfires can play in ecosystems, the severity and intensity of wildfires supersede that which should be occurring. Bureaucratic processes, burdensome regulations, external pressure, and judicial activism hamstrings our federal agencies from completing work on the ground in a timely manner. We must arm our federal land agencies with the tools they require to sustain the health and productivity of our nation's forests. Doing so is compatible with efforts to reduce emissions, as well-managed forests and the buildings constructed by the sustainable wood products that come from them have the potential to sequester carbon.

Figure 1: Recent letter to Congress from Western Congressional Caucus

Survival Analysis

“Time to Event”

- Time until an individual dies (or is cured!).
- Time until a kitten or puppy is adopted.
- Any time an outcome is a **duration**
- Requires a start date, end date, duration (time between), if the event occurred, and the time of observation.



Different approaches

- **Kaplan-Meier** -> non-parametric, predictive
- **Cox Proportional Hazards** -> semi-parametric, use to estimate effect of covariate, not predictive

Survival Analysis - Kaplan Meier

All data - Kaplan Meier (KM) Estimate

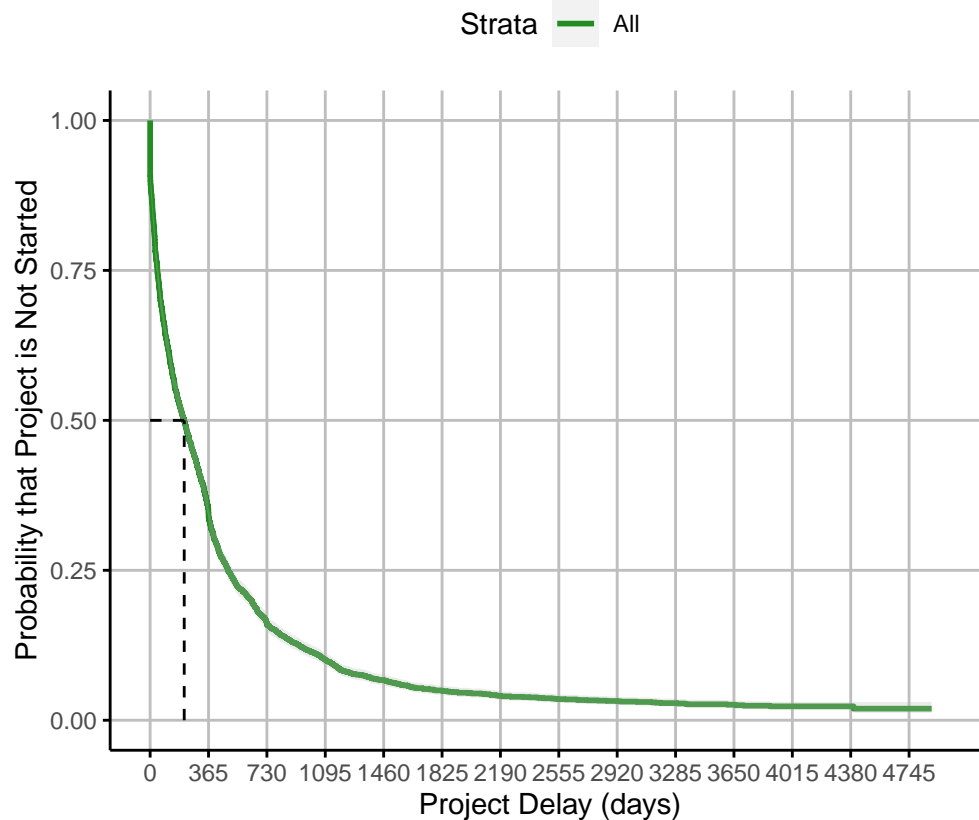
Using all the data pooled.

```
# Use the survfit function from the survival package to  
# calculate the KM estimate  
  
fit_delay_all <- survfit(Surv(proj.delay, INITIATED) ~ 1, data = df_fin)  
fit_delay_all  
  
## Call: survfit(formula = Surv(proj.delay, INITIATED) ~ 1, data = df_fin)  
##  
##          n  events  median 0.95LCL 0.95UCL  
##    3557    3289    213    194    228  
  
#summary(fit_delay_all)
```

Survival Analysis - All - Kaplan Meier Curves

All data

```
## ggsurvplot will create survival curves from the KM model created above.  
  
km_fit_all <- ggsurvplot(fit_delay_all,  
  conf.int = TRUE,  
  risk.table = FALSE,  
  risk.table.col = "strata",  
  surv.median.line = "hv",  
  break.time.by = 365,  
  risk.table.y.text=FALSE,  
  censor = FALSE,  
  ylab = "Probability that Project is Not Started",  
  xlab = "Project Delay (days)",  
  palette = c("forestgreen"),  
  surv.plot.height = 1,  
  ggtheme = theme(aspect.ratio = 0.75,  
    axis.line = element_line(colour = "black"),  
    panel.grid.major = element_line(colour = "grey"),  
    panel.border = element_blank(),  
    panel.background = element_blank()),  
  tables.theme = theme(aspect.ratio = 0.06)  
)  
print(km_fit_all)
```



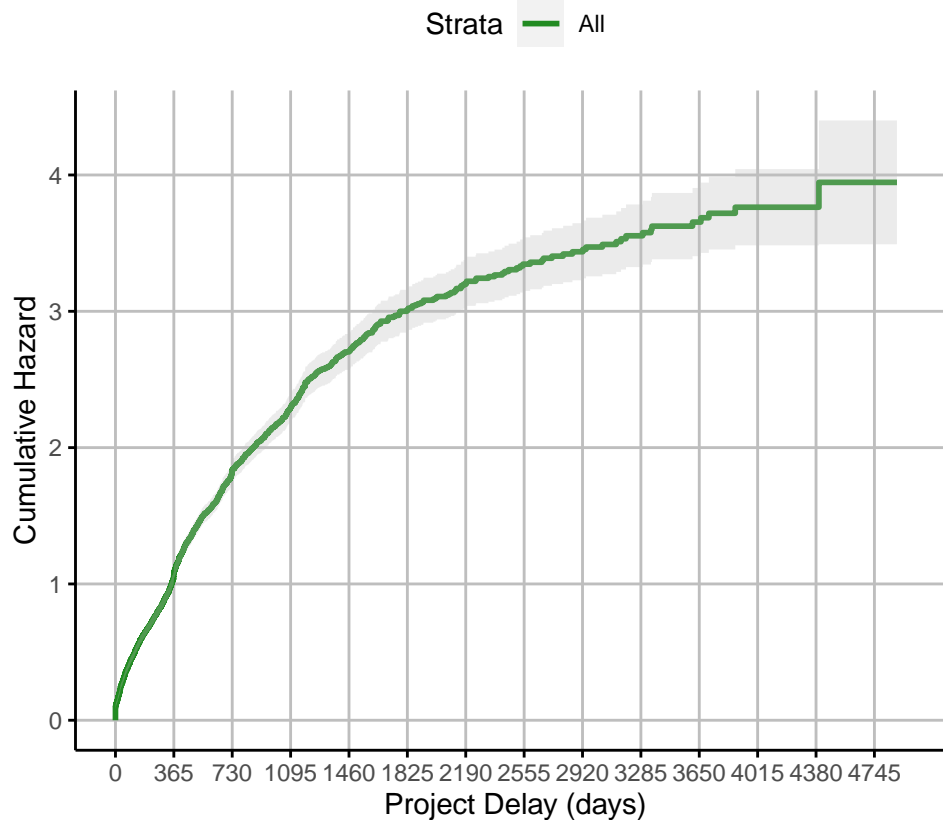
```
#ggsave( "/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/WRITING/FIGS/km_fit_all.pdf", print(km_fit
```

Survival Analysis - All - Kaplan Meier - Cumulative Hazard

All data

ggsurvplot will create the cumulative hazard plots based on the KM model created above.

```
km_haz_all <- ggsurvplot(fit_delay_all,
  conf.int = TRUE,
  risk.table = FALSE,
  risk.table.col = "strata",
  break.time.by = 365,
  risk.table.y.text=FALSE,
  censor = FALSE,
  ylab = "Cumulative Hazard",
  xlab = "Project Delay (days)",
  palette = c("forestgreen"),
  fun = "cumhaz",
  surv.plot.height = 1,
  ggtheme = theme(aspect.ratio = 0.75,
    axis.line = element_line(colour = "black"),
    panel.grid.major = element_line(colour = "grey"),
    panel.border = element_blank(),
    panel.background = element_blank()),
  tables.theme = theme(aspect.ratio = 0.06)
)
print(km_haz_all)
```



```
#ggsave( "/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/WRITING/FIGS/km_haz_all.pdf", print(km_haz
```

Survival Analysis - Appealed? - KM Estimate

Data grouped by appealed and non-appealed projects.

```
# Use the survfit function from the survival package to
# calculate the KM estimate for data grouped by appealed or non-appealed.
# This code chunk and the next two act as a template for the K-M estimation.
```

```
fit_app <- survfit(Surv(proj.delay, INITIATED) ~ APPEALED, data = df_fin)
fit_app_table <- summary(fit_app)
fit_app
```

```
## Call: survfit(formula = Surv(proj.delay, INITIATED) ~ APPEALED, data = df_fin)
##
##              n events median 0.95LCL 0.95UCL
## APPEALED=0 3184   2937   198     182     220
## APPEALED=1  373    352   303     258     365
```

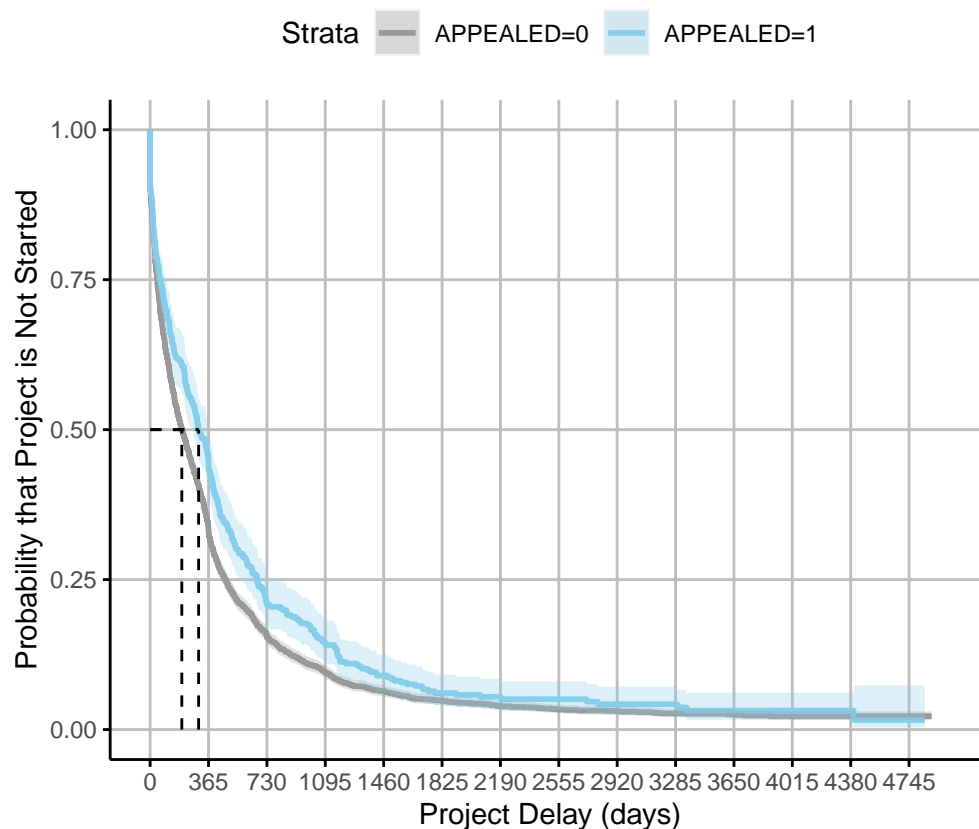
Survival Analysis - Appealed? - KM Curve

```
km_curv_app <- ggsurvplot(fit_app,
  conf.int = TRUE,
  risk.table = FALSE,
  risk.table.col = "strata",
  surv.median.line = "hv",
  break.time.by = 365,
```

```

risk.table.y.text=FALSE,
tables.height = 0.3,
censor = FALSE,
ylab = "Probability that Project is Not Started",
xlab = "Project Delay (days)",
palette = c("#999999", "skyblue"),
surv.plot.height = 1,
ggtheme = theme(aspect.ratio = 0.75,
                 axis.line = element_line(colour = "black"),
                 panel.grid.major = element_line(colour = "grey"),
                 panel.border = element_blank(),
                 panel.background = element_blank()),
tables.theme = theme(aspect.ratio = 0.06)
)
print(km_curv_app)

```



```
#ggsave( "/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/WRITING/FIGS/km_curv_app.pdf", print(km_curv_app) )
```

Survival Analysis - Appealed? - KM Cumulative Hazard

```

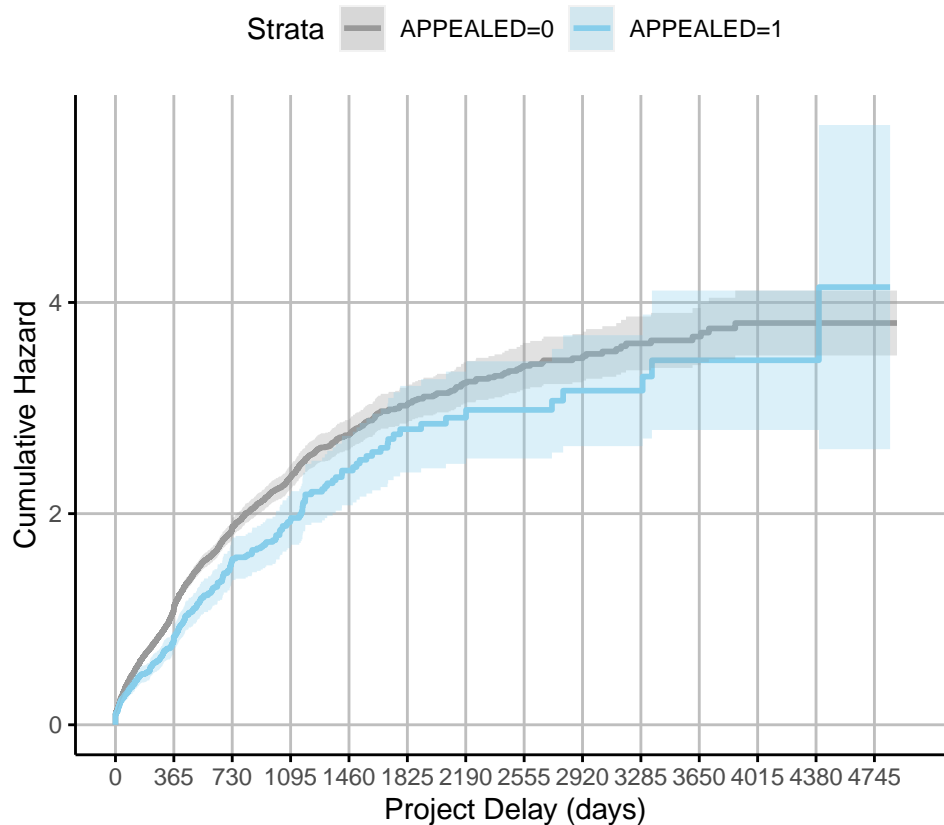
km_haz_app <- ggsurvplot(fit_app,
  conf.int = TRUE,
  risk.table = FALSE,
  risk.table.col = "strata",
  break.time.by = 365,
  risk.table.y.text=FALSE,
  censor = FALSE,

```

```

ylab = "Cumulative Hazard",
xlab = "Project Delay (days)",
palette = c("#999999", "skyblue"),
fun = "cumhaz",
surv.plot.height = 1,
ggtheme = theme(aspect.ratio = 0.75,
                 axis.line = element_line(colour = "black"),
                 panel.grid.major = element_line(colour = "grey"),
                 panel.border = element_blank(),
                 panel.background = element_blank()),
tables.theme = theme(aspect.ratio = 0.06)
)
print(km_haz_app)

```



```
#ggsave( "/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/WRITING/FIGS/km_haz_app.pdf", print(km_haz_app) )
```

Survival Analysis - Litigated? - KM Estimate

Data grouped by litigated and non-litigated projects.

```

fit_lit <- survfit(Surv(proj.delay, INITIATED) ~ LITIGATED, data = df_fin)
fit_lit_table <- summary(fit_lit)
fit_lit

```

```

## Call: survfit(formula = Surv(proj.delay, INITIATED) ~ LITIGATED, data = df_fin)
##
##              n events median 0.95LCL 0.95UCL
## LITIGATED=0 3445   3194   207     190    226

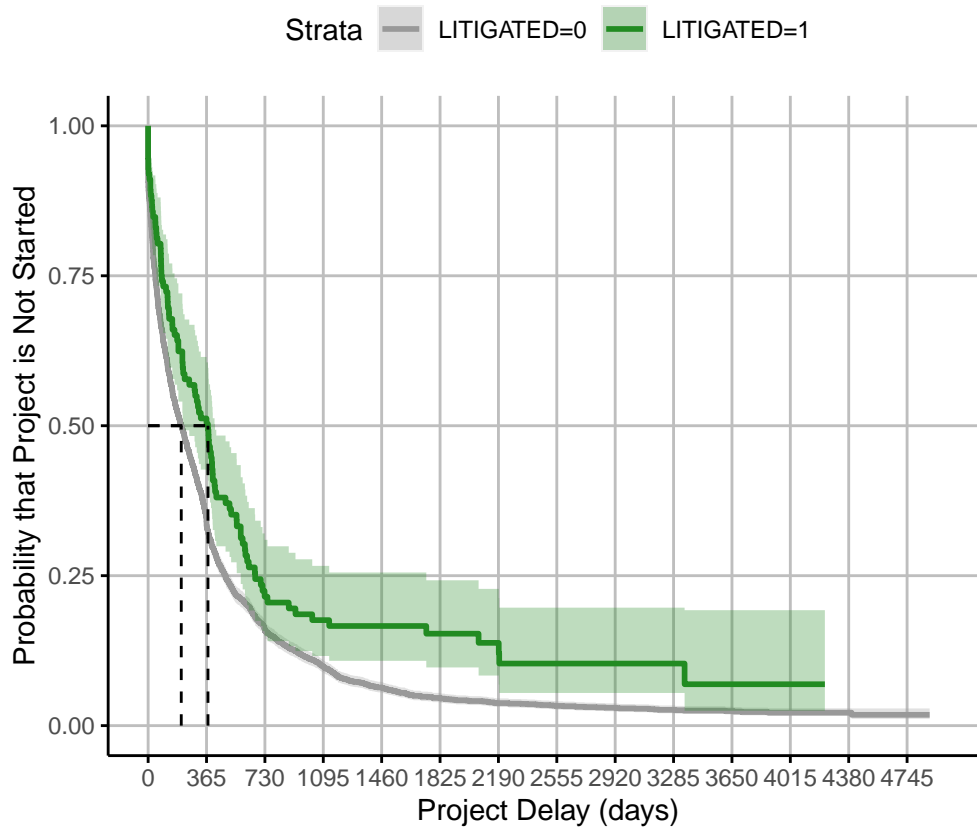
```



```
## LITIGATED=1 112 95 374 228 418
```

Survival Analysis - Litigated? - KM Curve

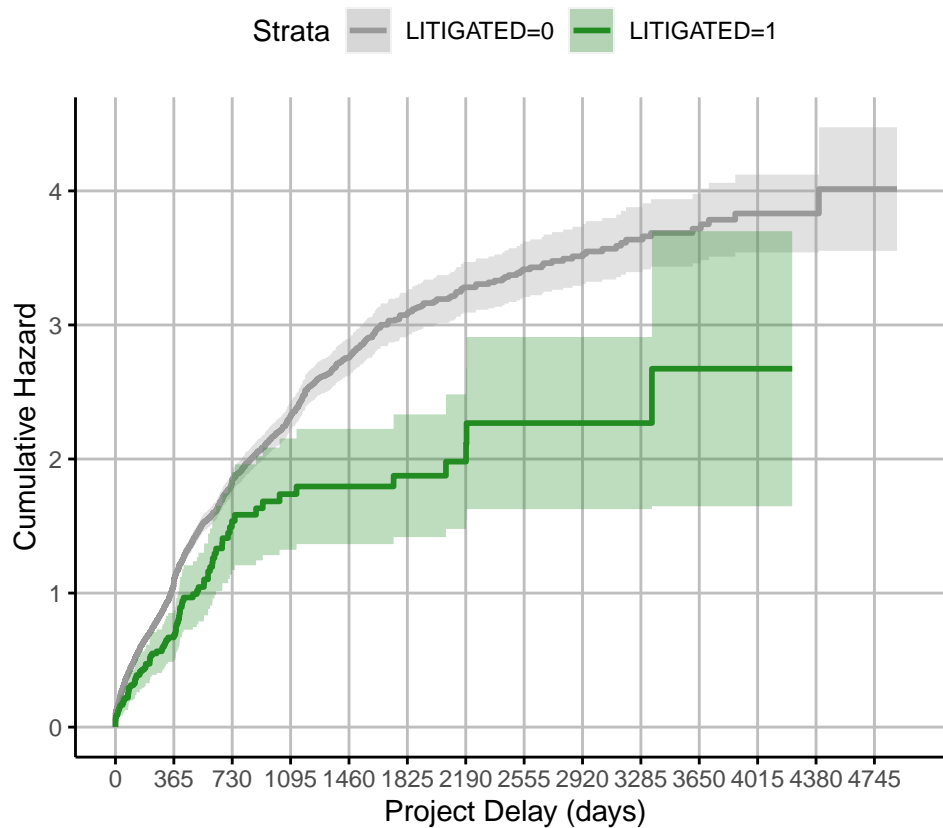
```
km_curv_lit <- ggsurvplot(fit_lit,
  conf.int = TRUE,
  risk.table = FALSE,
  risk.table.col = "strata",
  surv.median.line = "hv",
  break.time.by = 365,
  risk.table.y.text=FALSE,
  tables.height = 0.3,
  censor = FALSE,
  ylab = "Probability that Project is Not Started",
  xlab = "Project Delay (days)",
  palette = c("#999999", "forestgreen"),
  surv.plot.height = 1,
  ggtheme = theme(aspect.ratio = 0.75,
    axis.line = element_line(colour = "black"),
    panel.grid.major = element_line(colour = "grey"),
    panel.border = element_blank(),
    panel.background = element_blank()),
  tables.theme = theme(aspect.ratio = 0.06)
)
print(km_curv_lit)
```



```
#ggsave( "/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/WRITING/FIGS/km_curv_lit.pdf", print(km_cu
```

Survival Analysis - Litigated? - KM Cumulative Hazard

```
km_haz_lit <- ggsurvplot(fit_lit,
  conf.int = TRUE,
  risk.table = FALSE,
  risk.table.col = "strata",
  break.time.by = 365,
  risk.table.y.text=FALSE,
  censor = FALSE,
  ylab = "Cumulative Hazard",
  xlab = "Project Delay (days)",
  palette = c("#999999", "forestgreen"),
  fun = "cumhaz",
  surv.plot.height = 1,
  ggtheme = theme(aspect.ratio = 0.75,
    axis.line = element_line(colour = "black"),
    panel.grid.major = element_line(colour = "grey"),
    panel.border = element_blank(),
    panel.background = element_blank()),
  tables.theme = theme(aspect.ratio = 0.06)
)
print(km_haz_lit)
```



```
#ggsave( "/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/WRITING/FIGS/km_haz_lit.pdf", print(km_haz
```

Survival Analysis - NEPA Type - KM Estimate

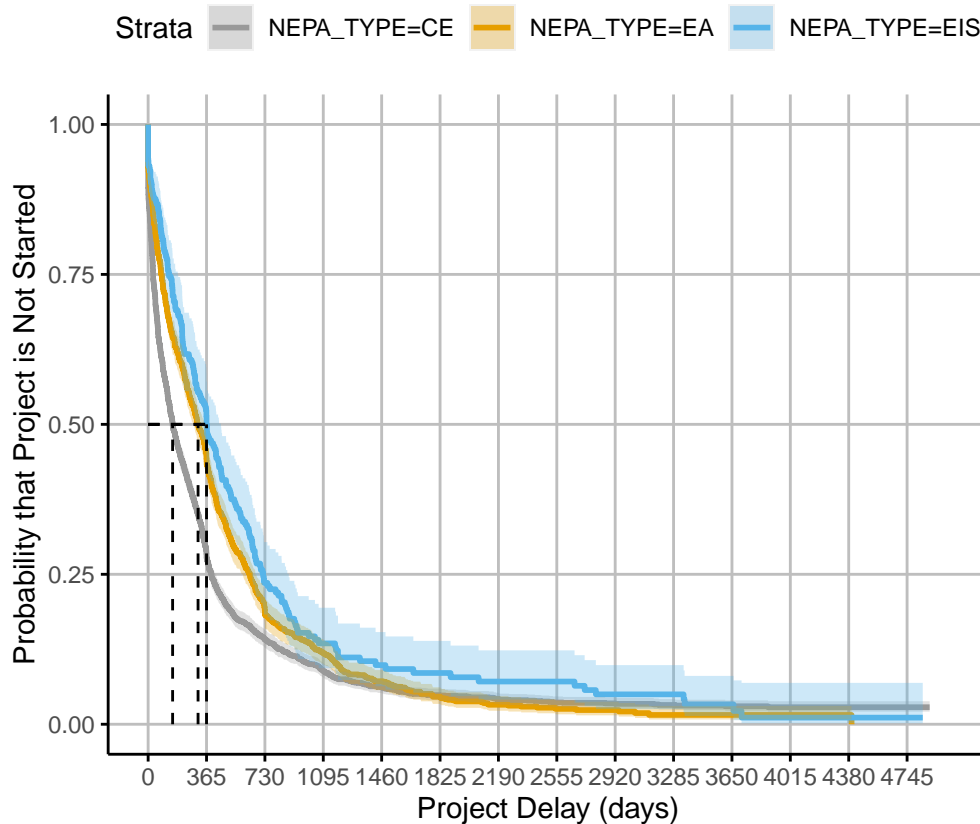
Data grouped by NEPA type for projects. *EIS* = Environmental Impact Statement, *EA* = Environmental Assessment, *CE* = Categorical Exclusion

```
fit_delay <- survfit(Surv(proj.delay, INITIATED) ~ NEPA_TYPE, data = df_fin)
fit_nepa_table <- summary(fit_delay)
fit_delay
```

```
## Call: survfit(formula = Surv(proj.delay, INITIATED) ~ NEPA_TYPE, data = df_fin)
##
##              n events median 0.95LCL 0.95UCL
## NEPA_TYPE=CE  2266   2110    153     139     171
## NEPA_TYPE=EA  1073    988    312     285     350
## NEPA_TYPE=EIS   218    191    365     303     455
```

Survival Analysis - NEPA Type - KM Curve

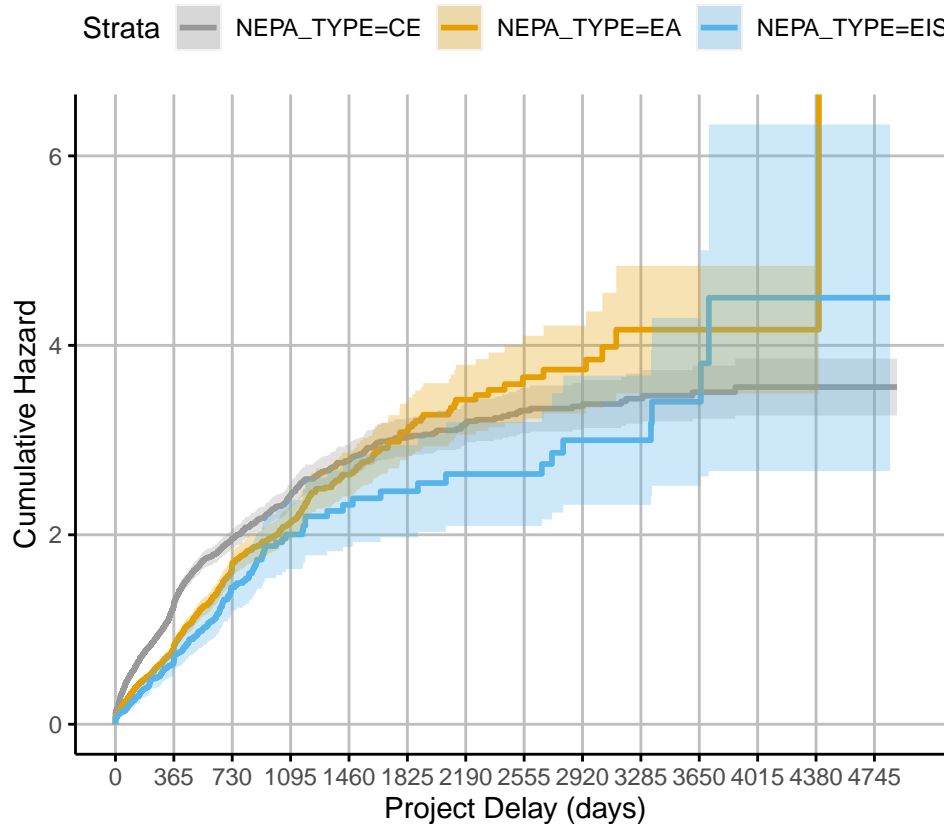
```
km_curv_nepa <- ggsurvplot(fit_delay,
  conf.int = TRUE,
  risk.table = FALSE,
  risk.table.col = "strata",
  surv.median.line = "hv",
  break.time.by = 365,
  risk.table.y.text=FALSE,
  tables.height = 0.3,
  censor = FALSE,
  ylab = "Probability that Project is Not Started",
  xlab = "Project Delay (days)",
  palette = c("#999999", "#E69F00", "#56B4E9"),
  surv.plot.height = 1,
  ggtheme = theme(aspect.ratio = 0.75,
    axis.line = element_line(colour = "black"),
    panel.grid.major = element_line(colour = "grey"),
    panel.border = element_blank(),
    panel.background = element_blank()),
  tables.theme = theme(aspect.ratio = 0.06)
)
print(km_curv_nepa)
```



```
#ggsave( "/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/WRITING/FIGS/km_curv_nepa.pdf", print(km_c
```

Survival Analysis - NEAP Type - KM Cumulative Hazard

```
km_haz_nepa <- ggsurvplot(fit_delay,
  conf.int = TRUE,
  risk.table = FALSE,
  risk.table.col = "strata",
  break.time.by = 365,
  risk.table.y.text=FALSE,
  tables.height = 0.3,
  censor = FALSE,
  ylab = "Cumulative Hazard",
  xlab = "Project Delay (days)",
  palette = c("#999999", "#E69F00", "#56B4E9"),
  fun = "cumhaz",
  surv.plot.height = 1,
  ggtheme = theme(aspect.ratio = 0.75,
    axis.line = element_line(colour = "black"),
    panel.grid.major = element_line(colour = "grey"),
    panel.border = element_blank(),
    panel.background = element_blank(),
    tables.theme = theme(aspect.ratio = 0.06)
  )
print(km_haz_nepa)
```



```
#ggsave( "/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/WRITING/FIGS/km_haz_nepa.pdf", print(km_ha
```

Survival Analysis - Regions - KM Estimate

Data grouped by project location. United States Forest Service (USFS) Regions 1-6

```
fit_reg <- survfit(Surv(proj.delay, INITIATED) ~ REGION, data = df_fin)
fit_reg_table <- summary(fit_reg)
fit_reg
```

```
## Call: survfit(formula = Surv(proj.delay, INITIATED) ~ REGION, data = df_fin)
##
##           n events median 0.95LCL 0.95UCL
## REGION=1  538    503   179    153    218
## REGION=2  455    405   245    197    323
## REGION=3  301    293   245    183    287
## REGION=4  540    507   126     94    173
## REGION=5  928    845   228    195    261
## REGION=6  795    736   245    205    294
```

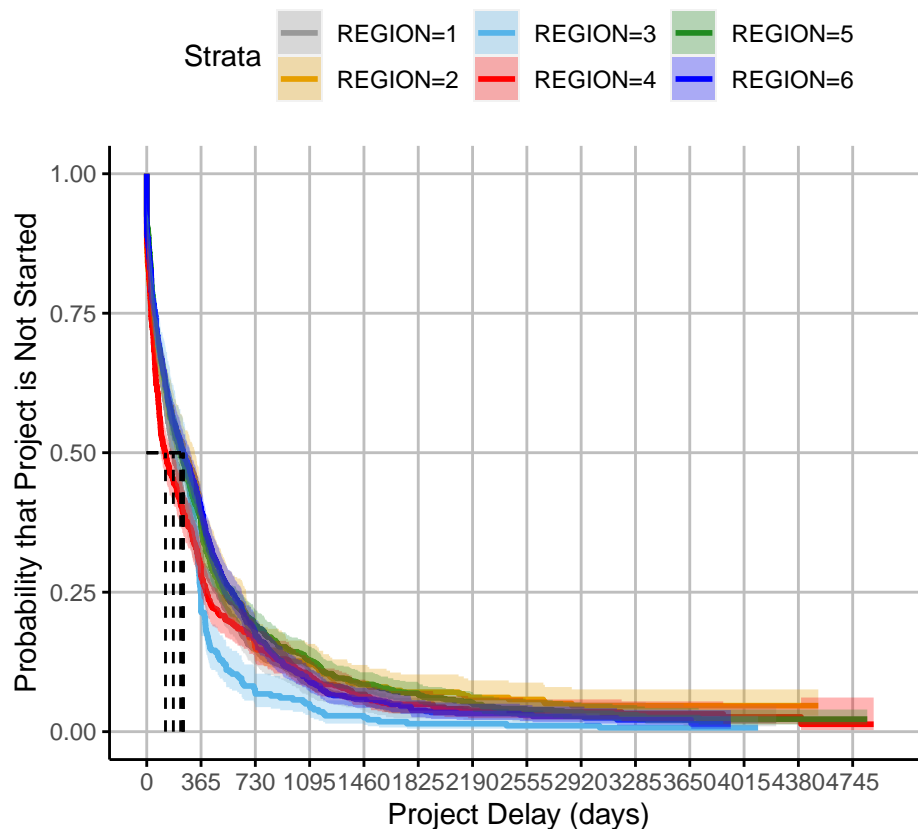
Survival Analysis - Regions - KM Curve

```
km_curv_reg <- ggsurvplot(fit_reg,
  conf.int = TRUE,
  risk.table = FALSE,
  risk.table.col = "strata",
  surv.median.line = "hv",
  break.time.by = 365,
```

```

risk.table.y.text=FALSE,
tables.height = 0.3,
censor = FALSE,
ylab = "Probability that Project is Not Started",
xlab = "Project Delay (days)",
palette = c("#999999", "#E69F00", "#56B4E9", "red", "forestgreen", "blue"),
surv.plot.height = 1,
ggtheme = theme(aspect.ratio = 0.75,
                 axis.line = element_line(colour = "black"),
                 panel.grid.major = element_line(colour = "grey"),
                 panel.border = element_blank(),
                 panel.background = element_blank()),
tables.theme = theme(aspect.ratio = 0.06)
)
print(km_curv_reg)

```



```
#ggsave( "/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/WRITING/FIGS/km_curv_reg.pdf", print(km_curv_reg) )
```

Survival Analysis - Regions - KM Cumulative Hazard

```

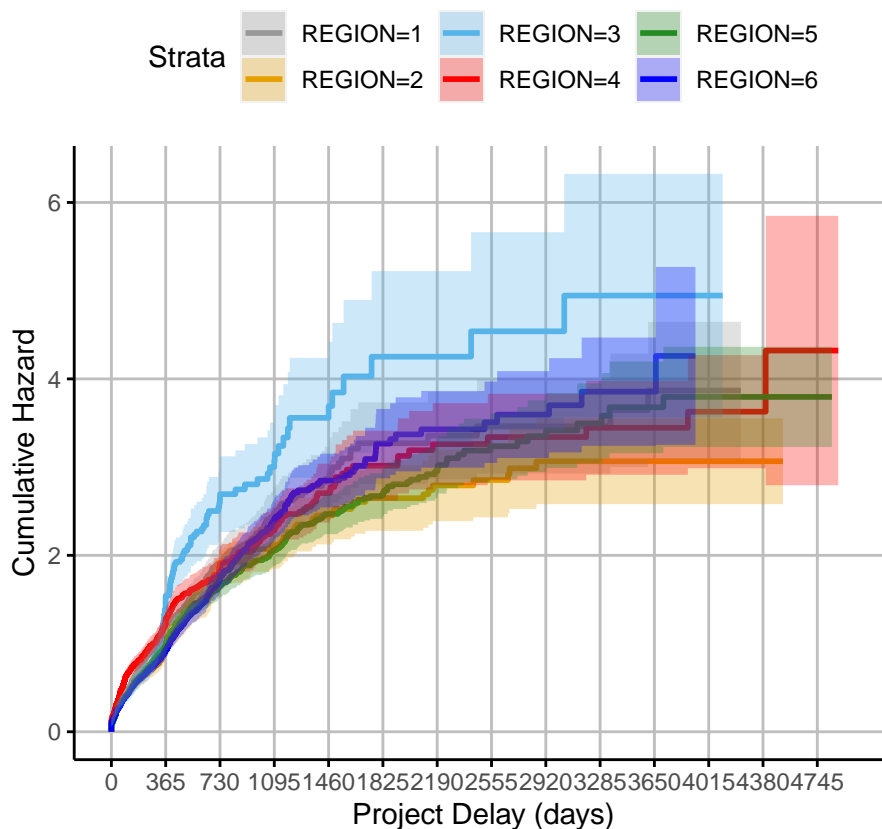
km_haz_reg <- ggsurvplot(fit_reg,
  conf.int = TRUE,
  risk.table = FALSE,
  risk.table.col = "strata",
  break.time.by = 365,
  risk.table.y.text=FALSE,
  tables.height = 0.3,

```

```

  censor = FALSE,
  ylab = "Cumulative Hazard",
  xlab = "Project Delay (days)",
  palette = c("#999999", "#E69F00", "#56B4E9", "red", "forestgreen", "blue"),
  fun = "cumhaz",
  surv.plot.height = 1,
  ggtheme = theme(aspect.ratio = 0.75,
    axis.line = element_line(colour = "black"),
    panel.grid.major = element_line(colour = "grey"),
    panel.border = element_blank(),
    panel.background = element_blank(),
    tables.theme = theme(aspect.ratio = 0.06)
  )
print(km_haz_reg)

```



```
#ggsave( "/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/WRITING/FIGS/km_haz_reg.pdf", print(km_haz_reg) )
```

Survival Analysis - Size - KM Estimate

Data grouped by a project's cumulative size. Size categories = small, medium, large, and extra-large.

Sizes are categorized based on quartile ranges + x-large > 2670 acres + large 768-2670 acres + medium 174-768 acres + small < 174 acres

```

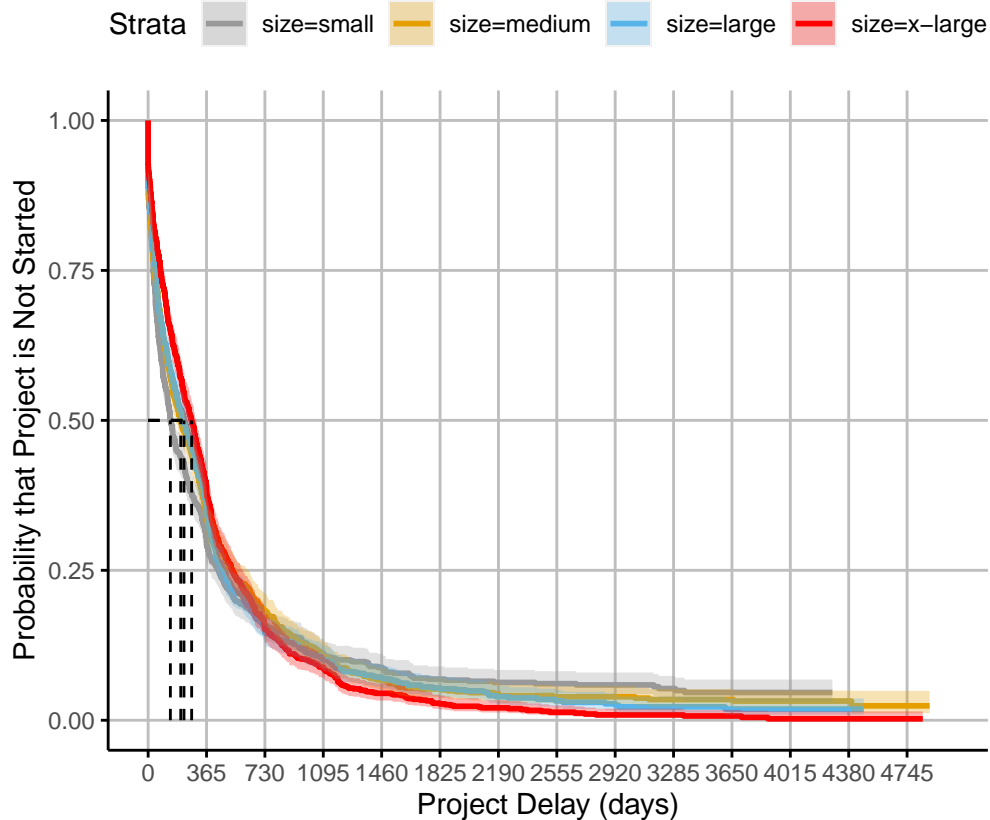
fit_size <- survfit(Surv(proj.delay, INITIATED) ~ size, data = df_fin)
fit_size_table <- summary(fit_size)
fit_size

```

```
## Call: survfit(formula = Surv(proj.delay, INITIATED) ~ size, data = df_fin)
##
##           n events median 0.95LCL 0.95UCL
## size=small  888    803    139    123    167
## size=medium 890    811    204    171    245
## size=large  888    814    224    187    261
## size=x-large 891    861    272    235    300
```

Survival Analysis - Size - KM Curve

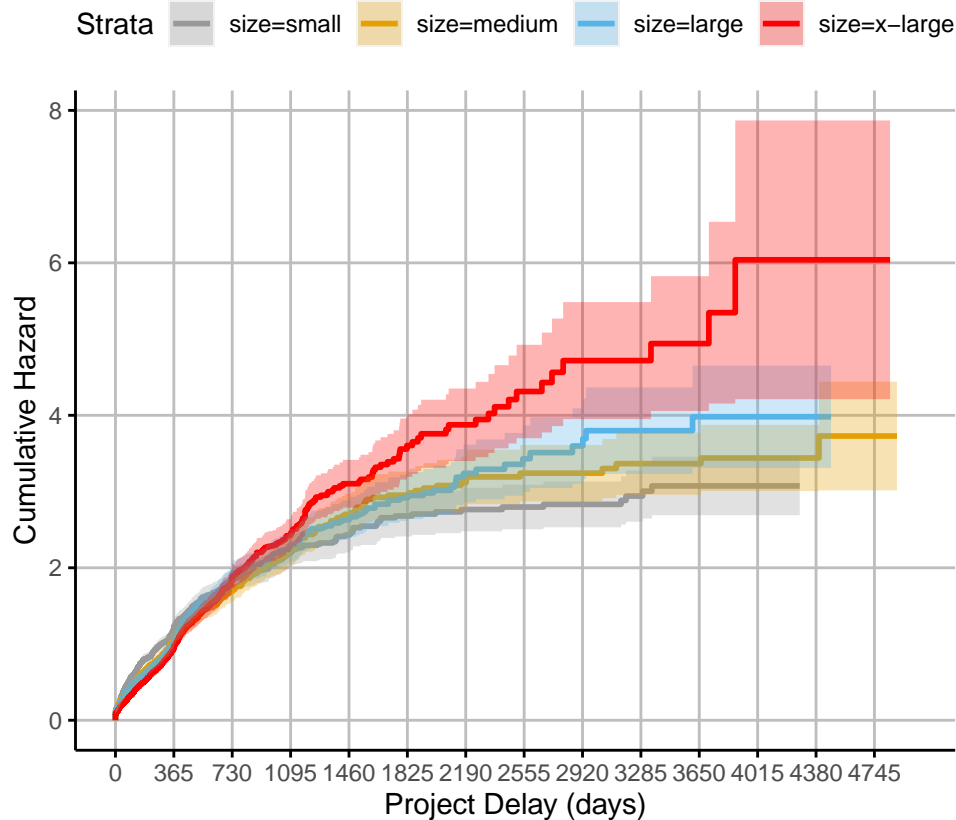
```
km_curv_size <- ggsurvplot(fit_size,
  conf.int = TRUE,
  risk.table = FALSE,
  risk.table.col = "strata",
  surv.median.line = "hv",
  break.time.by = 365,
  risk.table.y.text=FALSE,
  tables.height = 0.3,
  censor = FALSE,
  ylab = "Probability that Project is Not Started",
  xlab = "Project Delay (days)",
  palette = c("#999999", "#E69F00", "#56B4E9", "red"),
  surv.plot.height = 1,
  ggtheme = theme(aspect.ratio = 0.75,
    axis.line = element_line(colour = "black"),
    panel.grid.major = element_line(colour = "grey"),
    panel.border = element_blank(),
    panel.background = element_blank()),
  tables.theme = theme(aspect.ratio = 0.06)
)
print(km_curv_size)
```

```
#ggsave( "/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/WRITING/FIGS/km_curv_size.pdf", print(km_c
```

Survival Analysis - Size - KM Cumulative Hazard

```
km_haz_size <- ggsurvplot(fit_size,
  conf.int = TRUE,
  risk.table = FALSE,
  risk.table.col = "strata",
  break.time.by = 365,
  risk.table.y.text=FALSE,
  tables.height = 0.3,
  censor = FALSE,
  ylab = "Cumulative Hazard",
  xlab = "Project Delay (days)",
  palette = c("#999999", "#E69F00", "#56B4E9", "red"),
  fun = "cumhaz",
  surv.plot.height = 1,
  ggtheme = theme(aspect.ratio = 0.75,
    axis.line = element_line(colour = "black"),
    panel.grid.major = element_line(colour = "grey"),
    panel.border = element_blank(),
    panel.background = element_blank(),
    tables.theme = theme(aspect.ratio = 0.06)
  )
print(km_haz_size)
```



```
#ggsave( "/Users/kathrynmurenbeeld/Desktop/Survival_Analysis/WRITING/FIGS/km_haz_size.pdf", print(km_ha
```

Comparing survival curves using a log-rank approach

- Null hypothesis: no difference in survival between the groups
- Non-parametric test

Log-Rank - Appealed?

```
# The survdiff function in the survival package does this for you.
```

```
surv_diff_delay_app <- survdiff(Surv(proj.delay, INITIATED) ~ APPEALED, data = df_fin)
print(surv_diff_delay_app)
```

```
## Call:
## survdiff(formula = Surv(proj.delay, INITIATED) ~ APPEALED, data = df_fin)
##
##               N Observed Expected (O-E)^2/E (O-E)^2/V
## APPEALED=0 3184      2937      2870       1.58      12.6
## APPEALED=1  373       352       419      10.83      12.6
##
##  Chisq= 12.6  on 1 degrees of freedom, p= 4e-04
```

Log-Rank - Litigated?

```
surv_diff_delay_lit <- survdiff(Surv(proj.delay, INITIATED) ~ LITIGATED, data = df_fin)
print(surv_diff_delay_lit)
```

```
## Call:
## survdiff(formula = Surv(proj.delay, INITIATED) ~ LITIGATED, data = df_fin)
```

```
##
##           N Observed Expected (O-E)^2/E (O-E)^2/V
## LITIGATED=0 3445      3194      3154      0.504      12.4
## LITIGATED=1  112        95       135     11.787      12.4
##
## Chisq= 12.4 on 1 degrees of freedom, p= 4e-04

Log-Rank - NEPA
surv_diff_delay_nepa <- survdiff(Surv(proj.delay, INITIATED) ~ NEPA_TYPE, data = df_fin)
print(surv_diff_delay_nepa)

## Call:
## survdiff(formula = Surv(proj.delay, INITIATED) ~ NEPA_TYPE, data = df_fin)
##
##           N Observed Expected (O-E)^2/E (O-E)^2/V
## NEPA_TYPE=CE 2266      2110      1898      23.7      57.1
## NEPA_TYPE=EA 1073       988      1135      19.1      29.7
## NEPA_TYPE=EIS 218       191       256      16.4      18.0
##
## Chisq= 60.3 on 2 degrees of freedom, p= 8e-14

Log-Rank - Region
surv_diff_delay_reg <- survdiff(Surv(proj.delay, INITIATED) ~ REGION, data = df_fin)
print(surv_diff_delay_reg)

## Call:
## survdiff(formula = Surv(proj.delay, INITIATED) ~ REGION, data = df_fin)
##
##           N Observed Expected (O-E)^2/E (O-E)^2/V
## REGION=1 538       503       477       1.47       1.74
## REGION=2 455       405       441       2.93       3.43
## REGION=3 301       293       241      11.44      12.59
## REGION=4 540       507       457       5.41       6.37
## REGION=5 928       845       905       3.95       5.52
## REGION=6 795       736       769       1.41       1.87
##
## Chisq= 27.1 on 5 degrees of freedom, p= 6e-05

Log-Rank - size
surv_diff_delay_size <- survdiff(Surv(proj.delay, INITIATED) ~ size, data = df_fin)
print(surv_diff_delay_size)

## Call:
## survdiff(formula = Surv(proj.delay, INITIATED) ~ size, data = df_fin)
##
##           N Observed Expected (O-E)^2/E (O-E)^2/V
## size=small 888       803       772     1.25516     1.666100
## size=medium 890       811       827     0.29103     0.393744
## size=large 888       814       813     0.00047     0.000633
## size=x-large 891       861       877     0.30047     0.415975
##
## Chisq= 1.9 on 3 degrees of freedom, p= 0.6
```

Cox Proportional Hazards (Cox ph) Model

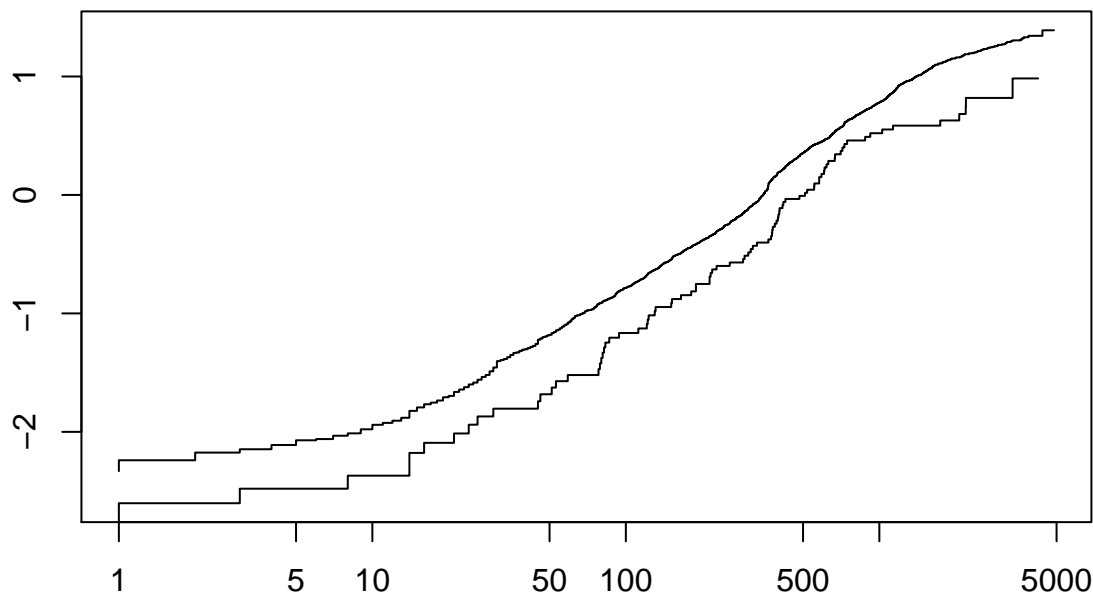
Test Cox proportional hazards assumptions

There are 5 assumptions for using a Cox PH model

1. Non-informative censoring
2. Survival times, t , are independent
3. Hazards are proportional
4. $\ln(\text{HR})$ is a linear function of *numerical* covariates (non in this study)
5. Covariate values don't change over time (eg. changing a treatment or dosage)
6. The baseline hazard (ie. hazard if all covariates were 0) is unspecified

Cox PH - Test proportionality assumption

c-log-log test Litigated?



```
## null device
##      1
```

c-log-log test Appealed?

```
## pdf
##      2
```

c-log-log test Region?

```
## pdf
##      2
```

c-log-log test Size?

```
## pdf
##      2
```

Determining the Cox Proportional Hazards Model

Here we used a likelihood ratios test (LRT) to determine the final form of the Cox proportional hazards model.

```
## Call:  coxph(formula = Surv(proj.delay, INITIATED) ~ 1, data = df_fin)
##
## Null model
##   log likelihood= -23950.14
##   n= 3557
```

Should we add NEPA type?

```
## Call:
## coxph(formula = Surv(proj.delay, INITIATED) ~ NEPA_TYPE, data = df_fin)
##
##               coef exp(coef) se(coef)      z      p
## NEPA_TYPEEEA -0.24888   0.77967  0.03865 -6.439 1.21e-10
## NEPA_TYPEEIS -0.40285   0.66841  0.07563 -5.327 1.00e-07
##
## Likelihood ratio test=62.32  on 2 df, p=2.934e-14
## n= 3557, number of events= 3289
```

```
# Here we use the anova function to complete the LRT test.
anova(cox.cat2, cox.test, test="LRT")
```

```
## Analysis of Deviance Table
## Cox model: response is Surv(proj.delay, INITIATED)
## Model 1: ~ NEPA_TYPE
## Model 2: ~ 1
##   loglik Chisq Df P(>|Chi|)
## 1 -23919
## 2 -23950 62.32  2 2.934e-14 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Yes

Should we include Litigated?

```
## Analysis of Deviance Table
## Cox model: response is Surv(proj.delay, INITIATED)
## Model 1: ~ NEPA_TYPE + LITIGATED
## Model 2: ~ NEPA_TYPE
##   loglik Chisq Df P(>|Chi|)
## 1 -23916
## 2 -23919 6.2551  1  0.01238 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Yes

Should we include Appealed?

```
## Analysis of Deviance Table
## Cox model: response is Surv(proj.delay, INITIATED)
## Model 1: ~ NEPA_TYPE + LITIGATED + APPEALED
## Model 2: ~ NEPA_TYPE + LITIGATED
##   loglik Chisq Df P(>|Chi|)
## 1 -23916
## 2 -23916 0.4222  1  0.5159
```

No

Should we include Region?

```
## Analysis of Deviance Table
## Cox model: response is Surv(proj.delay, INITIATED)
## Model 1: ~ NEPA_TYPE + LITIGATED + REGION
## Model 2: ~ NEPA_TYPE + LITIGATED
## loglik Chisq Df P(>|Chi|)
## 1 -23905
## 2 -23916 22.477 5 0.0004248 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Yes

Should we include size?

```
## Analysis of Deviance Table
## Cox model: response is Surv(proj.delay, INITIATED)
## Model 1: ~ NEPA_TYPE + LITIGATED + REGION + size
## Model 2: ~ NEPA_TYPE + LITIGATED + REGION
## loglik Chisq Df P(>|Chi|)
## 1 -23898
## 2 -23905 12.284 3 0.00647 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

The final Cox proportional hazards model should include all covariates except for appealed.

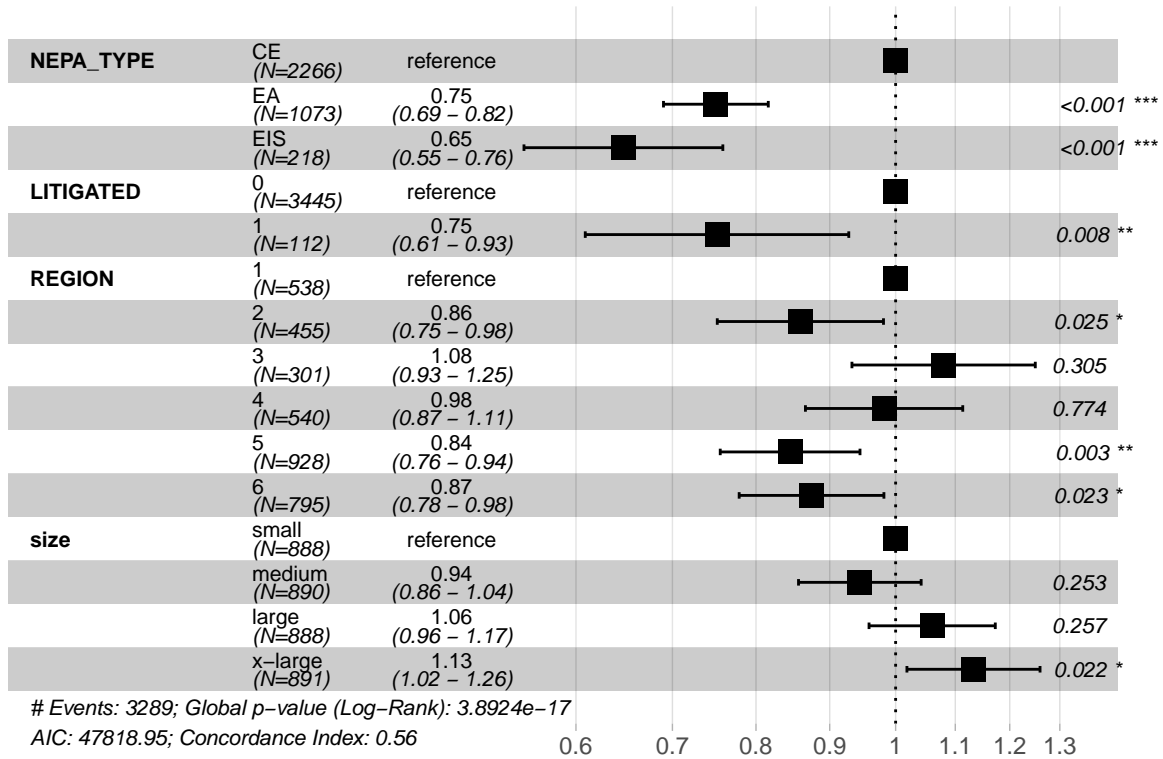
Need to create Cox models using each of the dataframes made with a different region set as the reference.

Create forest plots of Cox proportional hazards models.

Cox PH Model - Hazard Ratios - Region 1 as Ref

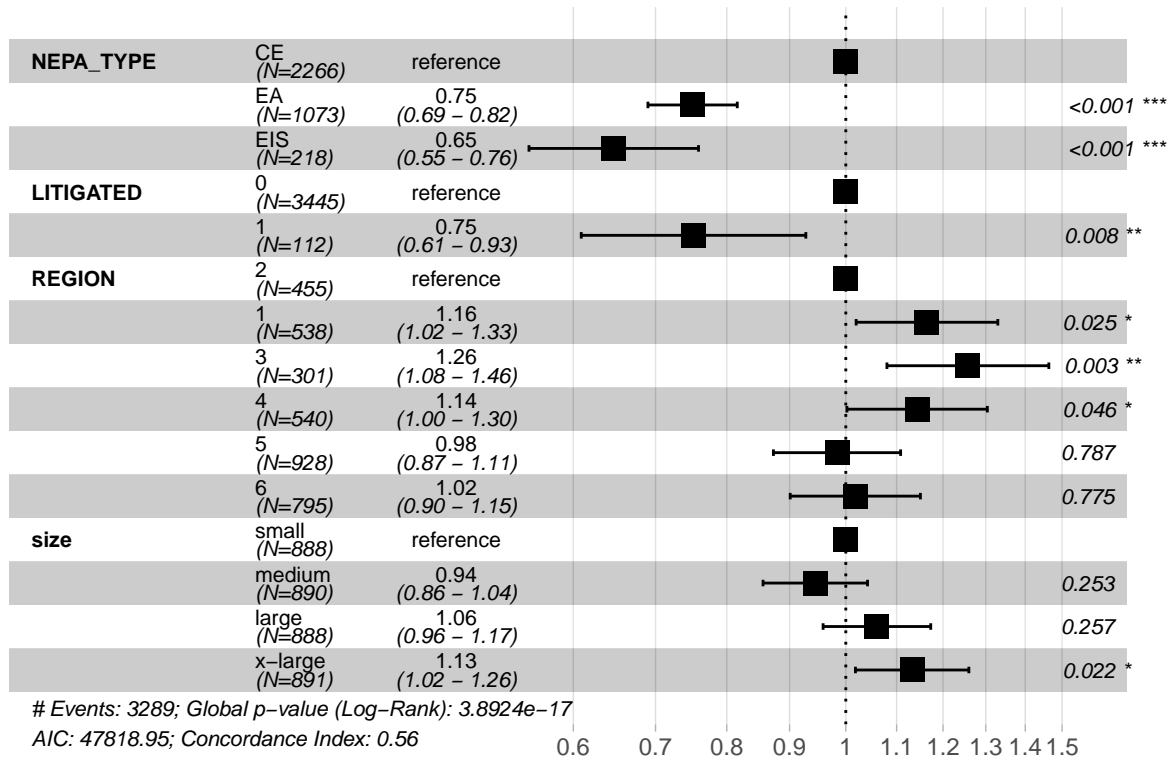
```
## Call:
## coxph(formula = Surv(proj.delay, INITIATED) ~ NEPA_TYPE + LITIGATED +
##       REGION + size, data = df_fin)
##
##               coef exp(coef) se(coef)      z      p
## NEPA_TYPEEEA -0.28722   0.75035  0.04272 -6.722 1.79e-11
## NEPA_TYPEEEIS -0.43501   0.64726  0.08107 -5.366 8.06e-08
## LITIGATED1    -0.28542   0.75170  0.10743 -2.657 0.00789
## REGION2       -0.15222   0.85880  0.06776 -2.246 0.02468
## REGION3        0.07668   1.07970  0.07474  1.026 0.30493
## REGION4       -0.01845   0.98172  0.06413 -0.288 0.77357
## REGION5       -0.16863   0.84482  0.05698 -2.960 0.00308
## REGION6       -0.13439   0.87425  0.05901 -2.277 0.02277
## sizemedium    -0.05720   0.94441  0.04999 -1.144 0.25251
## sizelarge      0.05836   1.06010  0.05146  1.134 0.25677
## sizex-large    0.12441   1.13248  0.05425  2.293 0.02183
##
## Likelihood ratio test=103.3 on 11 df, p=< 2.2e-16
## n= 3557, number of events= 3289
```

Hazard ratio

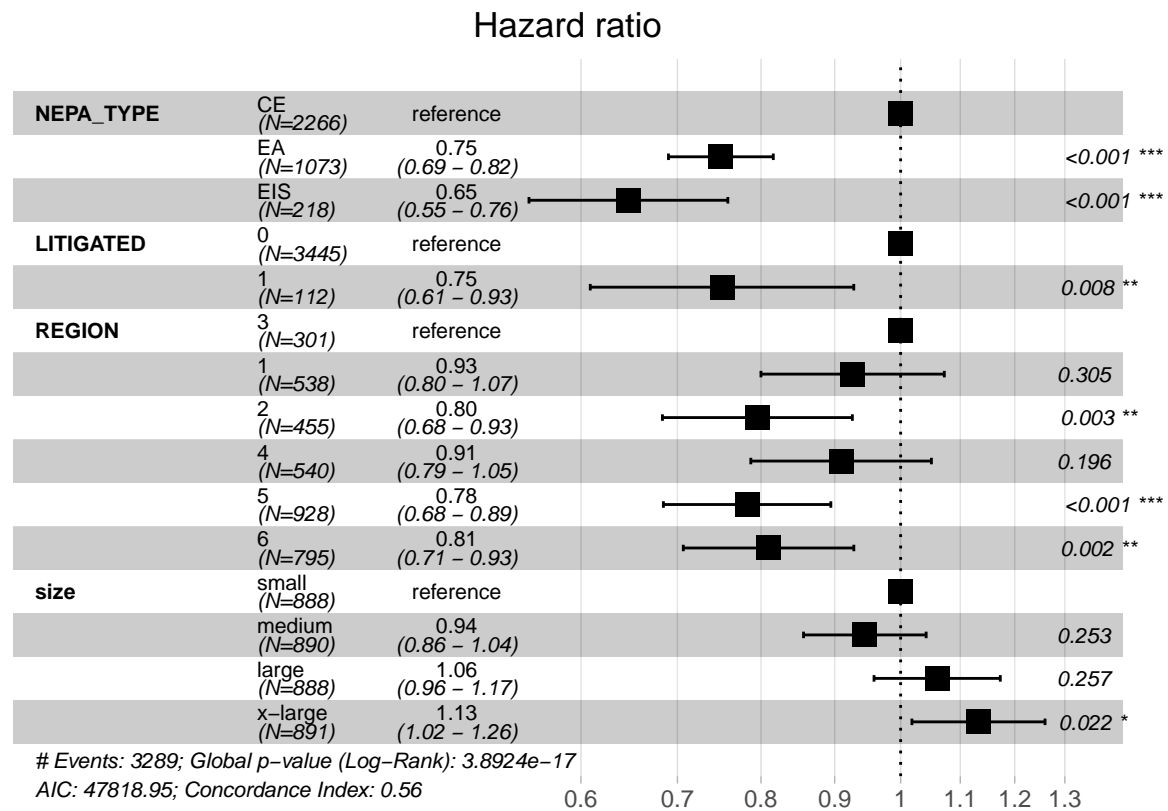


Cox PH Model - Hazard Ratios - Region 2 as Ref

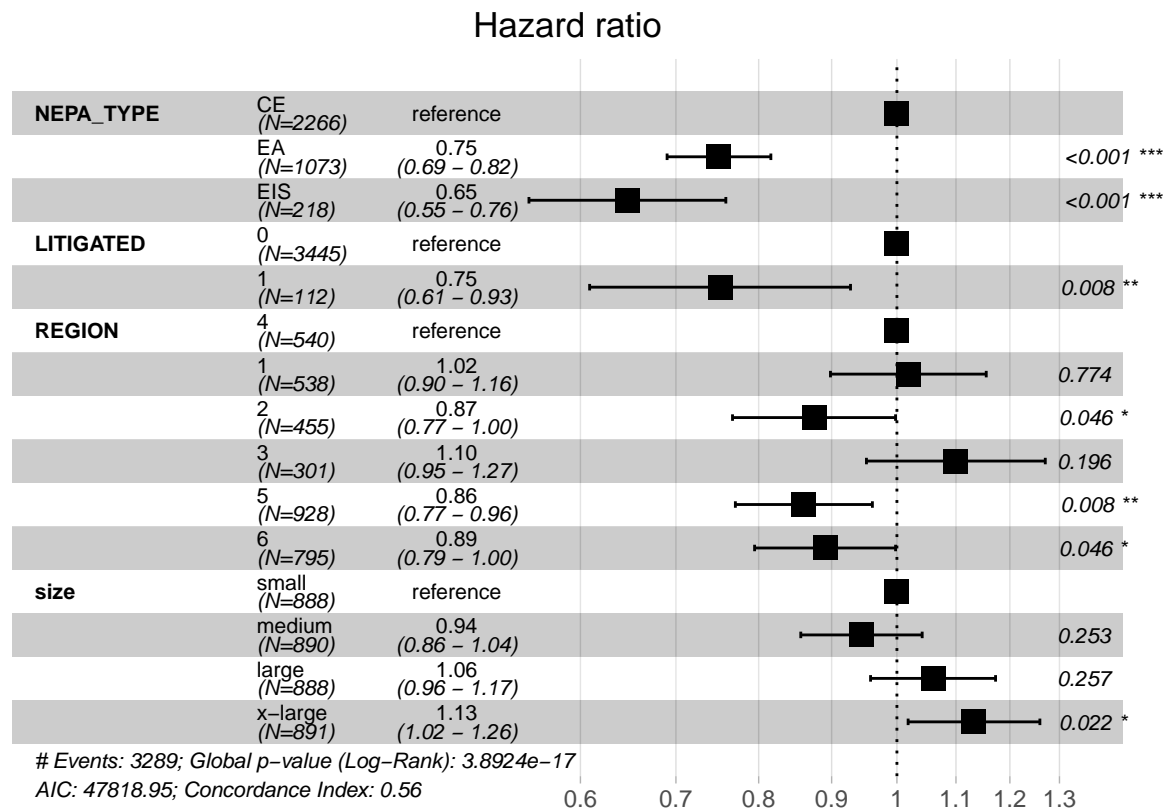
Hazard ratio



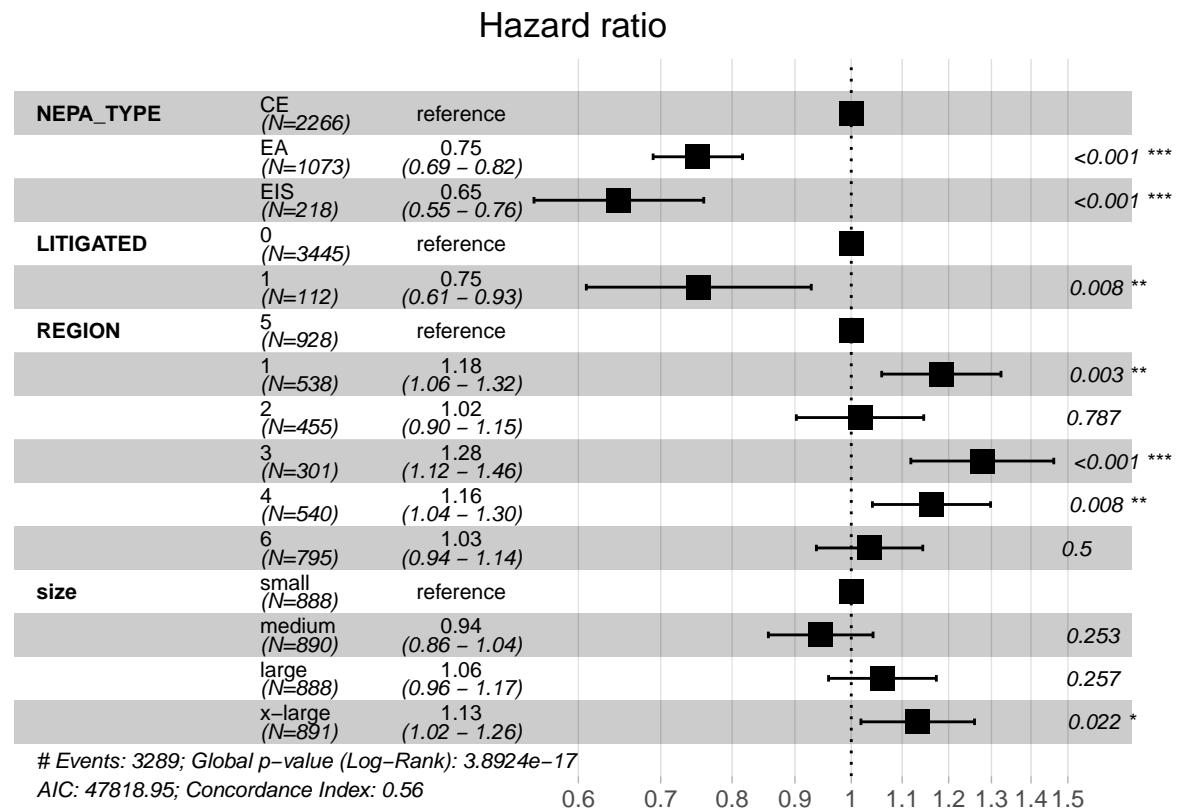
Cox PH Model - Hazard Ratios - Region 3 as Ref



Cox PH Model - Hazard Ratios - Region 4 as Ref



Cox PH Model - Hazard Ratios - Region 5 as Ref



Cox PH Model - Hazard Ratios - Region 6 as Ref

