Replicated Lab 3

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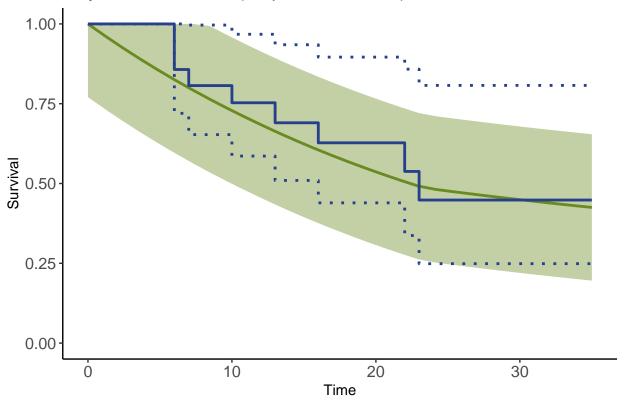
2023-11-28

```
knitr::opts_chunk$set(echo = TRUE)
#Packages
library(tidyr)
library(dplyr)
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
##
       filter, lag
## The following objects are masked from 'package:base':
##
       intersect, setdiff, setequal, union
library(ggplot2)
library(magrittr) # to get %$% operator
##
## Attaching package: 'magrittr'
## The following object is masked from 'package:tidyr':
##
##
       extract
library(survival)
library(KMsurv)
library(BayesSurvival) #our package
library(GGally) #for ggsurv fxn
## Registered S3 method overwritten by 'GGally':
##
    method from
     +.gg
            ggplot2
#Data Setup
data(drug6mp)
#create a Surv object using time=t2 and event=relapse
surv6mp <- as_tibble(drug6mp) %>% arrange(t2) %$% Surv(time=t2, event=relapse)
```

Kaplan-Meier Surivival Curves Here are the Kaplan-Meier survival curves.

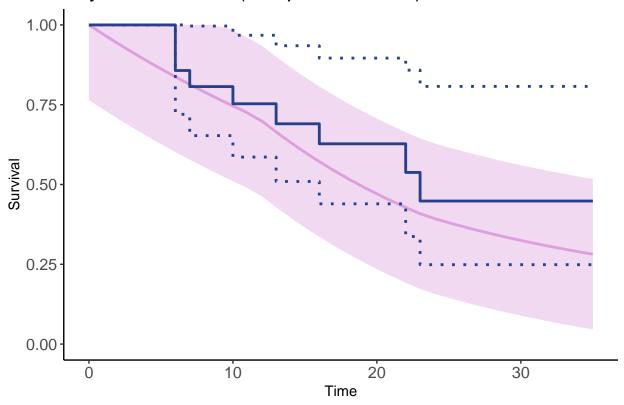
```
#KM Curve from Lab
  #survfit object
 lab_km_surv <- survfit(surv6mp~1, type="kaplan-meier")</pre>
  #dataframe of time and survival
  lab_km <- data.frame(t = lab_km_surv$time, surv = lab_km_surv$surv)</pre>
  #KM Curve
  km = ggsurv(lab_km_surv, main = 'KM Survival Curve From Lab',
              back.white = TRUE, size.est = 1, size.ci = 1,
              surv.col = 'royalblue4')
#Bayesian - Dependent
  #Bayesian Survival object
  bayes_dep <- BayesSurv(drug6mp, "t2", "relapse", prior = 'Dependent')</pre>
  #Plot Bayesian survival curve
  km_plot_bayes_dep <- PlotBayesSurv(bayes_dep, object = "survival",</pre>
               color = 'olivedrab4', legend = FALSE,
               ylab = 'Survival', xlab = 'Time',
               plot.title = 'Bayesian Survival (Dependent Prior) with KM Curve')
  #Add KM curve (+ CI) to Bayesian survival curve
  km_plot_bayes_dep <- km_plot_bayes_dep +</pre>
                       geom_step(data = km$data,aes(x = time, y = surv),
                                  col = 'royalblue4', linewidth = 1) +
                        geom_step(data = km$data, aes(x = time, y = up),
                                  col = 'royalblue4', linewidth = 1,
                                  linetype = 81) +
                        geom_step(data = km$data, aes(x = time, y = low),
                                  col = 'royalblue4', linewidth = 1,
                                  linetype = 81)
 km_plot_bayes_dep
```

Bayesian Survival (Dependent Prior) with KM Curve



```
#Bayesian - Independent
  #Bayesian Survival object
 bayes_indep <- BayesSurv(drug6mp, "t2", "relapse", prior = 'Independent')</pre>
 #Plot Bayesian survival curve
 km_plot_bayes_indep <- PlotBayesSurv(bayes_indep, object = "survival",</pre>
            color = 'plum', legend = FALSE,
            ylab = 'Survival', xlab = 'Time',
            plot.title = 'Bayesian Survival (Independent Prior) with KM Curve')
 #Add KM curve (+ CI) to Bayesian survival curve
 km_plot_bayes_indep <- km_plot_bayes_indep +</pre>
                         geom_step(data = km$data, aes(x = time, y = surv),
                                    col = 'royalblue4', linewidth = 1) +
                         geom_step(data = km$data, aes(x = time, y = up),
                                    col = 'royalblue4', linewidth = 1,
                                    linetype = 81) +
                         geom_step(data = km$data, aes(x = time, y = low),
                                    col = 'royalblue4', linewidth = 1,
                                    linetype = 81)
 km_plot_bayes_indep
```

Bayesian Survival (Independent Prior) with KM Curve



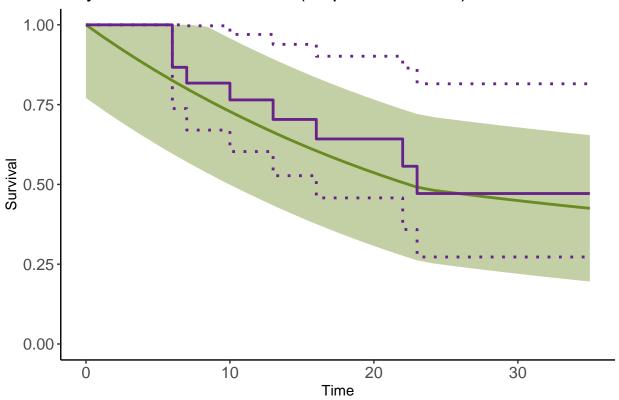
Fleming-Harrington Surivival Curves Here are the Fleming-Harrington survival curves.

```
#FH Curve from Lab
  #survfit object
 lab_fh_surv <- survfit(surv6mp~1, type="fleming-harrington")</pre>
  #dataframe of time and survival
  lab_fh <- data.frame(t = lab_fh_surv$time, surv = lab_fh_surv$surv)</pre>
  #FH Curve
  fh = ggsurv(lab_fh_surv, main = 'FH Survival Curve From Lab',
              back.white = TRUE, size.est = 1, size.ci = 1,
              surv.col = 'darkorchid4')
#Bayesian - Dependent
  #Plot Bayesian survival curve
  fh_plot_bayes_dep <- PlotBayesSurv(bayes_dep, object = "survival",</pre>
        color = 'olivedrab4', legend = FALSE,
        ylab = 'Survival', xlab = 'Time',
        plot.title = 'Bayesian Survival Curve (Dependent Prior) with FH Curve')
  #Add FH curve (+ CI) to Bayesian survival curve
  fh_plot_bayes_dep <- fh_plot_bayes_dep + geom_step(data = fh$data,</pre>
                                                       aes(x = time, y = surv),
                                                           col = 'darkorchid4',
                                                       linewidth = 1) +
                                            geom_step(data = fh$data,
                                                       aes(x = time, y = up),
```

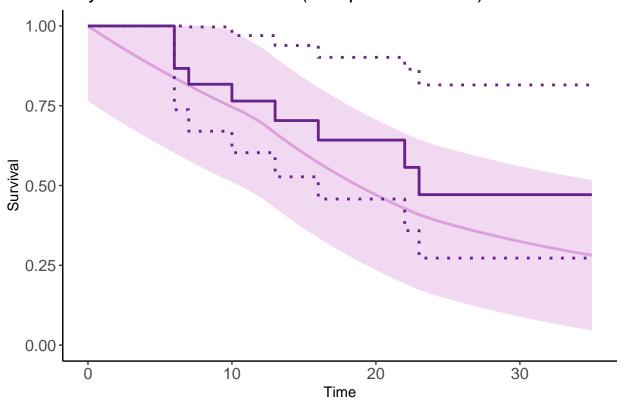
```
col = 'darkorchid4',
    linewidth = 1,
    linetype = 81) +
geom_step(data = fh$data,
    aes(x = time, y = low),
    col = 'darkorchid4',
    linewidth = 1,
    linetype = 81)

fh_plot_bayes_dep
```

Bayesian Survival Curve (Dependent Prior) with FH Curve

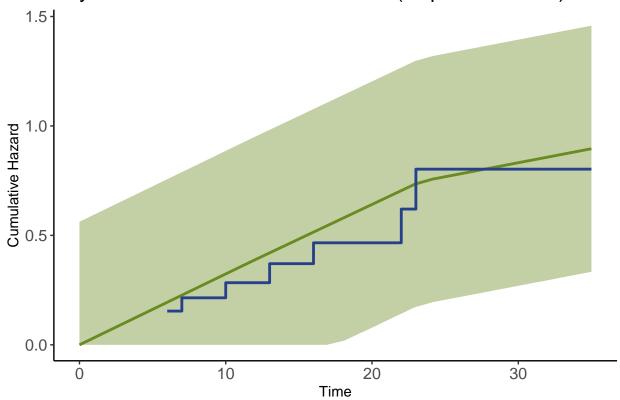


Bayesian Survival Curve (Independent Prior) with FH Curve

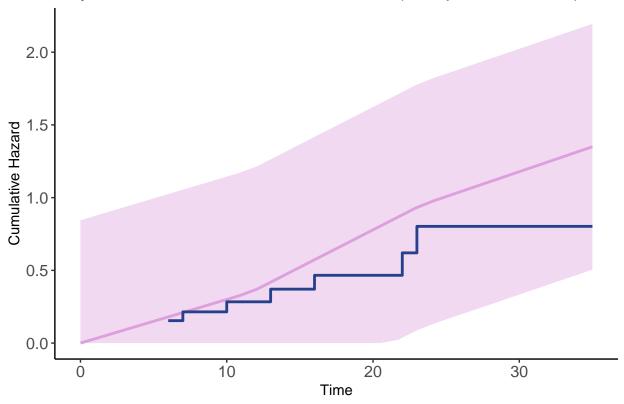


Kaplan-Meier Cumulative Hazard Curves Here are the Kaplan-Meier cumulative hazard curves.

Bayesian Cumulative Hazard Curve (Dependent Prior) with k



Bayesian Cumulative Hazard Curve (Independent Prior) with



Fleming-Harrington Cumulative Hazard Curves Here are the Kaplan-Meier cumulative hazard curves.

```
#FH Cumulative Hazard Curve from Lab
 lab_fh_chaz <- data.frame(t = lab_fh_surv$time, chaz = -log(lab_fh_surv$surv))</pre>
#FH Curve
  fh_chaz = ggplot() + geom_step(data = lab_fh_chaz, aes(x = t, y = chaz),
                                  col = 'royalblue4', linewidth = 1)
#Bayesian - Dependent
  #Plot Bayesian cumulative hazard curve
 fh_plot_bayes_dep <- PlotBayesSurv(bayes_dep, object = "cumhaz",</pre>
color = 'olivedrab4', legend = FALSE,
ylab = 'Cumulative Hazard', xlab = 'Time',
plot.title = 'Dependent Prior')
  #Add FH curve (+ CI) to Bayesian cumulative hazard curve
 fh_plot_bayes_dep <- fh_plot_bayes_dep + geom_step(data = lab_fh_chaz,</pre>
                                                       aes(x = t, y = chaz),
                                                       col = 'darkorchid4',
                                                       linewidth = 1)
  fh_plot_bayes_dep
```

Dependent Prior

