

War Recurrence

Libraries

```
# install.packages("readxl")
# install.packages("tidyverse")
# install.packages("ggplot2")
# install.packages("survminer")
# install.packages("timeROC")
# install.packages("caret")
# install.packages("timeROC")
# install.packages("caret")
# Install and load necessary packages
# install.packages("randomForestSRC")
# install.packages("tidyr")
#install.packages("Synth")

library(Synth)
```

```
##
```

```
## Synth Package: Implements Synthetic Control Methods.
```

```
## See https://web.stanford.edu/~jhain/synthpage.html for additional information.
```

```
library(pec)
```

```
Loading required package: prodlim
```

```
library(expss)
```

```
Loading required package: maditr
```

To aggregate all non-grouping columns: `take_all(mtcars, mean, by = am)`

Attaching package: 'maditr'

The following object is masked from 'package:base':

`sort_by`

```
library(tidyr)
```

Attaching package: 'tidyr'

The following objects are masked from 'package:expss':

`contains, nest`

```
library(caret)
```

Loading required package: ggplot2

Attaching package: 'ggplot2'

The following object is masked from 'package:expss':

`vars`

Loading required package: lattice

Attaching package: 'caret'

The following object is masked from 'package:pec':

`R2`

```
library(stargazer)
```

Please cite as:

Hlavac, Marek (2022). stargazer: Well-Formatted Regression and Summary Statistics Tables.

R package version 5.2.3. <https://CRAN.R-project.org/package=stargazer>

```
library(survival)
```

Attaching package: 'survival'

The following object is masked from 'package:caret':

cluster

```
library(survminer)
```

Loading required package: ggpubr

Attaching package: 'ggpubr'

The following object is masked from 'package:expss':

compare_means

Attaching package: 'survminer'

The following object is masked from 'package:survival':

myeloma

```
library(car)
```

Loading required package: carData

Attaching package: 'car'

The following object is masked from 'package:expss':

recode

```
library(carData)
library(readxl)
library(car)
library(carData)
library(date)
library(readxl)
library(scales)
library(dplyr)
```

Attaching package: 'dplyr'

The following object is masked from 'package:car':

recode

The following objects are masked from 'package:expss':

compute, contains, na_if, recode, vars, where

The following objects are masked from 'package:maditr':

between, coalesce, first, last

The following objects are masked from 'package:stats':

filter, lag

The following objects are masked from 'package:base':

intersect, setdiff, setequal, union

```
library(gridExtra)
```

Attaching package: 'gridExtra'

The following object is masked from 'package:dplyr':

combine

```
library(stargazer)
library(readxl)
library(car)
library(tidyverse)
```

-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --

v forcats	1.0.0	v readr	2.1.5
v lubridate	1.9.3	v stringr	1.5.1
v purrr	1.0.2	v tibble	3.2.1

-- Conflicts ----- tidyverse_conflicts() --

x dplyr::between()	masks	maditr::between()
x dplyr::coalesce()	masks	maditr::coalesce()
x readr::col_factor()	masks	scales::col_factor()
x readr::cols()	masks	maditr::cols()
x gridExtra::combine()	masks	dplyr::combine()
x dplyr::compute()	masks	expss::compute()
x dplyr::contains()	masks	tidyr::contains(), expss::contains()
x purrr::discard()	masks	scales::discard()
x dplyr::filter()	masks	stats::filter()
x dplyr::first()	masks	maditr::first()
x stringr::fixed()	masks	expss::fixed()
x purrr::keep()	masks	expss::keep()
x dplyr::lag()	masks	stats::lag()
x dplyr::last()	masks	maditr::last()
x purrr::lift()	masks	caret::lift()
x purrr::modify()	masks	expss::modify()

```

x purrr::modify_if()    masks expss::modify_if()
x dplyr::na_if()        masks expss::na_if()
x tidyr::nest()         masks expss::nest()
x dplyr::recode()       masks car::recode(), expss::recode()
x stringr::regex()      masks expss::regex()
x purrr::some()         masks car::some()
x purrr::transpose()    masks maditr::transpose()
x dplyr::vars()          masks ggplot2::vars(), expss::vars()
x purrr::when()         masks expss::when()
x dplyr::where()        masks expss::where()
i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become

```

```

library(survival)
library(survminer)
library(carData)
library(readxl)
library(carData)
options(scipen=999)
library(date)
library(readxl)
library(scales)
library(ggplot2)
library(gridExtra)
library(randomForestSRC)

```

randomForestSRC 3.2.3

Type `rfsrc.news()` to see new features, changes, and bug fixes.

Attaching package: 'randomForestSRC'

The following object is masked from 'package:purrr':

partial

```

library(readxl)
library(pec)
library(timeROC)

```

```

library(survivalROC)
library(ggplot2)
library(expss)
library(survival)
library(dplyr)
library(caret)
library(randomForestSRC)
library(readxl)
library(pec)
library(timeROC)
library(survivalROC)
library(ggplot2)
library(expss)
library(survival)
library(dplyr)
library(caret)

```

Data Load and Merge

Load Data

```

episodes <- read_excel("episodes.xlsx")

episodes$polity_scaled<- scale(episodes$p_polity2, center = TRUE, scale = TRUE)
episodes$gov_ter<-ifelse(episodes$wt==1, 1,0)
episodes$gov_war<-ifelse(episodes$wt==2, 1,0)
episodes$ter_war<-ifelse(episodes$wt==3, 1,0)
episodes$coal<- scale(episodes$W4, center = TRUE, scale = TRUE)

episodess <- subset(episodes, intensity_level == 2)
episodest<- subset(episodes, type_of_conflict ==3 )

peaces<- episodes %>% filter(peace== 1)

```

Data merge- recycled

```
# episodes_subset <- select(episodes, conflict_id, outcome, recur_any, recur_side, ps, recur)

#isq_subset <- select(ISQ_ACD, year, isq2015_id, recur)
```

```
#qog_vars <- c(
  # Military and conflict-related variables
  # "wdi_armimp",      # Arms imports (SIPRI trend indicator values)
  # "wvs_fight",      # Willingness to fight for country
  # "wdi_expml",      # Military expenditure (% of GDP)
  # "wdi_expmlge",    # Military expenditure (% of general government expenditure)
  # "bicc_hw",        # Heavy Weapons Index
  # "bicc_gmi",       # Global Militarization Index
  # "atop_ally",      # Member of an Alliance
  # "atop_number",    # Number of Alliances
  # "bicc_milexp",    # Military Expenditure Index
  # "fe_etfra",       # Ethnic Fractionalization
  # "year",
  # "cname",

  # Economic and political variables
  # "al_ethnic2000",  # Ethnic fractionalization index, 2000
  # "al_language2000", # Language fractionalization index, 2000
  # "al_religion2000", # Religious fractionalization index, 2000
  # "fe_cultdiv",     # Cultural diversity index
  # "p_durable",      # Political durability
  # "wdi_gdpcapcur",  # GDP per capita (current US$)
  # "p_polity2"       # Revised polity score
#)
```

```
qog_vars <- c(
  # New variables added
  "ht_colonial",      # Historical colonialism
  "vdem_gender",      # Gender equality index
  "gle_pop",          # Population data
  "gle_gdp",          # GDP data
  "fe_plural",        # Pluralism index
  "vdem_exbribe",     # Executive bribes index
  "vdem_execorr",     # Executive corruption index
  "vdem_exembezz",    # Executive embezzlement index
  "vdem_jucorrdc",    # Judicial corruption index
  "vdem_libdem",      # Liberal democracy index
```



```

"wdi_pop",          # Population data from World Development Indicators
"cname",            # Country name
"year"              # Year
)

#episodes<-merge(episodes, selected_qog_vars,by=c("cname","year"), all.x= TRUE)
#eps<-merge(episodes, NewWmeasure,by=c("country_name","year"), all.x= TRUE)

#epid<-merge(episodes,tem_sean,by=c("conflict_id","end_year"), all=TRUE)

#ACD_Sean_1000<-merge(episodes_1000,ISQ_ACD,by=c("conflict_id","year"), all=TRUE)

#ACD_Sean_1000<-merge(episodes_1000merge,ISQ_ACD_Merge,by=c("isq2015_id","year"), all=TRUE)

#ep_qog<-merge(episodes,qog_std_ts_jan23,by=c("cname","year"))
#ep<-merge(copy,ep_qog,by=c("conflict_id","end_year"), all=TRUE)
#clean_ep<-ep[, sapply(ep, function(col) length(na.omit(col))) >150]

#dur_acd<-merge(duration,uclp_brd_dyadic_221 ,by=c("cname", "year"))

#epidd$log_gdp<-log(epidd$wdi_gdpcapcur)

#fromqog= subset(qog_std_ts_jan23, select =c(cname, year,fe_etfra) )
#epid<-merge(episodes, fromqog,by=c("cname","year"))
#epidd<-merge(copy, epid,by=c("conflict_id","end_year"))

#episodes$yearr<- as.Date(episodes$start_date2)
#episodes$yea<-date.mdy(episodes$yearr)$yea
#episodes$yea<-episodes$yea+10
#episodes$end<- as.Date(episodes$ep_end_date)
#episodes$en<-date.mdy(episodes$end)$en
#episodes$en<-episodes$en+10

#X= subset(clean_o, select = -c(recur_a, conflict_id, end_year, cname, year, start_year, loc

```

```

#Y=clean_o$recur_a
#clean_ep<-ep[, sapply(ep, function(col) length(na.omit(col))) >150]
#clean_o<-na.omit(clean_ep)

#episodes$peace<-ifelse(episodes$outcome==1, 1,0)
#episodes$cease<-ifelse(episodes$outcome==2, 1,0)
#episodes$govvic<-ifelse(episodes$outcome==3, 1,0)
#episodes$rebvic<-ifelse(episodes$outcome==4, 1,0)
#episodes$lowac<-ifelse(episodes$outcome==5, 1,0)
#episodes$dis<-ifelse(episodes$outcome==6, 1,0)
#episodes$lowcease<-Recode(episodes$outcome, "1=0; 2=1; 3=0; 4=0; 5=1; 6=0")
#episodes$log_dur<-log(episodes$duration)

```

EDA

Distribution of Outcome

```

episodes$out_ <- as.character(episodes$outcome) # Convert factor to character if necessary

# Define a mapping for the outcomes
outcome_labels <- c("1" = "Peace Agreement",
                    "2" = "Ceasefire",
                    "3" = "Government Victory",
                    "4" = "Rebel Victory",
                    "5" = "Low Activity",
                    "6" = "Actors Ceases to Exist")

# Replace the coded values with the corresponding labels
episodes$out_ <- sapply(episodes$out_, function(x) outcome_labels[x])

# Replace NA values with "Ongoing Episodes"
episodes$out_[is.na(episodes$out_)] <- "Ongoing Episodes"

# Convert the 'outs' column back to factor and set the desired order
episodes$out_ <- factor(episodes$out_, levels = c("Peace Agreement", "Ceasefire", "Government Victory", "Rebel Victory", "Low Activity", "Actors Ceases to Exist"))

# Create the bar plot
outcome_freq <- ggplot(episodes, aes(x = reorder(outs, outs, function(x) -length(x)))) +
  geom_bar(fill = "blue", color = "black", alpha = 0.7) +
  labs(title = "Frequency of All Outcomes",

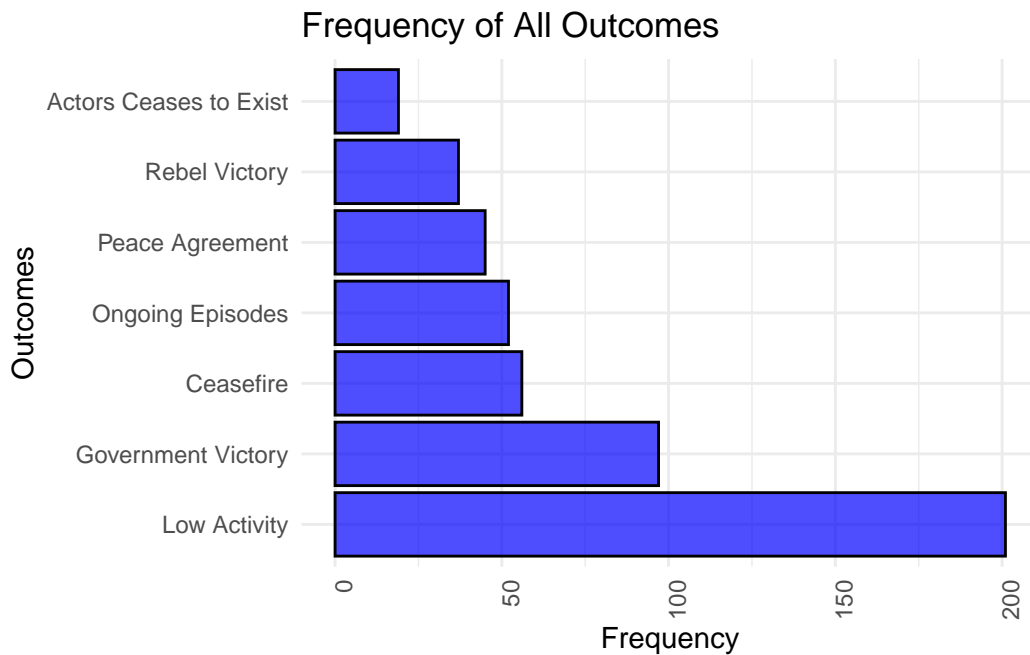
```

```

    x = "Outcomes",
    y = "Frequency") +
  theme_minimal() +
  coord_flip() +
  theme(axis.text.x = element_text(angle = 90, hjust = 1)) # Rotate x-axis labels for better

# Display the plot
print(outcome_freq)

```



Cross-Tab Viz

```

# Filter out rows with NA in the recurr column
episodes |>
  filter(!is.na(recur_any))

```

A tibble: 455 x 125

	episode_id	cname	year	cease	outcome	country_name	conflict_id	start_year
	<dbl>	<chr>	<dbl>	<dbl>	<dbl>	<chr>	<dbl>	<dbl>
1	1	Bolivia (~	1947	0	4	Bolivia	200	1946
2	2	Bolivia (~	1950	0	3	Bolivia	200	1949

3	3	Bolivia (~ 1953	0	4	Bolivia	200	1952
4	4	Bolivia (~ 1968	0	3	Bolivia	200	1967
5	5	China 1950	0	4	China	202	1946
6	6	Greece 1950	0	3	Greece	203	1946
7	7	Iran (Isl~ 1947	0	3	Iran	205	1946
8	8	Iran (Isl~ 1969	0	5	Iran	205	1966
9	9	Iran (Isl~ 1989	0	5	Iran	205	1979
10	10	Iran (Isl~ 1991	0	5	Iran	205	1990

i 445 more rows

i 117 more variables: end_year <dbl>, peace_cease <dbl>,
 # peace_cease_notes <chr>, wdi_expmilge <dbl>, wdi_armimp <dbl>,
 # wdi_expmil <dbl>, bicc_hw <dbl>, bicc_gmi <dbl>, atop_ally <dbl>,
 # atop_number <dbl>, bicc_milexp <dbl>, fe_etfra <dbl>, al_ethnic2000 <dbl>,
 # al_language2000 <dbl>, al_religion2000 <dbl>, fe_cultdiv <dbl>,
 # p_durable <dbl>, wdi_gdpcapcur <dbl>, p_polity2 <dbl>, pko_u <dbl>, ...

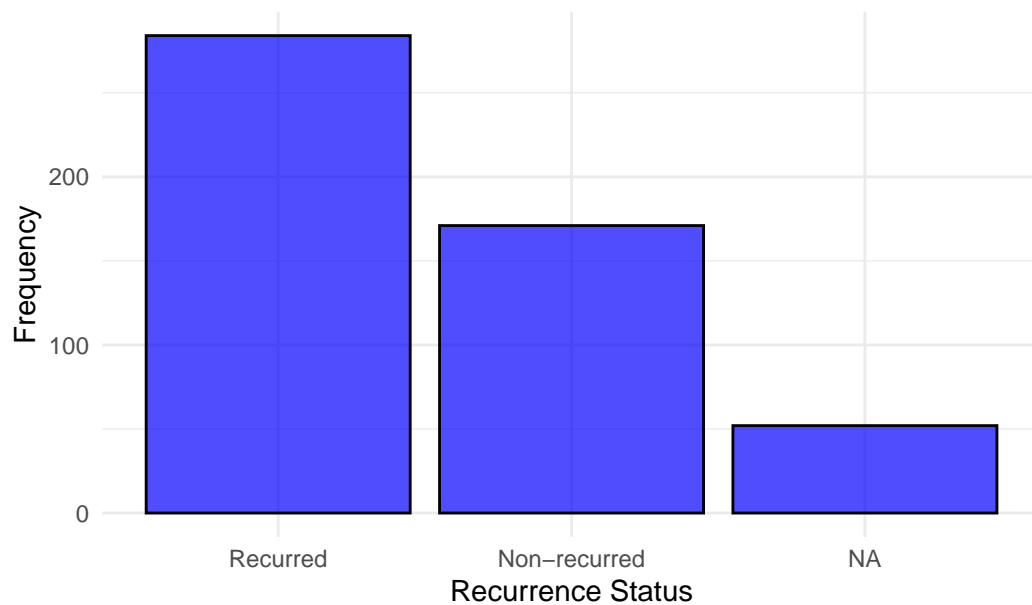
Recode the factor levels

```
episodes$recur0 <- recode_factor(episodes$recur_any, '1'="Recurred", '0'="Non-recurred")
```

Plot the bar chart

```
ggplot(episodes, aes(x=recur0)) +  
  geom_bar(fill="blue", color="black", alpha=0.7) +  
  labs(title="Distribution of War Recurrence Variable Based on ACD Conflict ID",  
        x="Recurrence Status",  
        y="Frequency") +  
  theme_minimal()
```

Distribution of War Recurrence Variable Based on ACD Conflic

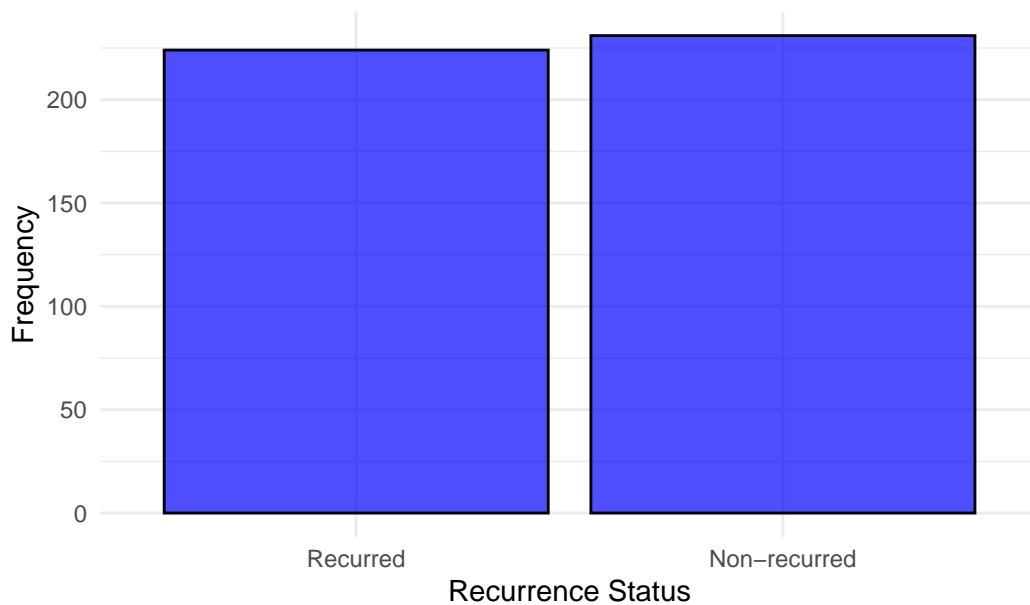


```
# Filter out rows with NA in the recurr column
episodes <- episodes %>% filter(!is.na(recur_side))

# Recode the factor levels
episodes$recur1 <- recode_factor(episodes$recur_side, '1'="Recurred", '0'="Non-recurred")

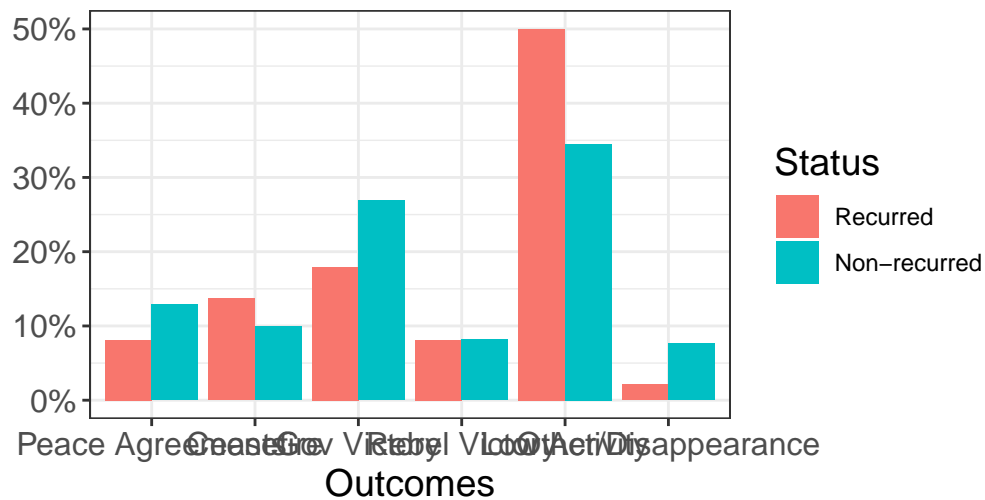
# Plot the bar chart
ggplot(episodes, aes(x=recur1)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Distribution of War Recurrence Variable Based on Sufficient Linkage",
       x="Recurrence Status",
       y="Frequency") +
  theme_minimal()
```

Distribution of War Recurrence Variable Based on Sufficient Li



```
recur_outcome<- table(episodes$recur0, episodes$outcome)
recur_outcome_p<- prop.table(recur_outcome, 1)
recur_outcome_df <- as.data.frame(recur_outcome_p)
names(recur_outcome_df) <- c("recur0", "outcome", "Frequency")
recur_outcome_df$outcome<-recode_factor(recur_outcome_df$outcome, '1'="Peace Agreements", '2'="Peace Agreements")
recur_by_outcome<-ggplot(recur_outcome_df, aes(x=outcome, y=Frequency, fill=recur0)) + geom_bar()
recur_by_outcome <- recur_by_outcome +
  scale_y_continuous(label=percent) +
  labs(title="ACD Conflict ID-Based Recurrence By Termination Outcomes",
        caption="",
        subtitle="",
        x="Outcomes", y="", fill="Status")+
  theme_bw() +
  theme(title=element_text(size=14), axis.text=element_text(size=12))
recur_by_outcome
```

ACD Conflict ID–Based Recurrence By Termination



```
# Create a frequency table
recur_outcome <- table(episodes$recur1, episodes$outcome)

# Convert the table to a data frame for ggplot
recur_outcome_df <- as.data.frame(recur_outcome)
names(recur_outcome_df) <- c("recur1", "outcome", "Frequency")

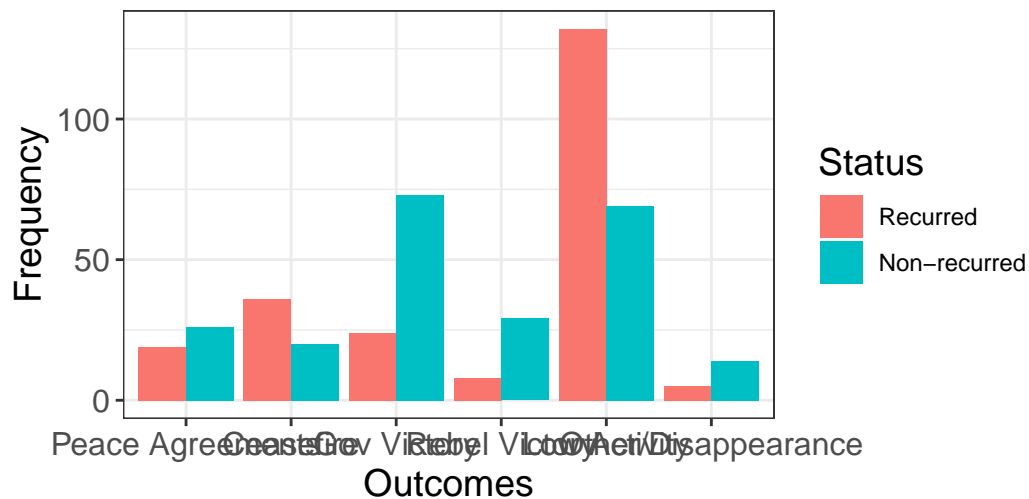
# Recode the outcome factor with descriptive names
recur_outcome_df$outcome <- recode_factor(recur_outcome_df$outcome,
                                          '1' = "Peace Agreements",
                                          '2' = "Ceasefire",
                                          '3' = "Gov Victory",
                                          '4' = "Rebel Victory",
                                          '5' = "Low Activity",
                                          '6' = "Other/Disappearance")

# Plot the bar chart using frequencies
recur_by_outcome <- ggplot(recur_outcome_df, aes(x = outcome, y = Frequency, fill = recur1))
  geom_col(position = "dodge") +
  labs(title = "Sufficient Linkage-Based Recurrence By Termination Outcomes",
       caption = "",
       subtitle = "",
       x = "Outcomes", y = "Frequency", fill = "Status") +
```

```
theme_bw() +
  theme(title = element_text(size = 14), axis.text = element_text(size = 12))

recur_by_outcome
```

Sufficient Linkage-Based Recurrence By Term



```
# Recode the outcome variable for better readability
episodes$outcome_label <- recode_factor(episodes$outcome,
                                         '1' = "Peace Agreements",
                                         '2' = "Ceasefire",
                                         '3' = "Gov Victory",
                                         '4' = "Rebel Victory",
                                         '5' = "Low Activity",
                                         '6' = "Other/Disappearance")

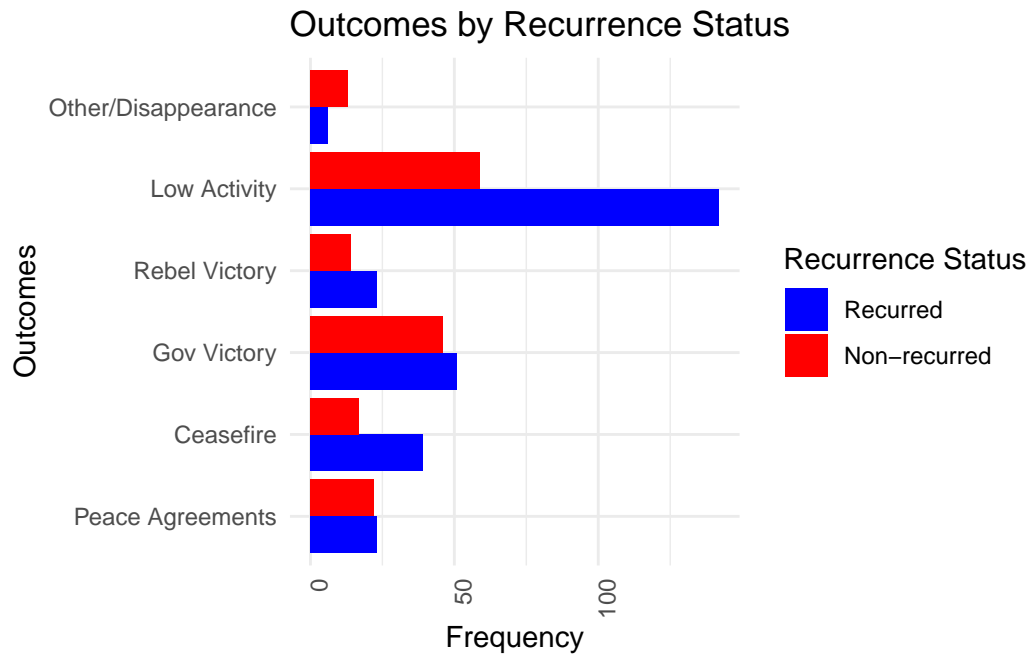
# Bar chart of the outcome variable separated by recur0
outcome_by_recur <- ggplot(episodes, aes(x=outcome_label, fill=factor(recur0))) +
  geom_bar(position="dodge") +
  labs(title="Outcomes by Recurrence Status",
       x="Outcomes",
       y="Frequency",
       fill="Recurrence Status") +
  scale_fill_manual(values=c("blue", "red")) + # Use manual colors if needed
```



```

theme_minimal() +
coord_flip()+
theme(axis.text.x = element_text(angle = 90, hjust = 1)) # Rotate x-axis labels for better
outcome_by_recur

```

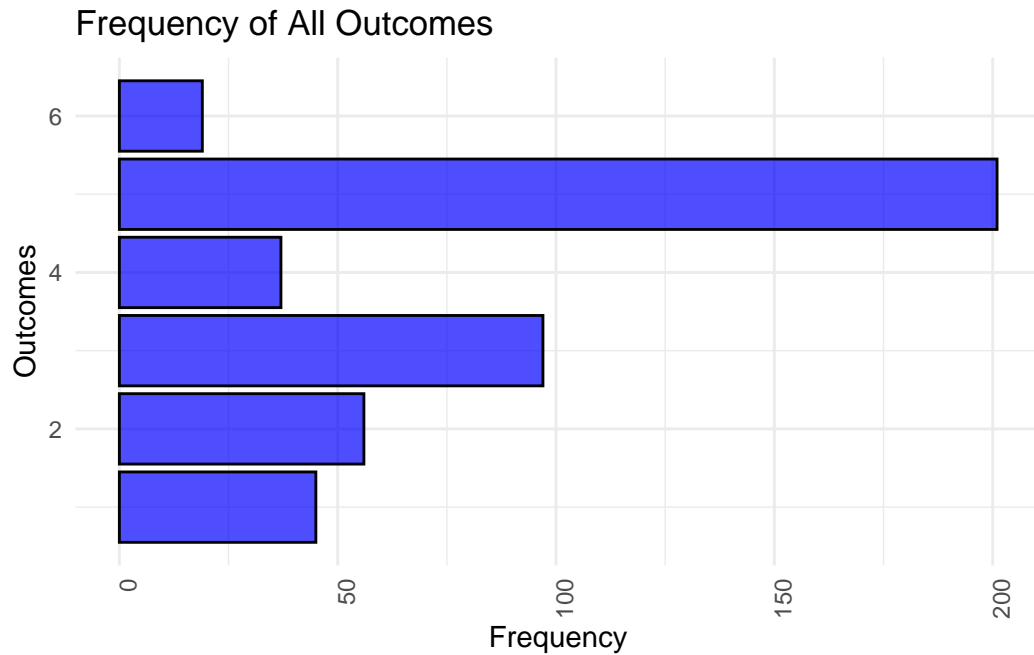


Cross-Tab Viz1

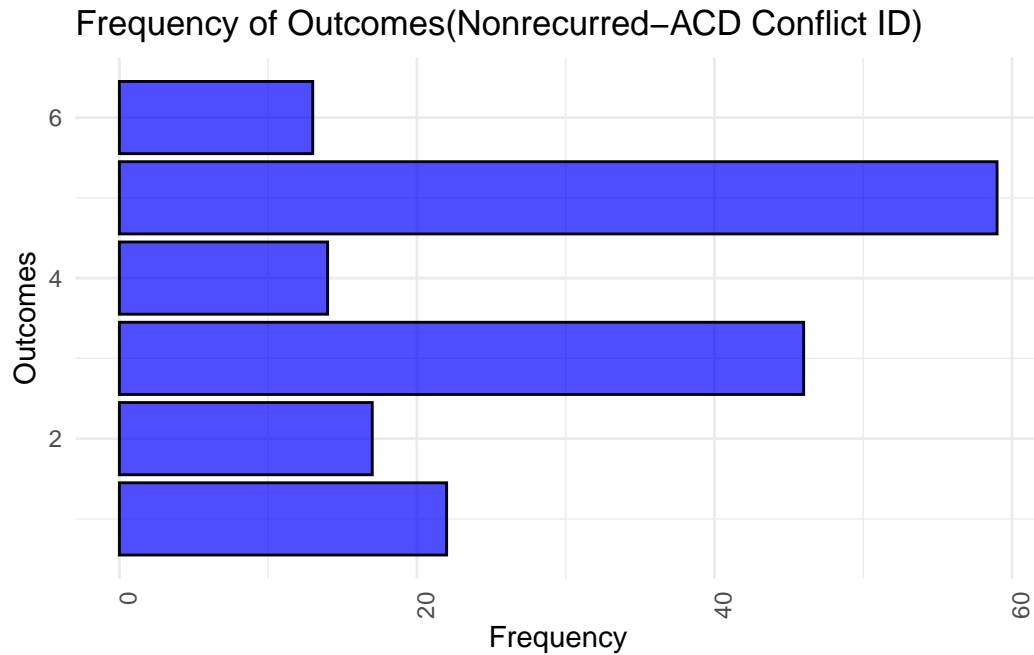
```

outcome_freq <- ggplot(episodes, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of All Outcomes",
        x="Outcomes",
        y="Frequency") +
  theme_minimal() +
  coord_flip()+
  theme(axis.text.x = element_text(angle = 90, hjust = 1)) # Rotate x-axis labels for better
outcome_freq

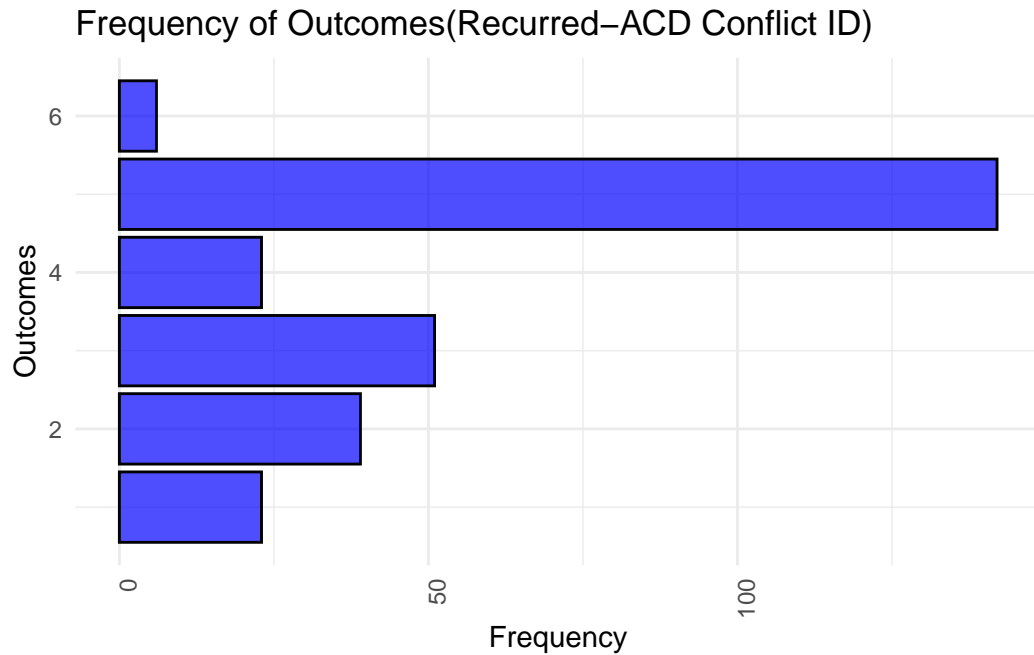
```



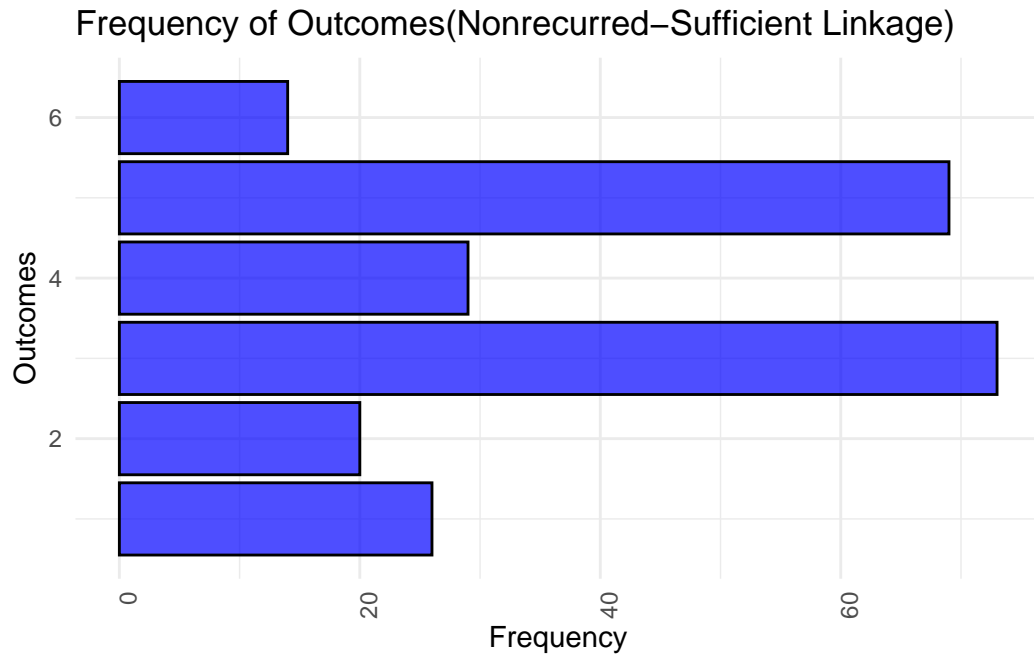
```
# Bar chart of the outcome variable without considering recur0
recur00_data<-subset(episodes, recur0 == "Non-recurred")
outcome_freq <- ggplot(recur00_data, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of Outcomes(Nonrecurred-ACD Conflict ID)",
       x="Outcomes",
       y="Frequency") +
  theme_minimal() +
  coord_flip()+
  theme(axis.text.x = element_text(angle = 90, hjust = 1)) # Rotate x-axis labels for better
outcome_freq
```



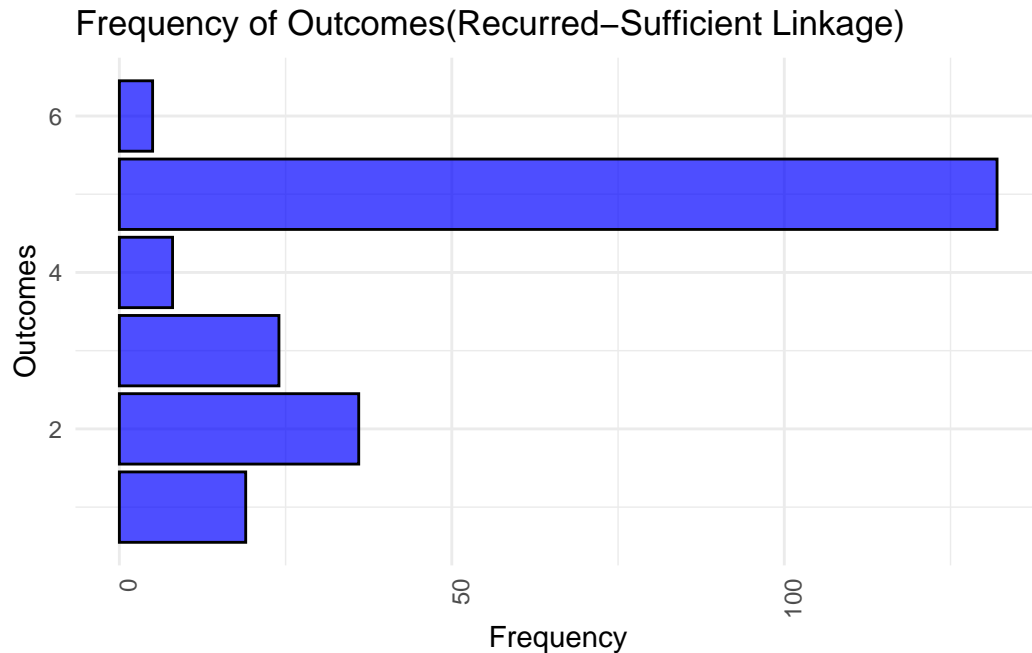
```
# Bar chart of the outcome variable without considering recur0
recur01_data<-subset(episodes, recur0 == "Recurred")
outcome_freq <- ggplot(recur01_data, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of Outcomes(Recurred-ACD Conflict ID)",
        x="Outcomes",
        y="Frequency") +
  theme_minimal() +
  coord_flip()+
  theme(axis.text.x = element_text(angle = 90, hjust = 1)) # Rotate x-axis labels for better
outcome_freq
```



```
# Bar chart of the outcome variable without considering recur0
recur10_data<-subset(episodes, recur1 == "Non-recurred")
outcome_freq <- ggplot(recur10_data, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of Outcomes(Nonrecurred-Sufficient Linkage)",
        x="Outcomes",
        y="Frequency") +
  theme_minimal() +
  coord_flip()+
  theme(axis.text.x = element_text(angle = 90, hjust = 1)) # Rotate x-axis labels for better
outcome_freq
```



```
# Bar chart of the outcome variable without considering recur0
recur11_data<-subset(episodes, recur1 == "Recurred")
outcome_freq <- ggplot(recur11_data, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of Outcomes(Recurred-Sufficient Linkage)",
        x="Outcomes",
        y="Frequency") +
  theme_minimal() +
  coord_flip()+
  theme(axis.text.x = element_text(angle = 90, hjust = 1)) # Rotate x-axis labels for better
outcome_freq
```



```
# Prepare the first plot (Non-recurred - ACD Conflict ID)
recur00_data <- subset(episodes, recur0 == "Non-recurred")
outcome_freq00 <- ggplot(recur00_data, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of Outcomes (Non-recurred - ACD Conflict ID)",
       x="Outcomes", y="Frequency") +
  theme_minimal() + coord_flip() +
  theme(axis.text.x = element_text(angle = 90, hjust = 1))

# Prepare the second plot (Recurred - ACD Conflict ID)
recur01_data <- subset(episodes, recur0 == "Recurred")
outcome_freq01 <- ggplot(recur01_data, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of Outcomes (Recurred - ACD Conflict ID)",
       x="Outcomes", y="Frequency") +
  theme_minimal() + coord_flip() +
  theme(axis.text.x = element_text(angle = 90, hjust = 1), axis.text.y = element_blank(), ax

# Prepare the third plot (Non-recurred - Sufficient Linkage)
recur10_data <- subset(episodes, recur1 == "Non-recurred")
outcome_freq10 <- ggplot(recur10_data, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of Outcomes (Non-recurred - Sufficient Linkage)",
```

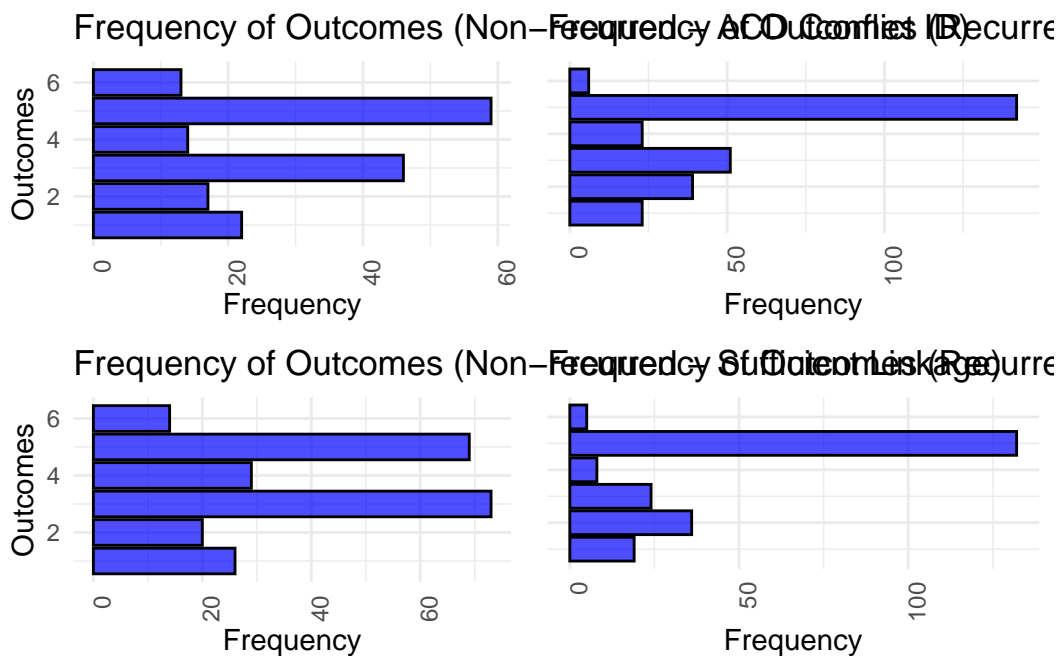
```

    x="Outcomes", y="Frequency") +
  theme_minimal() + coord_flip() +
  theme(axis.text.x = element_text(angle = 90, hjust = 1))

# Prepare the fourth plot (Recurred - Sufficient Linkage)
recur11_data <- subset(episodes, recur1 == "Recurred")
outcome_freq11 <- ggplot(recur11_data, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of Outcomes (Recurred - Sufficient Linkage)",
    x="Outcomes", y="Frequency") +
  theme_minimal() + coord_flip() +
  theme(axis.text.x = element_text(angle = 90, hjust = 1), axis.text.y = element_blank(), ax

# Combine all four plots into one
combined_plot <- grid.arrange(outcome_freq00, outcome_freq01, outcome_freq10, outcome_freq11

```



```
print(combined_plot)
```

```

TableGrob (2 x 2) "arrange": 4 grobs
  z    cells  name      grob
1 1 (1-1,1-1) arrange gtable[layout]
2 2 (1-1,2-2) arrange gtable[layout]

```

```
3 3 (2-2,1-1) arrange gtable[layout]
4 4 (2-2,2-2) arrange gtable[layout]
```

```
# Calculate the maximum frequency across all subsets
max_freq <- max(
  table(episodes$recur0, episodes$outcome),
  table(episodes$recur1, episodes$outcome)
)

# Prepare the first plot (Non-recurred - ACD Conflict ID)
recur00_data <- subset(episodes, recur0 == "Non-recurred")
outcome_freq00 <- ggplot(recur00_data, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of Outcomes (Non-recurred - ACD Conflict ID)",
       x="Outcomes", y="Frequency") +
  theme_minimal() +
  coord_flip() +
  theme(axis.text.x = element_text(angle = 90, hjust = 1)) +
  ylim(c(0, max_freq))

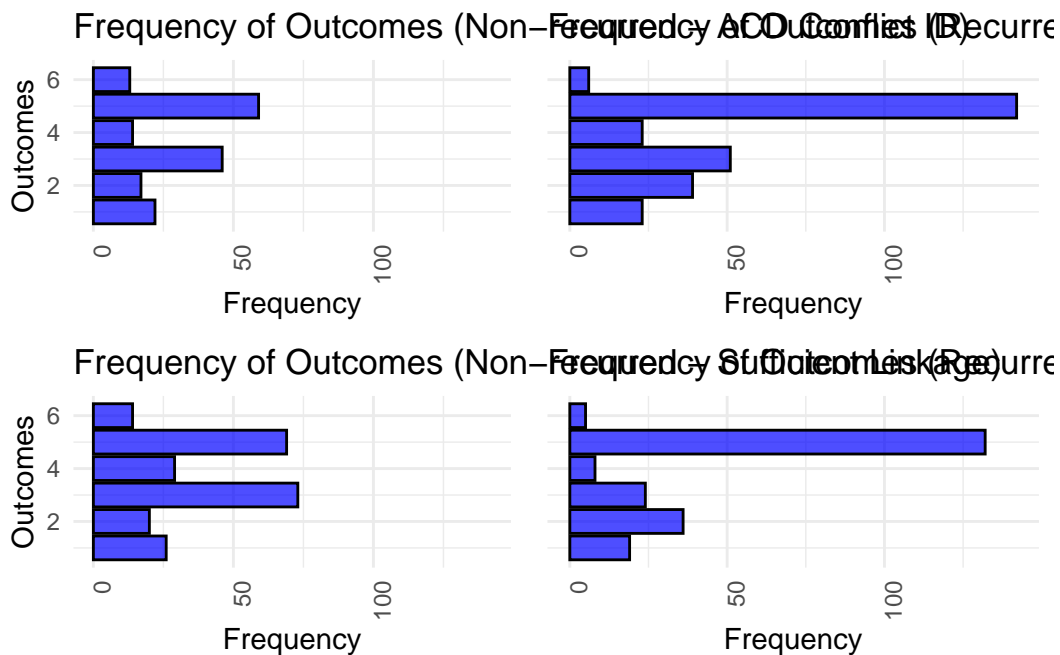
# Prepare the second plot (Recurred - ACD Conflict ID)
recur01_data <- subset(episodes, recur0 == "Recurred")
outcome_freq01 <- ggplot(recur01_data, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of Outcomes (Recurred - ACD Conflict ID)",
       x="Outcomes", y="Frequency") +
  theme_minimal() +
  coord_flip() +
  theme(axis.text.x = element_text(angle = 90, hjust = 1), axis.text.y = element_blank(), ax
  ylim(c(0, max_freq))

# Prepare the third plot (Non-recurred - Sufficient Linkage)
recur10_data <- subset(episodes, recur1 == "Non-recurred")
outcome_freq10 <- ggplot(recur10_data, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of Outcomes (Non-recurred - Sufficient Linkage)",
       x="Outcomes", y="Frequency") +
  theme_minimal() +
  coord_flip() +
  theme(axis.text.x = element_text(angle = 90, hjust = 1)) +
  ylim(c(0, max_freq))
```



```
# Prepare the fourth plot (Recurred - Sufficient Linkage)
recur11_data <- subset(episodes, recur1 == "Recurred")
outcome_freq11 <- ggplot(recur11_data, aes(x=outcome)) +
  geom_bar(fill="blue", color="black", alpha=0.7) +
  labs(title="Frequency of Outcomes (Recurred - Sufficient Linkage)",
       x="Outcomes", y="Frequency") +
  theme_minimal() +
  coord_flip() +
  theme(axis.text.x = element_text(angle = 90, hjust = 1), axis.text.y = element_blank(), ax
  ylim(c(0, max_freq))

# Combine all four plots into one
combined_plot <- grid.arrange(outcome_freq00, outcome_freq01, outcome_freq10, outcome_freq11
```



```
print(combined_plot)
```

```
TableGrob (2 x 2) "arrange": 4 grobs
  z   cells  name      grob
1 1 (1-1,1-1) arrange gtable[layout]
2 2 (1-1,2-2) arrange gtable[layout]
3 3 (2-2,1-1) arrange gtable[layout]
4 4 (2-2,2-2) arrange gtable[layout]
```

Kaplan-Meier survival curves

```
# Create a new dataframe from episodes

new_episodes <- episodes
new_episodes<-new_episodes[, c("outcome", "recur_side", "recur_any", "time_to_recur", "time_1
new_episodes<-na.omit(new_episodes)

#Convert the numerical outcome variable to a factor with appropriate labels
new_episodes$outcome <- factor(new_episodes$outcome, levels = c(1, 2, 3, 4, 5, 6),
                              labels = c("Peace Agreement", "Ceasefire", "Gov Victory", "Rel

# Create new variables for time in years
new_episodes$time_to_recur_any_years <- new_episodes$time_to_recur_any / 365.25
new_episodes$time_to_recur_years <- new_episodes$time_to_recur / 365.25

# Trim follow-up time to 20,000 days (approximately 54.8 years), and create corresponding va
new_episodes$time_to_recur_any_years <- pmin(new_episodes$time_to_recur_any_years, 20000 / 36
new_episodes$time_to_recur_years <- pmin(new_episodes$time_to_recur_years, 20000 / 365.25)

# Fit the Kaplan-Meier survival curves for first plot (Any Recurrence)
km_fit_outcome_any <- survfit(Surv(time_to_recur_any_years, recur_any) ~ outcome, data = new

# Plot the Kaplan-Meier survival curves with percentages in the risk table (Any Recurrence)
km_plot_any <- ggsurvplot(km_fit_outcome_any, data = new_episodes,
                          pval = TRUE, conf.int = FALSE,
                          risk.table = TRUE, risk.table.col = "strata",
                          risk.table.y.text.col = TRUE,
                          risk.table.height = 0.25,
                          risk.table.title = "Number at Risk",
                          xlim = c(0, 35), # Limit x-axis to 35 years
                          break.time.by = 5, # Break x-axis every 5 years
                          ggtheme = theme_classic() + # Use classic theme
                            theme(panel.grid.major = element_line(color = "gray", si
                              panel.grid.minor = element_line(color = "lightgray",
                          palette = c("red", "blue", "green", "purple", "orange", "brown"),
                          title = "KM Survival Curves for Outcomes (Any Recurrence)",
                          xlab = "Time (Years)", ylab = "Survival Probability",
                          tables.theme = theme_classic()) # Use classic theme for the risk t
```

Warning: The `size` argument of `element_line()` is deprecated as of ggplot2 3.4.0.

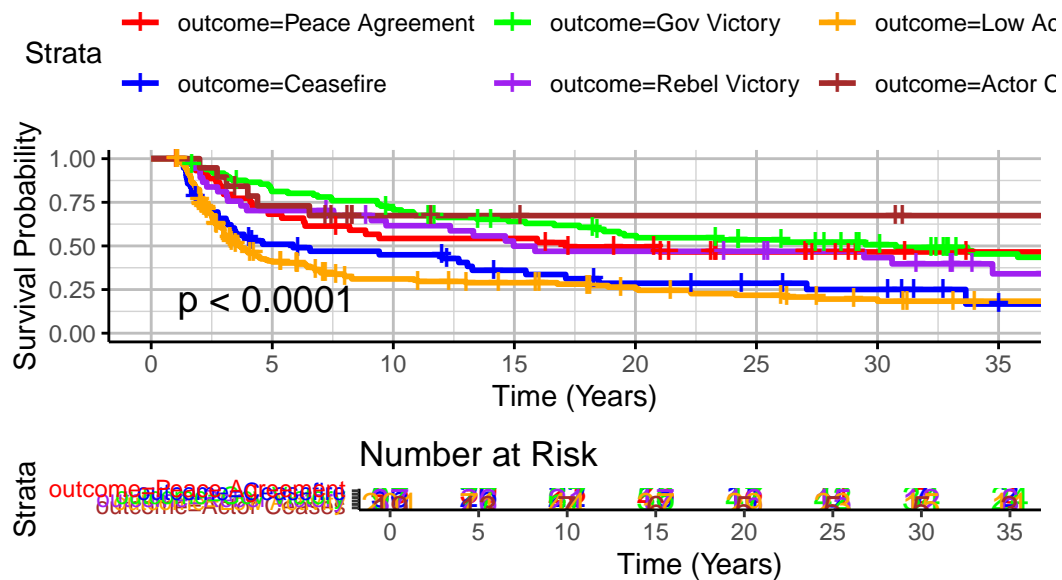
i Please use the ``linewidth`` argument instead.

```
# Fit the Kaplan-Meier survival curves for the second plot (Side Recurrence)
km_fit_outcome_side <- survfit(Surv(time_to_recur_years, recur_side) ~ outcome, data = new_episodes)

# Plot the Kaplan-Meier survival curves with percentages in the risk table for the second plot
km_plot_side <- ggsurvplot(km_fit_outcome_side, data = new_episodes,
                           pval = TRUE, conf.int = FALSE,
                           risk.table = TRUE, risk.table.col = "strata",
                           risk.table.y.text.col = TRUE,
                           risk.table.height = 0.25,
                           risk.table.title = "Number at Risk",
                           xlim = c(0, 35), # Limit x-axis to 35 years
                           break.time.by = 5, # Break x-axis every 5 years
                           ggtheme = theme_classic() + # Use classic theme
                             theme(panel.grid.major = element_line(color = "gray", size = 1),
                                   panel.grid.minor = element_line(color = "lightgray", size = 1)),
                           palette = c("red", "blue", "green", "purple", "orange", "brown"),
                           title = "KM Survival Curves for Outcomes (Side Recurrence)",
                           xlab = "Time (Years)", ylab = "Survival Probability",
                           tables.theme = theme_classic()) # Use classic theme for the risk table

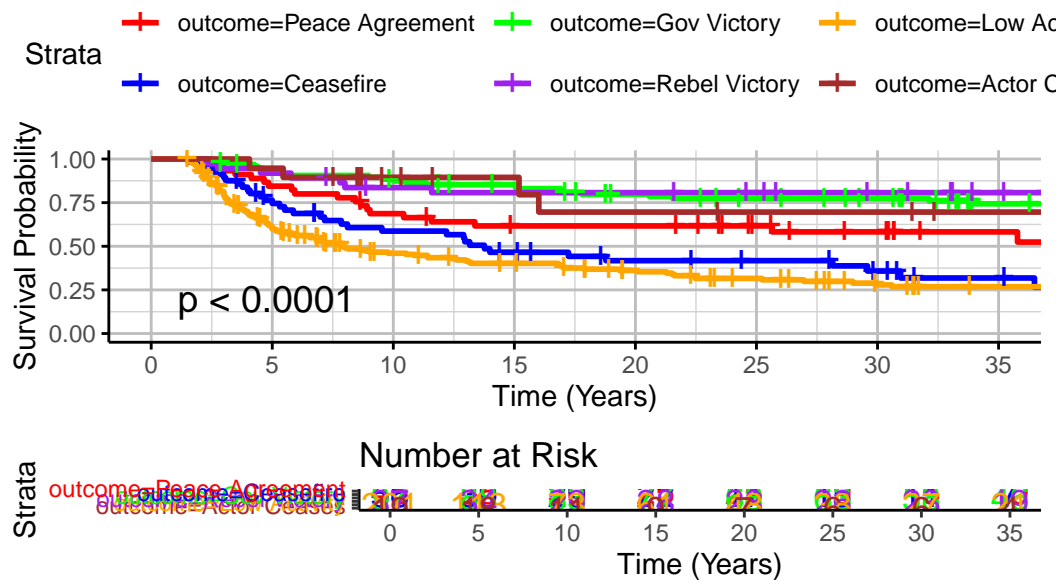
# Print plots
print(km_plot_any)
```

KM Survival Curves for Outcomes (Any Recurrence)



```
print(km_plot_side)
```

KM Survival Curves for Outcomes (Side Recurrence)



Cox Models

Cox P Models

```
m1<- coxph(Surv(time_to_recur_any, recur_any) ~dis+cease+govvic+rebvic+lowac, data = episodes,

m2<- coxph(Surv(time_to_recur_any, recur_any)~dis+cease+govvic+rebvic+lowac+pkou+log_dur+col

m3<- coxph(Surv(time_to_recur, recur_side) ~dis+cease+govvic+rebvic+lowac, data = episodes,

m4<- coxph(Surv(time_to_recur, recur_side)~dis+cease+govvic+rebvic+lowac+pkou+log_dur+cold_y

#stargazer(m1, m2, m3, m4, type = "text")

stargazer(m1, m2, m3, m4,
  ord.intercepts = TRUE,
  dep.var.labels = c("UCDP ID-Based War Recurrence", "UCDP ID-Based War Recurrence",
  covariate.labels = c("Actor Ceases", "Ceasefire", "Government Victory", "Rebel Victory")
```

Dependent variable:			
	UCDP ID-Based War Recurrence		UCDP ID-Based War Re
	(1)	(2)	(3)
Actor Ceases	-0.484 (0.459)	0.169 (0.606)	-0.447 (0.503)
Ceasefire	0.603** (0.263)	1.182*** (0.366)	0.658** (0.284)
Government Victory	-0.184 (0.252)	0.283 (0.377)	-0.757** (0.308)
Rebel Victory	0.105 (0.295)	0.822 (0.444)	-0.894** (0.422)

Low Activity	0.783*** (0.226)	1.186*** (0.335)	0.884*** (0.246)
Peacekeeping Missions		-0.585** (0.282)	
Log(Duration)		0.059* (0.029)	
Cold War		-0.492*** (0.175)	
Log(GDP per Capita)		-0.080 (0.057)	
Number of Veto Players		0.236*** (0.088)	
War over Government		-1.431*** (1.029)	
War over Territory		-1.321*** (1.031)	
Polity Score		-0.015 (0.022)	
Ethnic Fractionalization		1.508** (0.628)	
Coalition Size		0.028 (0.055)	
Log(Population)		-0.106 (0.365)	
Language Fractionalization		-1.130 (0.669)	
Power-Sharing		-0.192 (0.686)	

Religion Fractionalization

0.190
(0.247)

Observations	455	329	455
R2	0.107	0.209	0.197
Max. Possible R2	0.999	0.999	0.996
Log Likelihood	-1,550.938	-1,114.252	-1,219.034
Wald Test	48.250*** (df = 5)	264.530*** (df = 19)	94.070*** (df = 5)
LR Test	51.640*** (df = 5)	76.996*** (df = 19)	99.780*** (df = 5)
Score (Logrank) Test	51.438*** (df = 5)	78.548*** (df = 19)	95.665*** (df = 5)

Note:

*p<0.1; **p<0

```
#al_language2000
#al_religion2000
#fe_etfra
#fe_cultdiv
#al_ethnic2000
```

```
# Check proportional hazards assumption
m1_t<- cox.zph(m1)
m2_t<- cox.zph(m2)
m3_t<- cox.zph(m3)
m4_t<- cox.zph(m4)
print(m1_t)
```

	chisq	df	p
dis	0.0994	1	0.75260
cease	1.1805	1	0.27725
govvic	10.9580	1	0.00093
rebvic	1.1021	1	0.29381
lowac	6.0333	1	0.01404
GLOBAL	14.2584	5	0.01405

```
print(m2_t)
```

	chisq	df	p
--	-------	----	---

dis	0.0444332	1	0.833
cease	0.1483013	1	0.700
govvic	3.5989805	1	0.058
rebvic	0.0005874	1	0.981
lowac	0.8331863	1	0.361
pko_u	1.2850393	1	0.257
log_dur	5.4597689	1	0.019
cold_war	0.0324631	1	0.857
log_gdp	2.7998279	1	0.094
veto_u	1.6101650	1	0.204
ter_war	0.0000616	1	0.994
gov_war	0.0010633	1	0.974
p_polity2	4.4359281	1	0.035
fe_etfra	0.0239740	1	0.877
log_pop	0.0839223	1	0.772
al_religion2000	0.0111114	1	0.916
fe_cultdiv	1.2799303	1	0.258
W4	2.7115601	1	0.100
ps_original	3.4820684	1	0.062
GLOBAL	23.3819547	19	0.221

```
print(m3_t)
```

	chisq	df	p
dis	2.125	1	0.145
cease	3.704	1	0.054
govvic	0.645	1	0.422
rebvic	0.602	1	0.438
lowac	4.589	1	0.032
GLOBAL	8.251	5	0.143

```
print(m4_t)
```

	chisq	df	p
dis	2.69519262	1	0.1007
cease	4.56012230	1	0.0327
govvic	0.22149714	1	0.6379
rebvic	3.51213317	1	0.0609
lowac	2.44894340	1	0.1176
pko_u	0.01881588	1	0.8909
log_dur	25.99137111	1	0.00000034

cold_war	6.78720120	1	0.0092
log_gdp	0.04147132	1	0.8386
veto_u	5.16563745	1	0.0230
ter_war	0.79338533	1	0.3731
gov_war	0.84177544	1	0.3589
p_polity2	0.00027204	1	0.9868
fe_etfra	0.00000377	1	0.9985
log_pop	0.84916823	1	0.3568
al_religion2000	0.05014842	1	0.8228
fe_cultdiv	0.05344662	1	0.8172
W4	0.16978739	1	0.6803
ps_original	0.05276456	1	0.8183
GLOBAL	42.42831304	19	0.0016

```
#vif(m1)
#vif(m2)
#vif(m3)
#vif(m4)
```

```
library(corrplot)
```

corrplot 0.92 loaded

```
columns_of_interest <- episodes[, c("al_language2000", "al_religion2000", "fe_etfra", "fe_cultdiv")]
cor(columns_of_interest, use = "complete.obs")
```

	al_language2000	al_religion2000	fe_etfra	fe_cultdiv
al_language2000	1.0000000	0.2920441	0.7176872	0.7396354
al_religion2000	0.2920441	1.0000000	0.3563721	0.1479685
fe_etfra	0.7176872	0.3563721	1.0000000	0.7723940
fe_cultdiv	0.7396354	0.1479685	0.7723940	1.0000000

Coefficient Plots from Cox

```

# Extracting model coefficients and robust standard errors from the summary
coef_summary <- summary(m2)$coefficients

# Rename terms for better readability
terms_rename <- c(
  "dis" = "Actor Ceases",
  "cease" = "Ceasefire",
  "govvic" = "Government Victory",
  "rebvic" = "Rebel Victory",
  "lowac" = "Low Activity",
  "pko_u" = "Peacekeeping Missions",
  "log_dur" = "Log(Duration)",
  "cold_war" = "Cold War",
  "log_gdp" = "Log(GDP per Capita)",
  "veto_u" = "Number of Veto Players",
  "gov_war" = "War over Government",
  "ter_war" = "War over Territory",
  "p_polity2" = "Polity Score",
  "fe_etfra" = "Ethnic Fractionalization",
  "W4" = "Coalition Size",
  "log_pop" = "Log(Population)",
  "ps_original" = "Power-Sharing",
  "al_religion2000" = "Religion Fractionalization",
  "fe_cultdiv" = "Cultural Diversity"
)

# Create a dataframe for plotting
df_coef <- data.frame(
  Term = rownames(coef_summary),
  Estimate = coef_summary[, "coef"],
  StdErr = coef_summary[, "robust se"],
  Lower = coef_summary[, "coef"] - 1.96 * coef_summary[, "robust se"],
  Upper = coef_summary[, "coef"] + 1.96 * coef_summary[, "robust se"],
  p_value = coef_summary[, "Pr(>|z|)"]
)

# Apply the renaming to the Term column
df_coef$Term <- terms_rename[df_coef$Term]

# Order by Estimate from most negative to most positive
df_coef <- df_coef[order(df_coef$Estimate), ]

```

```

# Plotting using ggplot2
library(ggplot2)
coef_ucdp<-ggplot(df_coef, aes(x = Estimate, y = reorder(Term, Estimate))) +
  geom_point() +
  geom_errorbarh(aes(xmin = Lower, xmax = Upper), height = 0.2, color = ifelse(df_coef$p_val
  theme_minimal() +
  labs(title = "Coefficient Plot of Cox Model:UCDP",
        x = "Coefficient Value",
        y = "Variables")

# Extracting model coefficients and robust standard errors from the summary
coef_summary <- summary(m4)$coefficients

# Rename terms for better readability
terms_rename <- c(
  "dis" = "Actor Ceases",
  "cease" = "Ceasefire",
  "govvic" = "Government Victory",
  "rebvic" = "Rebel Victory",
  "lowac" = "Low Activity",
  "pko_u" = "Peacekeeping Missions",
  "log_dur" = "Log(Duration)",
  "cold_war" = "Cold War",
  "log_gdp" = "Log(GDP per Capita)",
  "veto_u" = "Number of Veto Players",
  "gov_war" = "War over Government",
  "ter_war" = "War over Territory",
  "p_polity2" = "Polity Score",
  "fe_etfra" = "Ethnic Fractionalization",
  "W4" = "Coalition Size",
  "log_pop" = "Log(Population)",
  "ps_original" = "Power-Sharing",
  "al_religion2000" = "Religion Fractionalization",
  "fe_cultdiv" = "Cultural Diversity"
)

# Create a dataframe for plotting
df_coef <- data.frame(
  Term = rownames(coef_summary),
  Estimate = coef_summary[, "coef"],
  StdErr = coef_summary[, "robust se"],

```

```

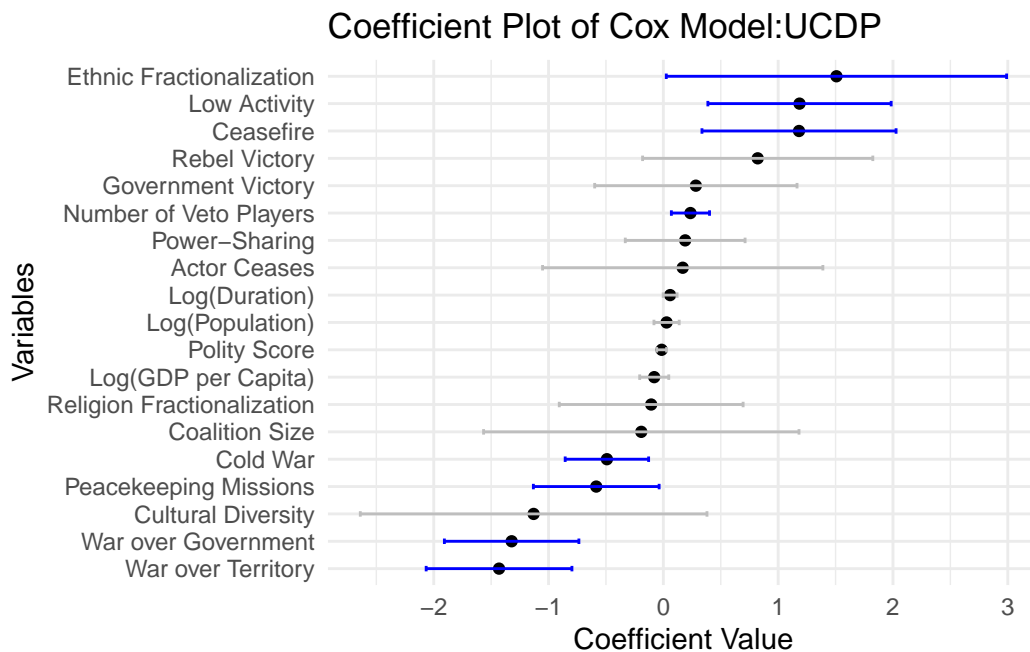
Lower = coef_summary[, "coef"] - 1.96 * coef_summary[, "robust se"],
Upper = coef_summary[, "coef"] + 1.96 * coef_summary[, "robust se"],
p_value = coef_summary[, "Pr(>|z|)"]
)

# Apply renaming to the Term column
df_coef$Term <- terms_rename[df_coef$Term]

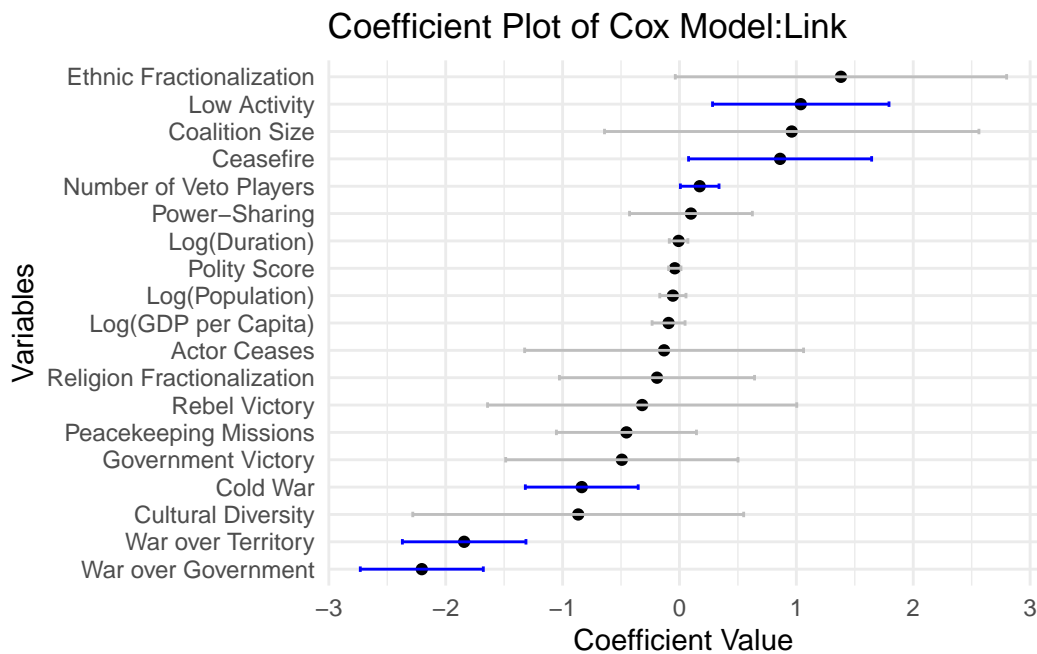
# Order by Estimate from most negative to most positive
df_coef <- df_coef[order(df_coef$Estimate), ]

# Plotting using ggplot2
library(ggplot2)
coef_link<-ggplot(df_coef, aes(x = Estimate, y = reorder(Term, Estimate))) +
  geom_point() +
  geom_errorbarh(aes(xmin = Lower, xmax = Upper), height = 0.2, color = ifelse(df_coef$p_value < 0.05, "blue", "grey")) +
  theme_minimal() +
  labs(title = "Coefficient Plot of Cox Model:Link",
       x = "Coefficient Value",
       y = "Variables")
coef_ucdp

```



coef_link



Survival Rates from Cox

```
library(survival)
library(dplyr)
library(survminer)
library(ggplot2)

# Define factor levels and labels
outcome_levels <- c("Peace Agreement", "Ceasefire", "Gov Victory", "Rebel Victory", "Low Act.")

# Convert outcome variable to factor with specified levels in the original dataset
episodes$outcomes <- factor(episodes$outcome, levels = 1:6, labels = outcome_levels)

# Convert the time variables from days to years in the original dataset
episodes$time_to_recur_any_years <- episodes$time_to_recur_any / 365.25
episodes$time_to_recur_years <- episodes$time_to_recur / 365.25

# Fit Cox proportional hazards models with time in years
```

```

m2 <- coxph(Surv(time_to_recur_any_years, recur_any) ~ as.factor(outcomes) + pko_u + log_dur

m4 <- coxph(Surv(time_to_recur_years, recur_side) ~ as.factor(outcomes) + pko_u + log_dur +

# Create representative data without al_language2000
representative_data <- episodes %>%
  summarise(
    pko_u = median(as.numeric(pko_u), na.rm = TRUE),
    log_dur = mean(log_dur, na.rm = TRUE),
    cold_war = median(cold_war, na.rm = TRUE),
    log_gdp = mean(log_gdp, na.rm = TRUE),
    veto_u = median(veto_u, na.rm = TRUE),
    gov_war = median(gov_war, na.rm = TRUE),
    ter_war = median(ter_war, na.rm = TRUE),
    p_polity2 = mean(p_polity2, na.rm = TRUE),
    fe_etfra = mean(fe_etfra, na.rm = TRUE),
    W4 = mean(W4, na.rm = TRUE),
    log_pop = mean(log_pop, na.rm = TRUE),
    ps_original = median(ps_original, na.rm = TRUE),
    al_religion2000 = mean(al_religion2000, na.rm = TRUE),
    fe_cultdiv = mean(fe_cultdiv, na.rm = TRUE)
  )

# Expand the data to include different outcomes
expanded_data <- do.call(rbind, replicate(6, representative_data, simplify = FALSE))
expanded_data$outcomes <- factor(rep(outcome_levels, each = 1), levels = outcome_levels)

# Generate survival curves based on the Cox model for m2
surv_fits_any <- survfit(m2, newdata = expanded_data)

# Define colors for the plot
colors <- c("Peace Agreement" = "red", "Ceasefire" = "blue", "Gov Victory" = "green", "Rebel

# Plotting survival curves for Any Recurrence with x-axis in years
plot_any <- ggsurvplot(
  surv_fits_any,
  data = expanded_data,
  pval = TRUE, # Show p-value for log-rank test
  conf.int = FALSE, # Show confidence intervals
  xlab = "Time in Years",
  ylab = "Survival Probability",
  ggtheme = theme_bw(),

```

```

xlim = c(0, 30), # Set the x-axis limit in years (adjust as needed)
break.time.by = 5, # Break x-axis every 5 years
risk.table = TRUE, # Show number at risk table
surv.scale = "percent",
legend.title = "Outcome",
legend.labs = levels(expanded_data$outcomes), # Automatically use factor labels
title = "Survival Probability Over Time for War Recurrence Based on ACD Conflict ID",
subtitle = "",
caption = "",
palette = colors,
size = 1.2, # Increase line width for better visibility
linetype = "solid" # Set line type to solid for all curves
)

```

Warning: `gather_()` was deprecated in tidyr 1.2.0.

i Please use `gather()` instead.

i The deprecated feature was likely used in the survminer package.

Please report the issue at <<https://github.com/kassambara/survminer/issues>>.

Warning in .pvalue(fit, data = data, method = method, pval = pval, pval.coord = pval.coord,
This is a null model.

Warning: No shared levels found between `names(values)` of the manual scale and the
data's fill values.

Warning in (function (survsummary, times, survtable = c("cumevents",
"risk.table", : The length of legend.labs should be 1

```

# Customize the legend font size and axis titles for Any Recurrence plot
plot_any$plot <- plot_any$plot +
  theme(legend.text = element_text(size = 12), # Increase legend text size
        legend.title = element_text(size = 14), # Increase legend title size
        axis.title.x = element_text(size = 10), # X-axis title size
        axis.title.y = element_text(size = 10), # Y-axis title size
        axis.text.x = element_text(size = 10, angle = 45, hjust = 1), # X-axis text size and
        axis.text.y = element_text(size = 10), # Y-axis text size
        plot.title = element_text(size = 12)) # Plot title size

# Customize the risk table text for Any Recurrence plot
plot_any$table <- plot_any$table +
  scale_x_continuous(breaks = seq(0, 30, by = 5), labels = seq(0, 30, by = 5)) + # Breaks and
  theme(axis.text.x = element_text(size = 10, angle = 45, hjust = 1)) # Rotate risk table t

```

Scale for x is already present.

Adding another scale for x, which will replace the existing scale.

```
# Generate survival curves based on the Cox model for m4
surv_fits_side <- survfit(m4, newdata = expanded_data)

# Plotting survival curves for Side Recurrence with x-axis in years
plot_side <- ggsurvplot(
  surv_fits_side,
  data = expanded_data,
  pval = TRUE, # Show p-value for log-rank test
  conf.int = FALSE, # Show confidence intervals
  xlab = "Time in Years",
  ylab = "Survival Probability",
  ggtheme = theme_bw(),
  xlim = c(0, 30), # Set the x-axis limit in years (adjust as needed)
  break.time.by = 5, # Break x-axis every 5 years
  risk.table = TRUE, # Show number at risk table
  surv.scale = "percent",
  legend.title = "Outcome",
  legend.labs = levels(expanded_data$outcomes), # Automatically use factor labels
  title = "Survival Probability Over Time for War Recurrence Based on Sufficient Linkage",
  subtitle = "",
  caption = "",
  palette = colors,
  size = 1.2, # Increase line width for better visibility
  linetype = "solid" # Set line type to solid for all curves
)
```

```
Warning in .pvalue(fit, data = data, method = method, pval = pval, pval.coord = pval.coord,
  This is a null model.
```

```
Warning: No shared levels found between `names(values)` of the manual scale and the
data's fill values.
```

```
Warning in (function (survsummary, times, survtable = c("cumevents",
"risk.table", : The length of legend.labs should be 1
```

```
# Customize the legend font size and axis titles for Side Recurrence plot
plot_side$plot <- plot_side$plot +
  theme(legend.text = element_text(size = 12), # Increase legend text size
```



```

    legend.title = element_text(size = 14), # Increase legend title size
    axis.title.x = element_text(size = 10), # X-axis title size
    axis.title.y = element_text(size = 10), # Y-axis title size
    axis.text.x = element_text(size = 10, angle = 45, hjust = 1), # X-axis text size and
    axis.text.y = element_text(size = 10), # Y-axis text size
    plot.title = element_text(size = 12)) # Plot title size

# Customize the risk table text for Side Recurrence plot
plot_side$table <- plot_side$table +
  scale_x_continuous(breaks = seq(0, 30, by = 5), labels = seq(0, 30, by = 5)) + # Breaks and
  theme(axis.text.x = element_text(size = 10, angle = 45, hjust = 1)) # Rotate risk table text

```

Scale for x is already present.

Adding another scale for x, which will replace the existing scale.

```

# Print the plots
print(plot_any)

```

Warning: No shared levels found between `names(values)` of the manual scale and the data's fill values.

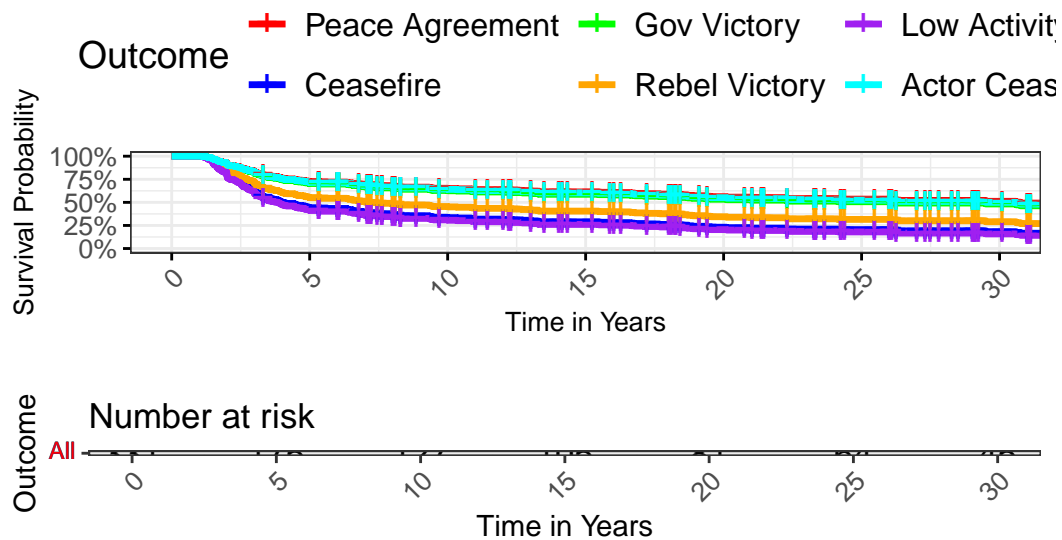
Warning: No shared levels found between `names(values)` of the manual scale and the data's fill values.

No shared levels found between `names(values)` of the manual scale and the data's fill values.

Warning: No shared levels found between `names(values)` of the manual scale and the data's colour values.

Warning: No shared levels found between `names(values)` of the manual scale and the data's fill values.

Survival Probability Over Time for War Recurrence Based on ACF



```
print(plot_side)
```

Warning: No shared levels found between `names(values)` of the manual scale and the data's fill values.

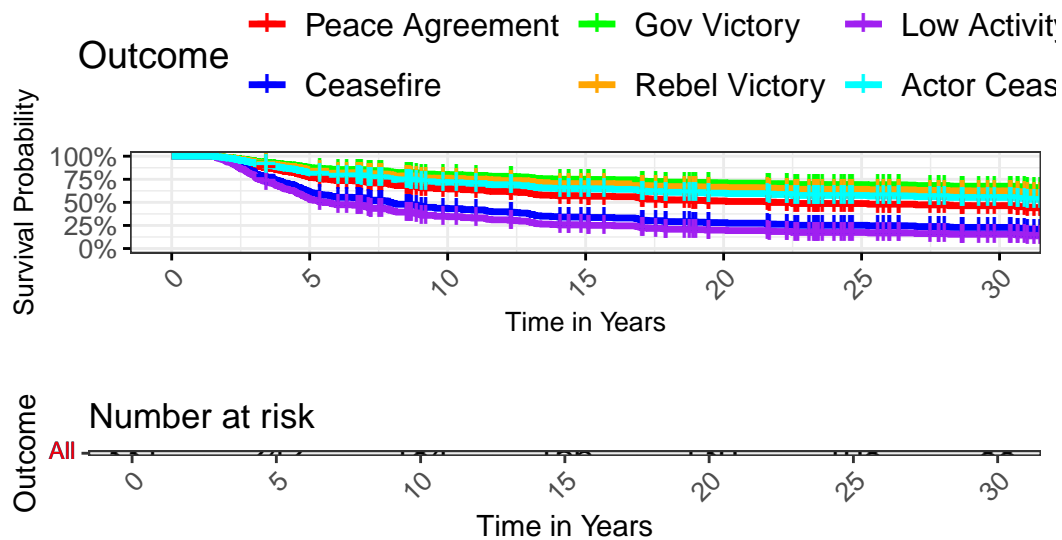
No shared levels found between `names(values)` of the manual scale and the data's fill values.

No shared levels found between `names(values)` of the manual scale and the data's fill values.

Warning: No shared levels found between `names(values)` of the manual scale and the data's colour values.

Warning: No shared levels found between `names(values)` of the manual scale and the data's fill values.

Survival Probability Over Time for War Recurrence Based on Suff



RSF Models

Var Imp with RSF: Any

```
# Applying labels
episodes_a <- apply_labels(episodes,
                           time_to_recur_any = "Time to Recur Any",
                           recur_any = "War Recurrence Any",
                           dis = "Actor Ceases",
                           cease = "Ceasefire",
                           govvic = "Government Victory",
                           rebvic = "Rebel Victory",
                           lowac = "Low Activity",
                           W4 = "Coalition Size",
                           pko_u = "Peacekeeping Operations",
                           log_dur = "Log Duration",
                           cold_war = "Cold War",
                           p_polity2 = "Polity Score",
                           fe_etfra = "Ethnic Fractionalization",
                           log_gdp = "Log GDP per Capita",
                           log_pop = "Log Population",
```

```

        ps_original = "Power-Sharing",
        al_religion2000 = "Religion Fractionalization",
        fe_cultdiv = "Cultural Diversity",
        veto_u = "Veto Players",
        gov_war = "War over Government",
        ter_war = "War over Territory")

# Subsetting data
X <- subset(episodes_a, select = c(time_to_recur_any, recur_any, dis, cease, govvic, rebvic,
X <- na.omit(X)

# Ensure numeric variables are correctly formatted
X <- X %>%
  mutate(across(everything(), as.numeric))

# Define cross-validation folds
set.seed(123) # For reproducibility
folds <- createFolds(X$recur_any, k = 5, list = TRUE)

# Initialize a list to store variable importance scores
importance_list <- vector("list", length(folds))

# Perform cross-validation
for (i in seq_along(folds)) {
  # Define training and validation sets
  train_indices <- unlist(folds[-i])
  test_indices <- unlist(folds[i])

  train_data <- X[train_indices, ]
  test_data <- X[test_indices, ]

  # Train the Random Survival Forest model
  rsf <- rfsrc(Surv(time_to_recur_any, recur_any) ~ ., data = train_data, ntree = 1000, nodes

  # Store the variable importance scores
  importance_list[[i]] <- rsf$importance
}

# Aggregate variable importance scores
importance_df <- as.data.frame(do.call(cbind, importance_list))
importance_df$Variable <- rownames(importance_df)
rownames(importance_df) <- NULL

```

```

importance_df$Importance <- rowMeans(importance_df[, -ncol(importance_df)])
importance_df <- importance_df[, c("Variable", "Importance")]

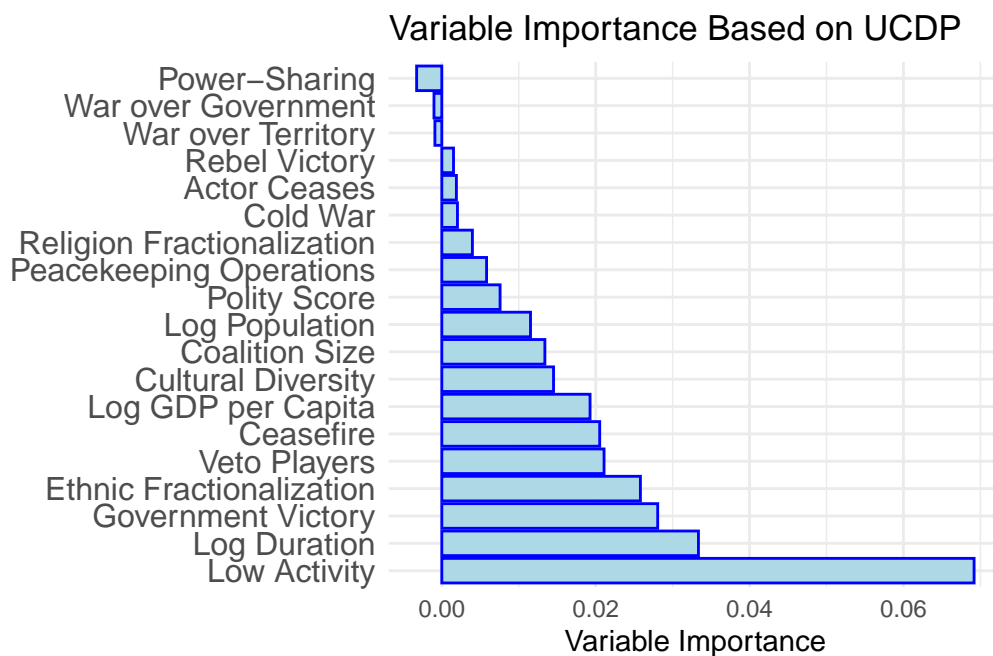
# Relabeling variables for plot
var_labels <- c("Actor Ceases", "Ceasefire", "Government Victory", "Rebel Victory", "Low Act.
names(var_labels) <- c("dis", "cease", "govvic", "rebtvic", "lowac", "pko_u", "log_dur", "col

# Adding variable labels
importance_df$Variable <- factor(importance_df$Variable, levels = names(var_labels), labels =

# Order by importance
importance_df <- importance_df %>% arrange(desc(Importance))
importance_df$Variable <- factor(importance_df$Variable, levels = importance_df$Variable)

# Plotting using ggplot2
ggplot(importance_df, aes(x = Importance, y = Variable)) +
  geom_bar(stat = "identity", fill = "lightblue", color = "blue") +
  theme_minimal() +
  labs(title = "Variable Importance Based on UCDP", x = "Variable Importance", y = "") +
  theme(axis.text.y = element_text(size = 12))

```



Var Imp with RSF: Link

```
# Applying labels
episodes_a <- apply_labels(episodes,
                           time_to_recur = "Time to Recur",
                           recur_side = "War Recurrence",
                           dis = "Actor Ceases",
                           cease = "Ceasefire",
                           govvic = "Government Victory",
                           rebvic = "Rebel Victory",
                           lowac = "Low Activity",
                           W4 = "Coalition Size",
                           pko_u = "Peacekeeping Operations",
                           log_dur = "Log Duration",
                           cold_war = "Cold War",
                           p_polity2 = "Polity Score",
                           fe_etfra = "Ethnic Fractionalization",
                           log_gdp = "Log GDP per Capita",
                           log_pop = "Log Population",
                           ps_original = "Power-Sharing",
                           al_religion2000 = "Religion Fractionalization",
                           fe_cultdiv = "Cultural Diversity",
                           veto_u = "Veto Players",
                           gov_war = "War over Government",
                           ter_war = "War over Territory")

# Subsetting data
X <- subset(episodes_a, select = c(time_to_recur, recur_side, dis, cease, govvic, rebvic, lowac, W4, pko_u, log_dur, cold_war, p_polity2, fe_etfra, log_gdp, log_pop, ps_original, al_religion2000, fe_cultdiv, veto_u, gov_war, ter_war))
X <- na.omit(X)

# Ensure numeric variables are correctly formatted
X <- X %>%
  mutate(across(everything(), as.numeric))

# Define cross-validation folds
set.seed(123) # For reproducibility
folds <- createFolds(X$recur_side, k = 5, list = TRUE)

# Initialize a list to store variable importance scores
importance_list <- vector("list", length(folds))

# Perform cross-validation
```

```

for (i in seq_along(folds)) {
  # Define training and validation sets
  train_indices <- unlist(folds[-i])
  test_indices <- unlist(folds[i])

  train_data <- X[train_indices, ]
  test_data <- X[test_indices, ]

  # Train the Random Survival Forest model
  rsf <- rfsrc(Surv(time_to_recur, recur_side) ~ ., data = train_data, ntree = 1000, nodesize = 1)

  # Store the variable importance scores
  importance_list[[i]] <- rsf$importance
}

# Aggregate variable importance scores
importance_df <- as.data.frame(do.call(cbind, importance_list))
importance_df$Variable <- rownames(importance_df)
rownames(importance_df) <- NULL
importance_df$Importance <- rowMeans(importance_df[, -ncol(importance_df)])
importance_df <- importance_df[, c("Variable", "Importance")]

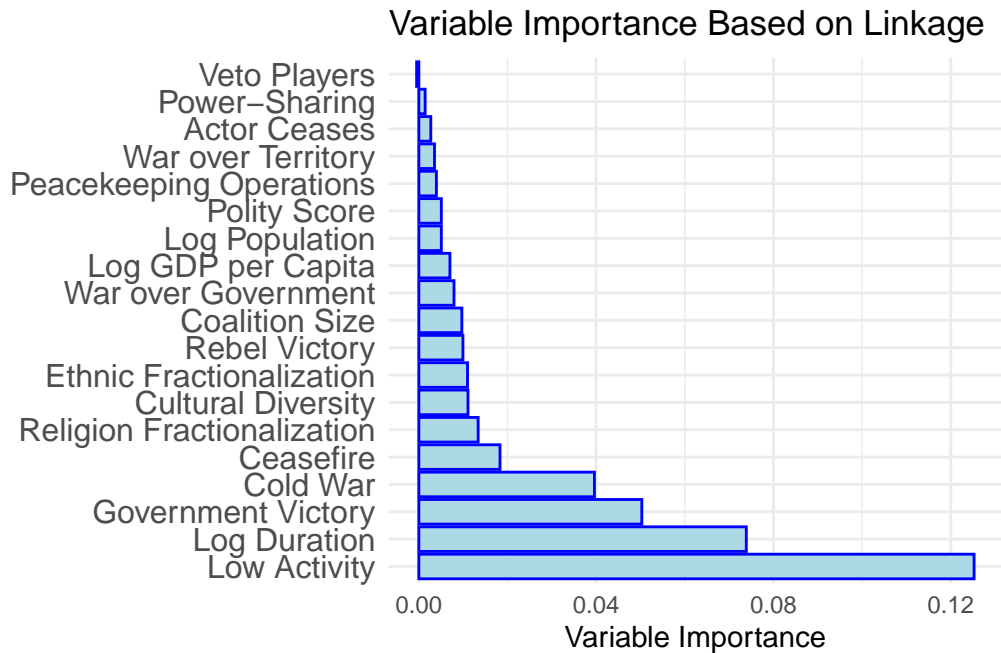
# Relabeling variables for plot
var_labels <- c("Actor Ceases", "Ceasefire", "Government Victory", "Rebel Victory", "Low Act.", "Log Duration", "Pko_u", "Reb_u", "Reb_v", "Gov_v", "Dis")
names(var_labels) <- c("dis", "cease", "govvic", "rebvic", "lowac", "pko_u", "log_dur", "col", "reb_u", "gov_u", "dis")

# Adding variable labels
importance_df$Variable <- factor(importance_df$Variable, levels = names(var_labels), labels = var_labels)

# Order by importance
importance_df <- importance_df %>% arrange(desc(Importance))
importance_df$Variable <- factor(importance_df$Variable, levels = importance_df$Variable)

# Plotting using ggplot2
ggplot(importance_df, aes(x = Importance, y = Variable)) +
  geom_bar(stat = "identity", fill = "lightblue", color = "blue") +
  theme_minimal() +
  labs(title = "Variable Importance Based on Linkage", x = "Variable Importance", y = "") +
  theme(axis.text.y = element_text(size = 12))

```



RSF- Any Survival with No CV

```
# Subsetting and cleaning data
X <- subset(episodes, select = c(time_to_recur_any, recur_any, dis, cease, govvic, rebvic, l
                                cold_war, p_polity2, fe_etfra, log_gdp, log_pop, ps_origi
                                al_religion2000, fe_cultdiv, veto_u, gov_war, ter_war))

X <- na.omit(X)
X <- X %>% mutate(across(everything(), as.numeric))

# Convert time to years
X$time_to_recur_any <- X$time_to_recur_any / 365.25

# Train the Random Survival Forest model on the entire dataset
fit <- rfsrc(Surv(time_to_recur_any, recur_any) ~ ., data = X, ntree = 1000, nodesize = 5, n

# Predict survival probabilities for the entire dataset
pred <- predict(fit, X, OOB = TRUE, type = "response")
survival_probs <- as.data.frame(pred$survival)

# Extract time points (already in years)
time_points <- fit$time.interest
```



```

colnames(survival_probs) <- paste("Time", round(time_points, 2), sep = "_")
survival_probs$id <- 1:nrow(survival_probs)
X$id <- 1:nrow(X)

# Convert survival probabilities to long format for plotting
survival_probs_long <- survival_probs %>%
  pivot_longer(cols = -id, names_to = "Time", values_to = "Probability") %>%
  mutate(Time = as.numeric(gsub("Time_", "", Time)))

# Combine with the original data to get the binary outcome variables
survival_probs_long <- left_join(survival_probs_long, X, by = "id")

# Function to calculate mean and confidence intervals
calculate_summary <- function(data) {
  data %>%
    group_by(Time) %>%
    summarise(
      Mean_Probability = mean(Probability, na.rm = TRUE),
      SD = sd(Probability, na.rm = TRUE),
      Lower_CI = Mean_Probability - qt(0.975, length(Probability)-1) * SD / sqrt(length(Probability)),
      Upper_CI = Mean_Probability + qt(0.975, length(Probability)-1) * SD / sqrt(length(Probability))
    )
}

# Calculate summary statistics for each binary outcome variable
dis_summary <- calculate_summary(filter(survival_probs_long, dis == 1))
cease_summary <- calculate_summary(filter(survival_probs_long, cease == 1))
govvic_summary <- calculate_summary(filter(survival_probs_long, govvic == 1))
rebvic_summary <- calculate_summary(filter(survival_probs_long, rebvic == 1))
lowac_summary <- calculate_summary(filter(survival_probs_long, lowac == 1))
peace_summary <- calculate_summary(filter(survival_probs_long, peace == 1))

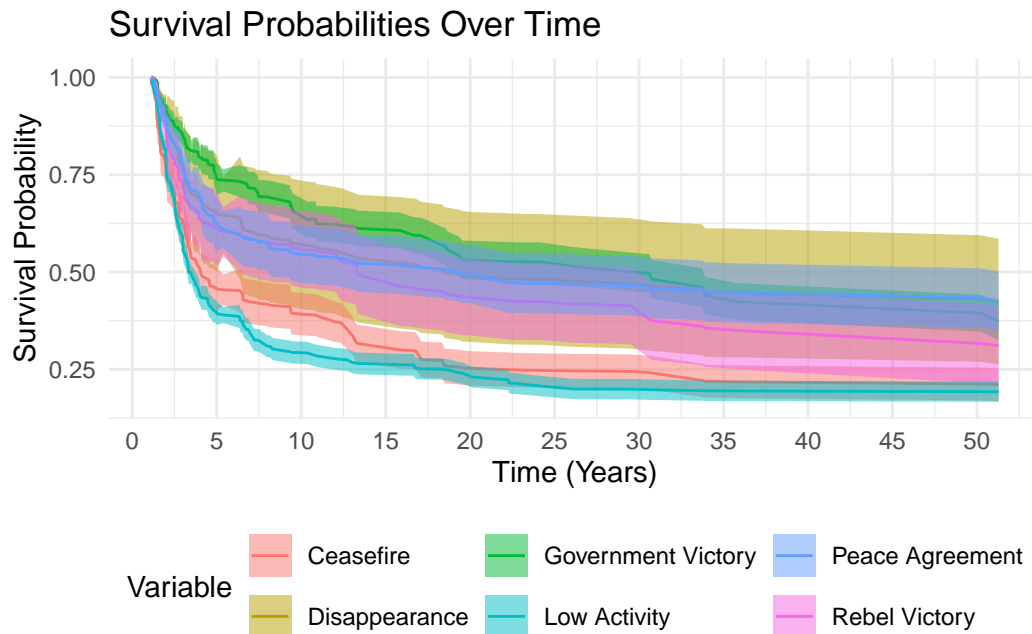
# Plot using ggplot2, one line per variable of interest with confidence intervals
ggplot() +
  geom_line(data = dis_summary, aes(x = Time, y = Mean_Probability, color = "Disappearance")) +
  geom_ribbon(data = dis_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Disappearance")) +
  geom_line(data = cease_summary, aes(x = Time, y = Mean_Probability, color = "Ceasefire")) +
  geom_ribbon(data = cease_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Ceasefire")) +
  geom_line(data = govvic_summary, aes(x = Time, y = Mean_Probability, color = "Government Victory")) +
  geom_ribbon(data = govvic_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Government Victory")) +
  geom_line(data = rebvic_summary, aes(x = Time, y = Mean_Probability, color = "Rebel Victory")) +
  geom_ribbon(data = rebvic_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Rebel Victory"))

```

```

geom_line(data = lowac_summary, aes(x = Time, y = Mean_Probability, color = "Low Activity"))
geom_ribbon(data = lowac_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Low Activity"))
geom_line(data = peace_summary, aes(x = Time, y = Mean_Probability, color = "Peace Agreement"))
geom_ribbon(data = peace_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Peace Agreement"))
scale_x_continuous(breaks = seq(0, max(survival_probs_long$Time), by = 5)) + # Set breaks
labs(title = "Survival Probabilities Over Time", x = "Time (Years)", y = "Survival Probability")
theme_minimal() +
theme(legend.position = "bottom")

```



RSF- Side Survival with No CV

```

# Subsetting and cleaning data
X <- subset(episodes, select = c(time_to_recur, recur_side, dis, cease, govvic, rebvic, lowac,
                                cold_war, p_polity2, fe_etfra, log_gdp, log_pop, ps_origin,
                                al_religion2000, fe_cultdiv, veto_u, gov_war, ter_war))

X <- na.omit(X)
X <- X %>% mutate(across(everything(), as.numeric))

# Convert time to years
X$time_to_recur <- X$time_to_recur / 365.25

```

```

# Train the Random Survival Forest model on the entire dataset
fit <- rfsrc(Surv(time_to_recur, recur_side) ~ ., data = X, ntree = 1000, nodesize = 5, nspl

# Predict survival probabilities for the entire dataset
pred <- predict(fit, X, OOB = TRUE, type = "response")
survival_probs <- as.data.frame(pred$survival)

# Extract time points (already in years)
time_points <- fit$time.interest
colnames(survival_probs) <- paste("Time", round(time_points, 2), sep = "_")
survival_probs$id <- 1:nrow(survival_probs)
X$id <- 1:nrow(X)

# Convert survival probabilities to long format for plotting
survival_probs_long <- survival_probs %>%
  pivot_longer(cols = -id, names_to = "Time", values_to = "Probability") %>%
  mutate(Time = as.numeric(gsub("Time_", "", Time)))

# Combine with the original data to get the binary outcome variables
survival_probs_long <- left_join(survival_probs_long, X, by = "id")

# Function to calculate mean and confidence intervals
calculate_summary <- function(data) {
  data %>%
    group_by(Time) %>%
    summarise(
      Mean_Probability = mean(Probability, na.rm = TRUE),
      SD = sd(Probability, na.rm = TRUE),
      Lower_CI = Mean_Probability - qt(0.975, length(Probability)-1) * SD / sqrt(length(Probab
      Upper_CI = Mean_Probability + qt(0.975, length(Probability)-1) * SD / sqrt(length(Probab
    )
}

# Calculate summary statistics for each binary outcome variable
dis_summary <- calculate_summary(filter(survival_probs_long, dis == 1))
cease_summary <- calculate_summary(filter(survival_probs_long, cease == 1))
govvic_summary <- calculate_summary(filter(survival_probs_long, govvic == 1))
rebvic_summary <- calculate_summary(filter(survival_probs_long, rebvic == 1))
lowac_summary <- calculate_summary(filter(survival_probs_long, lowac == 1))
peace_summary <- calculate_summary(filter(survival_probs_long, peace == 1))

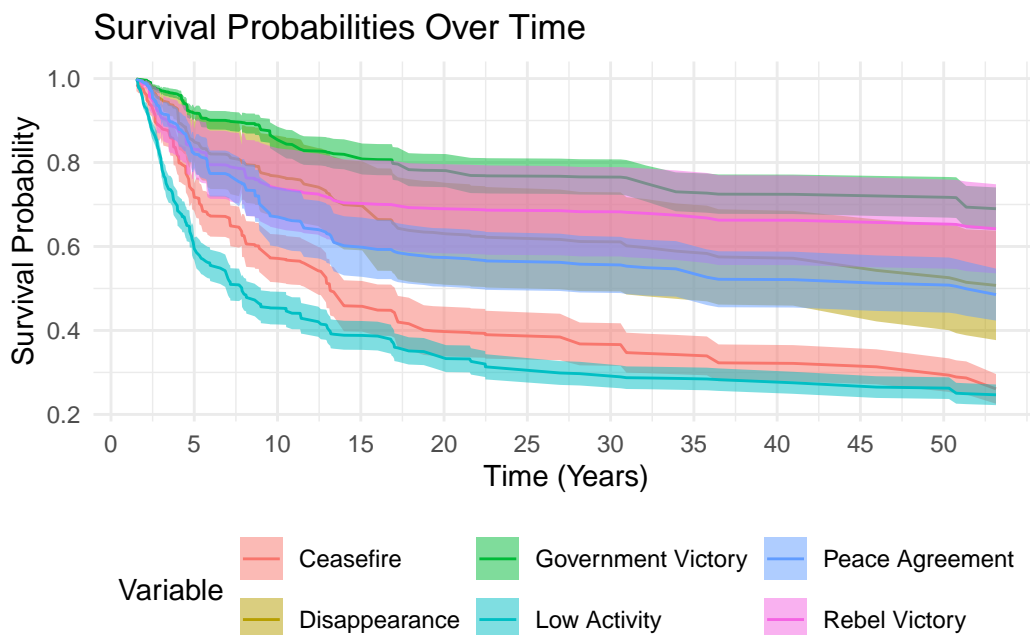
# Plot using ggplot2, one line per variable of interest with confidence intervals

```

```

ggplot() +
  geom_line(data = dis_summary, aes(x = Time, y = Mean_Probability, color = "Disappearance")) +
  geom_ribbon(data = dis_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Disappearance")) +
  geom_line(data = cease_summary, aes(x = Time, y = Mean_Probability, color = "Ceasefire")) +
  geom_ribbon(data = cease_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Ceasefire")) +
  geom_line(data = govvic_summary, aes(x = Time, y = Mean_Probability, color = "Government Victory")) +
  geom_ribbon(data = govvic_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Government Victory")) +
  geom_line(data = rebvic_summary, aes(x = Time, y = Mean_Probability, color = "Rebel Victory")) +
  geom_ribbon(data = rebvic_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Rebel Victory")) +
  geom_line(data = lowac_summary, aes(x = Time, y = Mean_Probability, color = "Low Activity")) +
  geom_ribbon(data = lowac_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Low Activity")) +
  geom_line(data = peace_summary, aes(x = Time, y = Mean_Probability, color = "Peace Agreement")) +
  geom_ribbon(data = peace_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Peace Agreement")) +
  scale_x_continuous(breaks = seq(0, max(survival_probs_long$Time), by = 5)) + # Set breaks
  labs(title = "Survival Probabilities Over Time", x = "Time (Years)", y = "Survival Probability") +
  theme_minimal() +
  theme(legend.position = "bottom")

```



No Low Ac

```
### remove low activity cases
episodes <- episodes %>% filter(lowac != 1)
```

Kaplan-Mier Curves

```
# Create a new dataframe from episodes
new_episodes <- episodes

# Convert the numerical outcome variable to a factor with appropriate labels, excluding "Low
new_episodes$outcome <- factor(new_episodes$outcome, levels = c(1, 2, 3, 4, 6),
                              labels = c("Peace Agreement", "Ceasefire", "Gov Victory", "Rel

# Check the levels of the outcome variable
print(levels(new_episodes$outcome))
```

```
[1] "Peace Agreement" "Ceasefire"          "Gov Victory"      "Rebel Victory"
[5] "Actor Ceases"
```

```
# Inspect the distribution of survival times
summary(new_episodes$time_to_recur_any)
```

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
369	1365	4705	6941	10768	27774

```
summary(new_episodes$time_to_recur)
```

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
560	3189	9357	10573	16567	28034

```
# Trim follow-up time to 20,000 days
new_episodes$time_to_recur_any <- pmin(new_episodes$time_to_recur_any, 20000)
new_episodes$time_to_recur <- pmin(new_episodes$time_to_recur, 20000)

# Fit the Kaplan-Meier survival curves for the first plot
km_fit_outcome_any <- survfit(Surv(time_to_recur_any, recur_any) ~ outcome, data = new_episodes)
```

```

# Plot the Kaplan-Meier survival curves with percentages in the risk table
km_plot_any <- ggsurvplot(km_fit_outcome_any, data = new_episodes,
  pval = TRUE, conf.int = FALSE,
  risk.table = TRUE, risk.table.col = "strata",
  risk.table.y.text.col = TRUE,
  risk.table.height = 0.25,
  risk.table.title = "Number at Risk",
  ggtheme = theme_classic(),
  palette = c("red", "blue", "green", "purple", "brown"),
  title = "KM Survival Curves for Outcomes (Any Recurrence)",
  xlab = "Time", ylab = "Survival Probability",
  tables.y.text = FALSE)

# Customize the survival plot to add grid lines
km_plot_any$plot <- km_plot_any$plot +
  theme(legend.text = element_text(size = 10), # Increase legend text size
    legend.title = element_text(size = 12), # Increase legend title size
    axis.title.x = element_text(size = 10), # X-axis title size
    axis.title.y = element_text(size = 10), # Y-axis title size
    axis.text.x = element_text(size = 10), # X-axis text size
    axis.text.y = element_text(size = 10), # Y-axis text size
    plot.title = element_text(size = 12), # Plot title size
    panel.grid.major = element_line(color = "gray", size = 0.5), # Add major grid lines
    panel.grid.minor = element_line(color = "lightgray", size = 0.25)) # Add minor grid lines

# Remove grid lines from the risk table
km_plot_any$risk.table <- km_plot_any$risk.table +
  theme(panel.grid.major = element_blank(),
    panel.grid.minor = element_blank())

# Fit the Kaplan-Meier survival curves for the second plot
km_fit_outcome_side <- survfit(Surv(time_to_recur, recur_side) ~ outcome, data = new_episodes)

# Plot the Kaplan-Meier survival curves with percentages in the risk table for the second plot
km_plot_side <- ggsurvplot(km_fit_outcome_side, data = new_episodes,
  pval = TRUE, conf.int = FALSE,
  risk.table = TRUE, risk.table.col = "strata",
  risk.table.y.text.col = TRUE,
  risk.table.height = 0.25,
  risk.table.title = "Number at Risk",
  ggtheme = theme_classic(),
  palette = c("red", "blue", "green", "purple", "brown"),

```

```

        title = "KM Survival Curves for Outcomes (Side Recurrence)",
        xlab = "Time", ylab = "Survival Probability",
        tables.y.text = FALSE)

# Customize the survival plot to add grid lines
km_plot_side$plot <- km_plot_side$plot +
  theme(legend.text = element_text(size = 10), # Increase legend text size
        legend.title = element_text(size = 12), # Increase legend title size
        axis.title.x = element_text(size = 10), # X-axis title size
        axis.title.y = element_text(size = 10), # Y-axis title size
        axis.text.x = element_text(size = 10), # X-axis text size
        axis.text.y = element_text(size = 10), # Y-axis text size
        plot.title = element_text(size = 12), # Plot title size
        panel.grid.major = element_line(color = "gray", size = 0.5), # Add major grid line
        panel.grid.minor = element_line(color = "lightgray", size = 0.25)) # Add minor grid line

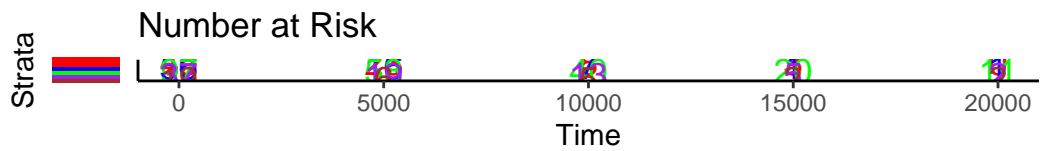
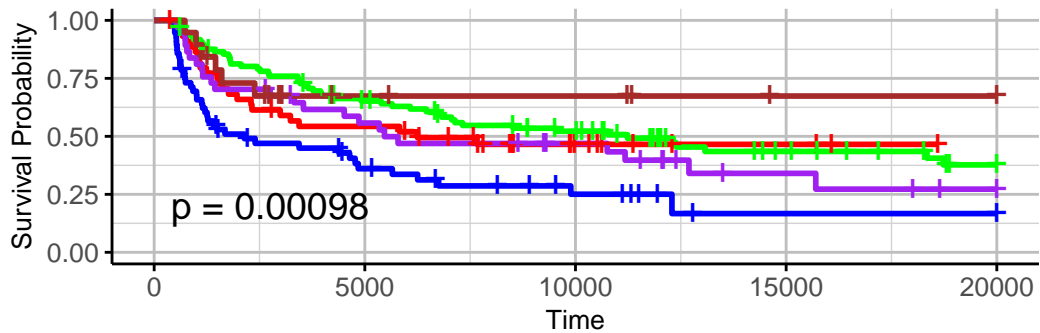
# Remove grid lines from the risk table
km_plot_side$risk.table <- km_plot_side$risk.table +
  theme(panel.grid.major = element_blank(),
        panel.grid.minor = element_blank())

# Print the plots
print(km_plot_any)

```

KM Survival Curves for Outcomes (Any Recurrence)

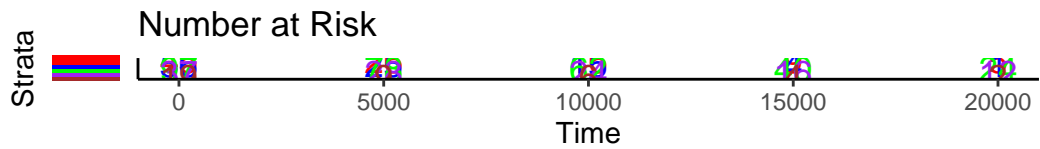
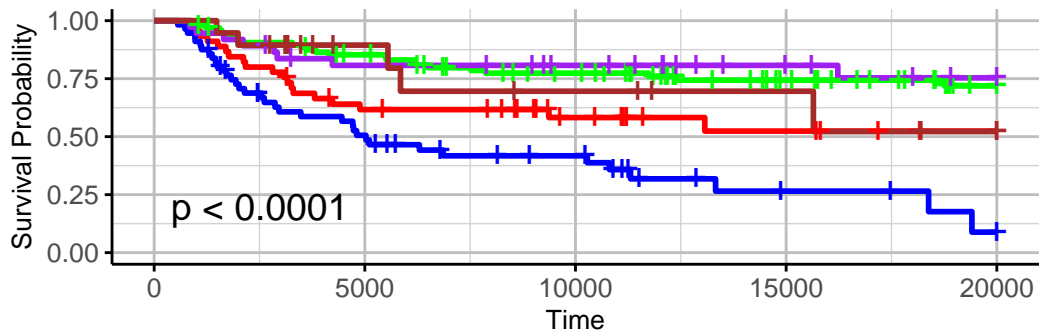
Agreement + outcome=Ceasefire + outcome=Gov Victory + outcome=Rebel



```
print(km_plot_side)
```

KM Survival Curves for Outcomes (Side Recurrence)

Agreement + outcome=Ceasefire + outcome=Gov Victory + outcome=Rebel



Cox models

```
## Models

m1 <- coxph(Surv(time_to_recur_any, recur_any) ~ dis + cease + govvic + rebvic, data = episodes)
m2 <- coxph(Surv(time_to_recur_any, recur_any) ~ dis + cease + govvic + rebvic + pko_u + log_dur)
m3 <- coxph(Surv(time_to_recur, recur_side) ~ dis + cease + govvic + rebvic, data = episodes)
m4 <- coxph(Surv(time_to_recur, recur_side) ~ dis + cease + govvic + rebvic + pko_u + log_dur)

stargazer(m1, m2, m3, m4, type = "text")
```

=====				
	Dependent variable:			
	time_to_recur_any		time_to_recur	
	(1)	(2)	(3)	(4)

dis	-0.472 (0.459)	0.349 (0.667)	-0.440 (0.503)	-0.160 (0.747)
cease	0.620** (0.263)	1.351** (0.445)	0.678** (0.284)	0.778 (0.471)
govvic	-0.222 (0.253)	0.370 (0.455)	-0.790** (0.309)	-0.252 (0.528)
rebvic	0.086 (0.296)	0.831 (0.521)	-0.924** (0.423)	-0.227 (0.696)
pko_u		-0.552 (0.462)		-0.691 (0.527)
log_dur		0.082* (0.045)		0.065 (0.061)
cold_war		-0.420 (0.275)		-1.074** (0.358)

log_gdp		0.003 (0.090)		-0.110 (0.113)
veto_u		0.390** (0.146)		0.344* (0.166)
gov_war		0.401 (0.266)		-0.264 (0.302)
ter_war			(0.000)	(0.000)
p_polity2		0.010 (0.037)		-0.045 (0.046)
fe_etfra		0.496 (1.001)		-0.451 (1.168)
W4		-0.861 (1.122)		1.523 (1.487)
log_pop		0.117* (0.071)		-0.032 (0.088)
al_language2000		1.737** (0.700)		2.493*** (0.891)
ps_original		0.055 (0.349)		-0.137 (0.418)
al_religion2000		-0.690 (0.544)		-0.916 (0.663)
fe_cultdiv		-1.729 (0.995)		-1.586 (1.263)

Observations	254	173	254	173
R2	0.062	0.251	0.134	0.291
Max. Possible R2	0.996	0.995	0.977	0.977
Log Likelihood	-705.245	-440.117	-458.157	-297.331
Wald Test	15.440*** (df = 4)	64.930*** (df = 18)	42.750*** (df = 4)	64.860*** (df = 4)

LR Test 16.391*** (df = 4) 49.916*** (df = 18) 36.449*** (df = 4) 59.464*** (df = 18)
 Score (Logrank) Test 18.506*** (df = 4) 54.963*** (df = 18) 43.412*** (df = 4) 61.667*** (df = 18)

Note:

*p<0.1; **p<0.05; ***p<0.01

```
stargazer(m1, m2, m3, m4,
  ord.intercepts = TRUE,
  dep.var.labels = c("UCDP ID-Based War Recurrence", "UCDP ID-Based War Recurrence",
  covariate.labels = c("Actor Ceases", "Ceasefire", "Government Victory", "Rebel Victory",
  type = "text")
```

Dependent variable:				
	UCDP ID-Based War Recurrence (1)	UCDP ID-Based War Recurrence (2)	UCDP ID-Based War Recurrence (3)	UCDP ID-Based War Recurrence (4)
Actor Ceases	-0.472 (0.459)	0.349 (0.667)	-0.440 (0.503)	-0.440 (0.503)
Ceasefire	0.620** (0.263)	1.351** (0.445)	0.678** (0.284)	0.678** (0.284)
Government Victory	-0.222 (0.253)	0.370 (0.455)	-0.790** (0.309)	-0.790** (0.309)
Rebel Victory	0.086 (0.296)	0.831 (0.521)	-0.924** (0.423)	-0.924** (0.423)
Peacekeeping Missions		-0.552 (0.462)		-0.552 (0.462)
Log(Duration)		0.082* (0.045)		0.082* (0.045)
Cold War		-0.420 (0.275)		-0.420 (0.275)
Log(GDP per Capita)		0.003 (0.090)		0.003 (0.090)

Number of Veto Players	0.390** (0.146)	0 (0)
War over Government	0.401 (0.266)	-0 (0)
War over Territory	(0.000)	(0)
Polity Score	0.010 (0.037)	-0 (0)
Ethnic Fractionalization	0.496 (1.001)	-0 (0)
Coalition Size	-0.861 (1.122)	0 (0)
Log(Population)	0.117* (0.071)	-0 (0)
Language Fractionalization	1.737** (0.700)	2.4 (0)
Power-Sharing	0.055 (0.349)	-0 (0)
Religion Fractionalization	-0.690 (0.544)	-0 (0)
Cultural Diversity	-1.729 (0.995)	-0 (0)

Observations	254	173	254	
R2	0.062	0.251	0.134	0
Max. Possible R2	0.996	0.995	0.977	0
Log Likelihood	-705.245	-440.117	-458.157	-25
Wald Test	15.440*** (df = 4)	64.930*** (df = 18)	42.750*** (df = 4)	64.860***
LR Test	16.391*** (df = 4)	49.916*** (df = 18)	36.449*** (df = 4)	59.464***
Score (Logrank) Test	18.506*** (df = 4)	54.963*** (df = 18)	43.412*** (df = 4)	61.667***

Note:

*p<0.1; **p<0.05

Coefficient Plots fom Cox

```
# Extracting model coefficients and robust standard errors from the summary
coef_summary <- summary(m2)$coefficients

# Rename terms for better readability
terms_rename <- c(
  "dis" = "Actor Ceases",
  "cease" = "Ceasefire",
  "govvic" = "Government Victory",
  "rebvic" = "Rebel Victory",
  "pko_u" = "Peacekeeping Missions",
  "log_dur" = "Log(Duration)",
  "cold_war" = "Cold War",
  "log_gdp" = "Log(GDP per Capita)",
  "veto_u" = "Number of Veto Players",
  "gov_war" = "War over Government",
  "ter_war" = "War over Territory",
  "p_polity2" = "Polity Score",
  "fe_etfra" = "Ethnic Fractionalization",
  "W4" = "Coalition Size",
  "log_pop" = "Log(Population)",
  "al_language2000" = "Language Fractionalization",
  "ps_original" = "Power-Sharing",
  "al_religion2000" = "Religion Fractionalization",
  "fe_cultdiv" = "Cultural Diversity"
)

# Create a dataframe for plotting
df_coef <- data.frame(
  Term = rownames(coef_summary),
  Estimate = coef_summary[, "coef"],
  StdErr = coef_summary[, "robust se"],
  Lower = coef_summary[, "coef"] - 1.96 * coef_summary[, "robust se"],
  Upper = coef_summary[, "coef"] + 1.96 * coef_summary[, "robust se"],
  p_value = coef_summary[, "Pr(>|z|)"]
)

# Apply the renaming to the Term column
df_coef$Term <- terms_rename[df_coef$Term]

# Plotting using ggplot2
```

```

library(ggplot2)
cofp_any<-ggplot(df_coef, aes(x = Estimate, y = Term)) +
  geom_point() +
  geom_errorbarh(aes(xmin = Lower, xmax = Upper), height = 0.2, color = ifelse(df_coef$p_val
  theme_minimal() +
  labs(title = "Coefficient Plot of Cox Model: UCDP",
        x = "Coefficient Value",
        y = "Variables")

# Extracting model coefficients and robust standard errors from the summary
coef_summary <- summary(m4)$coefficients

# Create a dataframe for plotting
df_coef <- data.frame(
  Term = rownames(coef_summary),
  Estimate = coef_summary[, "coef"],
  StdErr = coef_summary[, "robust se"],
  Lower = coef_summary[, "coef"] - 1.96 * coef_summary[, "robust se"],
  Upper = coef_summary[, "coef"] + 1.96 * coef_summary[, "robust se"],
  p_value = coef_summary[, "Pr(>|z|)"]
)

# Apply the renaming to the Term column
df_coef$Term <- terms_rename[df_coef$Term]

# Plotting using ggplot2
cofp_side<-ggplot(df_coef, aes(x = Estimate, y = Term)) +
  geom_point() +
  geom_errorbarh(aes(xmin = Lower, xmax = Upper), height = 0.2, color = ifelse(df_coef$p_val
  theme_minimal() +
  labs(title = "Coefficient Plot of Cox Model: Link",
        x = "Coefficient Value",
        y = "Variables")

```

Survival rates from Cox

```

# Define factor levels and labels without "Low Activity"
outcome_levels <- c("Peace Agreement", "Ceasefire", "Gov Victory", "Rebel Victory", "Actor C

# Convert outcome variable to factor with specified levels in the original dataset

```

```

episodes$outcomes <- factor(episodes$outcome, levels = c(1, 2, 3, 4, 6), labels = outcome_levels)

m2 <- coxph(Surv(time_to_recur_any, recur_any) ~ as.factor(outcomes) + pko_u + log_dur + cold_war)

m4 <- coxph(Surv(time_to_recur, recur_side) ~ as.factor(outcomes) + pko_u + log_dur + cold_war)

representative_data <- episodes %>%
  summarise(
    pko_u = median(as.numeric(pko_u), na.rm = TRUE),
    log_dur = mean(log_dur, na.rm = TRUE),
    cold_war = median(cold_war, na.rm = TRUE),
    log_gdp = mean(log_gdp, na.rm = TRUE),
    veto_u = median(veto_u, na.rm = TRUE),
    gov_war = median(gov_war, na.rm = TRUE),
    ter_war = median(ter_war, na.rm = TRUE),
    p_polity2 = mean(p_polity2, na.rm = TRUE),
    fe_etfra = mean(fe_etfra, na.rm = TRUE),
    W4 = mean(W4, na.rm = TRUE),
    log_pop = mean(log_pop, na.rm = TRUE),
    al_language2000 = mean(al_language2000, na.rm = TRUE),
    ps_original = median(ps_original, na.rm = TRUE),
    al_religion2000 = mean(al_religion2000, na.rm = TRUE),
    fe_cultdiv = mean(fe_cultdiv, na.rm = TRUE)
  )

# Expand the data to include different outcomes
expanded_data <- do.call(rbind, replicate(5, representative_data, simplify = FALSE))
expanded_data$outcomes <- factor(rep(outcome_levels, each = 1), levels = outcome_levels)

# Generate survival curves based on the Cox model for m2
surv_fits_any <- survfit(m2, newdata = expanded_data)

# Define colors for the plot
colors <- c("Peace Agreement" = "red", "Ceasefire" = "blue", "Gov Victory" = "green", "Rebel Victory" = "red")

# Plotting survival curves for Any Recurrence
plot_any <- ggsurvplot(
  surv_fits_any,
  data = expanded_data,
  pval = TRUE, # Show p-value for log-rank test
  conf.int = FALSE, # Show confidence intervals
  xlab = "Time in Days",

```

```

ylab = "Survival Probability",
ggtheme = theme_bw(),
xlim = c(0, 20000),
break.time.by = 1000,
risk.table = TRUE, # Show number at risk table
surv.scale = "percent",
legend.title = "Outcome",
legend.labs = levels(expanded_data$outcomes), # Automatically use factor labels
title = "Survival Probability Over Time for War Recurrence Based on ACD Conflict ID",
subtitle = "",
caption = "",
palette = colors,
size = 1.2, # Increase line width for better visibility
linetype = "solid" # Set line type to solid for all curves
)

```

Warning in .pvalue(fit, data = data, method = method, pval = pval, pval.coord = pval.coord,
This is a null model.

Warning: No shared levels found between `names(values)` of the manual scale and the
data's fill values.

Warning in (function (survsummary, times, survtable = c("cumevents",
"risk.table", : The length of legend.labs should be 1

```

# Customize the legend font size and axis titles for Any Recurrence plot
plot_any$plot <- plot_any$plot +
  theme(legend.text = element_text(size = 12), # Increase legend text size
        legend.title = element_text(size = 14), # Increase legend title size
        axis.title.x = element_text(size = 10), # X-axis title size
        axis.title.y = element_text(size = 10), # Y-axis title size
        axis.text.x = element_text(size = 10, angle = 45, hjust = 1), # X-axis text size and
        axis.text.y = element_text(size = 10), # Y-axis text size
        plot.title = element_text(size = 12)) # Plot title size

# Customize the risk table text for Any Recurrence plot
plot_any$table <- plot_any$table +
  theme(axis.text.x = element_text(size = 10, angle = 45, hjust = 1)) # Rotate risk table text

# Generate survival curves based on the Cox model for m4
surv_fits_side <- survfit(m4, newdata = expanded_data)

```



```
# Plotting survival curves for Side Recurrence
plot_side <- ggsurvplot(
  surv_fits_side,
  data = expanded_data,
  pval = TRUE, # Show p-value for log-rank test
  conf.int = FALSE, # Show confidence intervals
  xlab = "Time in Days",
  ylab = "Survival Probability",
  ggtheme = theme_bw(),
  xlim = c(0, 20000),
  break.time.by = 1000,
  risk.table = TRUE, # Show number at risk table
  surv.scale = "percent",
  legend.title = "Outcome",
  legend.labs = levels(expanded_data$outcomes), # Automatically use factor labels
  title = "Survival Probability Over Time for War Recurrence Based on Sufficient Linkage",
  subtitle = "",
  caption = "",
  palette = colors,
  size = 1.2, # Increase line width for better visibility
  linetype = "solid" # Set line type to solid for all curves
)
```

Warning in .pvalue(fit, data = data, method = method, pval = pval, pval.coord = pval.coord,
This is a null model.

Warning: No shared levels found between `names(values)` of the manual scale and the
data's fill values.

Warning in (function (survsummary, times, survtable = c("cumevents",
"risk.table", : The length of legend.labs should be 1

```
# Customize the legend font size and axis titles for Side Recurrence plot
plot_side$plot <- plot_side$plot +
  theme(legend.text = element_text(size = 12), # Increase legend text size
        legend.title = element_text(size = 14), # Increase legend title size
        axis.title.x = element_text(size = 10), # X-axis title size
        axis.title.y = element_text(size = 10), # Y-axis title size
        axis.text.x = element_text(size = 10, angle = 45, hjust = 1), # X-axis text size and
        axis.text.y = element_text(size = 10), # Y-axis text size)
```

```

    plot.title = element_text(size = 12))      # Plot title size

# Customize the risk table text for Side Recurrence plot
plot_side$table <- plot_side$table +
  theme(axis.text.x = element_text(size = 10, angle = 45, hjust = 1)) # Rotate risk table text

# Print the plots
print(plot_any)

```

Warning: No shared levels found between `names(values)` of the manual scale and the data's fill values.

Warning: No shared levels found between `names(values)` of the manual scale and the data's fill values.

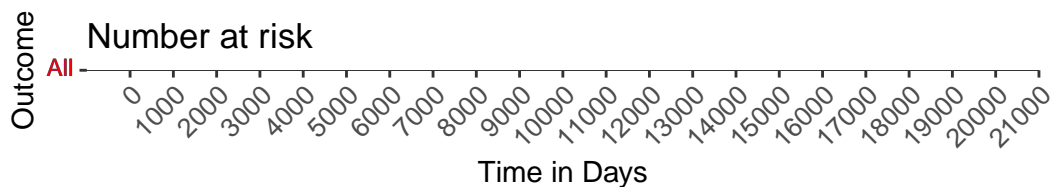
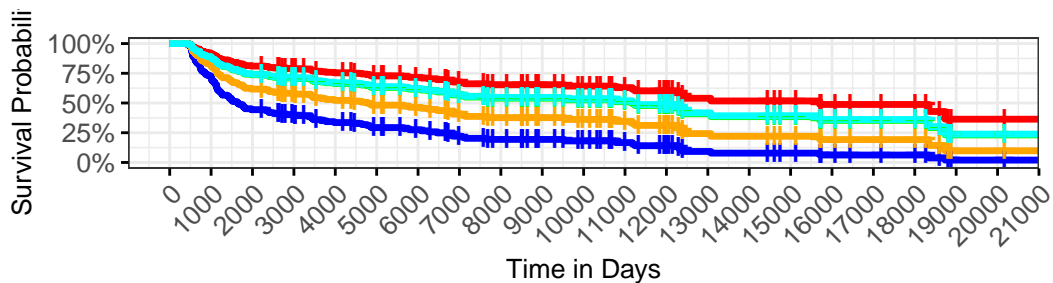
No shared levels found between `names(values)` of the manual scale and the data's fill values.

Warning: No shared levels found between `names(values)` of the manual scale and the data's colour values.

Warning: No shared levels found between `names(values)` of the manual scale and the data's fill values.

Survival Probability Over Time for War Recurrence Based on ACF

+ Peace Agreement
 + Ceasefire
 + Gov Victory
 + Rebel Victory



```
print(plot_side)
```

Warning: No shared levels found between `names(values)` of the manual scale and the data's fill values.

No shared levels found between `names(values)` of the manual scale and the data's fill values.

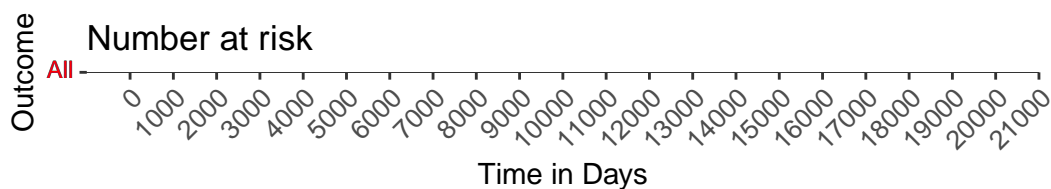
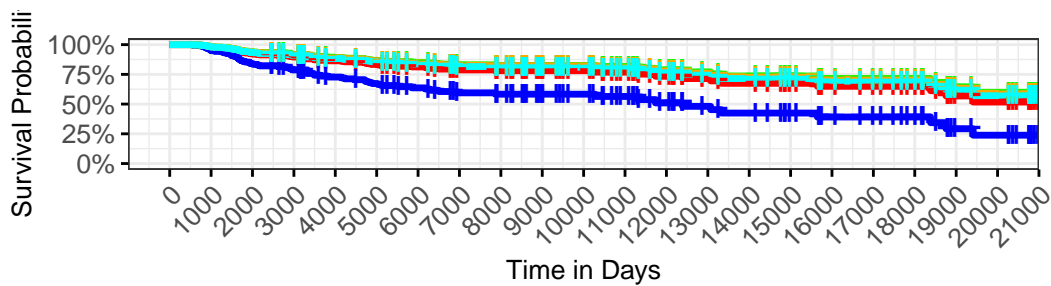
No shared levels found between `names(values)` of the manual scale and the data's fill values.

Warning: No shared levels found between `names(values)` of the manual scale and the data's colour values.

Warning: No shared levels found between `names(values)` of the manual scale and the data's fill values.

Survival Probability Over Time for War Recurrence Based on Suff

—+— Peace Agreement —+— Ceasefire —+— Gov Victory —+— Rebel Victory



RSF- Var Im-Any

```

# Install and load necessary packages
# install.packages("randomForestSRC")
# install.packages("timeROC")
# install.packages("caret")
library(randomForestSRC)
library(readxl)
library(pec)
library(timeROC)
library(survivalROC)
library(ggplot2)
library(expss)
library(survival)
library(dplyr)
library(caret)

# Assuming 'episodes' is already loaded

# Applying labels
episodes_a <- apply_labels(episodes,
                           time_to_recur_any = "Time to Recur Any",
                           recur_any = "War Recurrence Any",
                           dis = "Actor Ceases",
                           cease = "Ceasefire",
                           govvic = "Government Victory",
                           rebvic = "Rebel Victory",
                           lowac = "Low Activity",
                           W4 = "Coalition Size",
                           pko_u = "Peacekeeping Operations",
                           log_dur = "Log Duration",
                           cold_war = "Cold War",
                           p_polity2 = "Polity Score",
                           fe_etfra = "Ethnic Fractionalization",
                           log_gdp = "Log GDP per Capita",
                           log_pop = "Log Population",
                           al_language2000 = "Language Fractionalization",
                           ps_original = "Power-Sharing",
                           al_religion2000 = "Religion Fractionalization",
                           fe_cultdiv = "Cultural Diversity",
                           veto_u = "Veto Players",
                           gov_war = "War over Government",
                           ter_war = "War over Territory")

```

```

# Subsetting data and removing lowac
X <- subset(episodes_a, select = c(time_to_recur_any, recur_any, dis, cease, govvic, rebvic,
X <- na.omit(X)

# Ensure numeric variables are correctly formatted
X <- X %>%
  mutate(across(everything(), as.numeric))

# Define cross-validation folds
set.seed(123) # For reproducibility
folds <- createFolds(X$recur_any, k = 5, list = TRUE)

# Initialize a list to store variable importance scores
importance_list <- vector("list", length(folds))

# Perform cross-validation
for (i in seq_along(folds)) {
  # Define training and validation sets
  train_indices <- unlist(folds[-i])
  test_indices <- unlist(folds[i])

  train_data <- X[train_indices, ]
  test_data <- X[test_indices, ]

  # Train the Random Survival Forest model
  rsf <- rfsrc(Surv(time_to_recur_any, recur_any) ~ ., data = train_data, ntree = 1000, nodes

  # Store the variable importance scores
  importance_list[[i]] <- rsf$importance
}

# Aggregate variable importance scores
importance_df <- as.data.frame(do.call(cbind, importance_list))
importance_df$Variable <- rownames(importance_df)
rownames(importance_df) <- NULL
importance_df$Importance <- rowMeans(importance_df[, -ncol(importance_df)])
importance_df <- importance_df[, c("Variable", "Importance")]

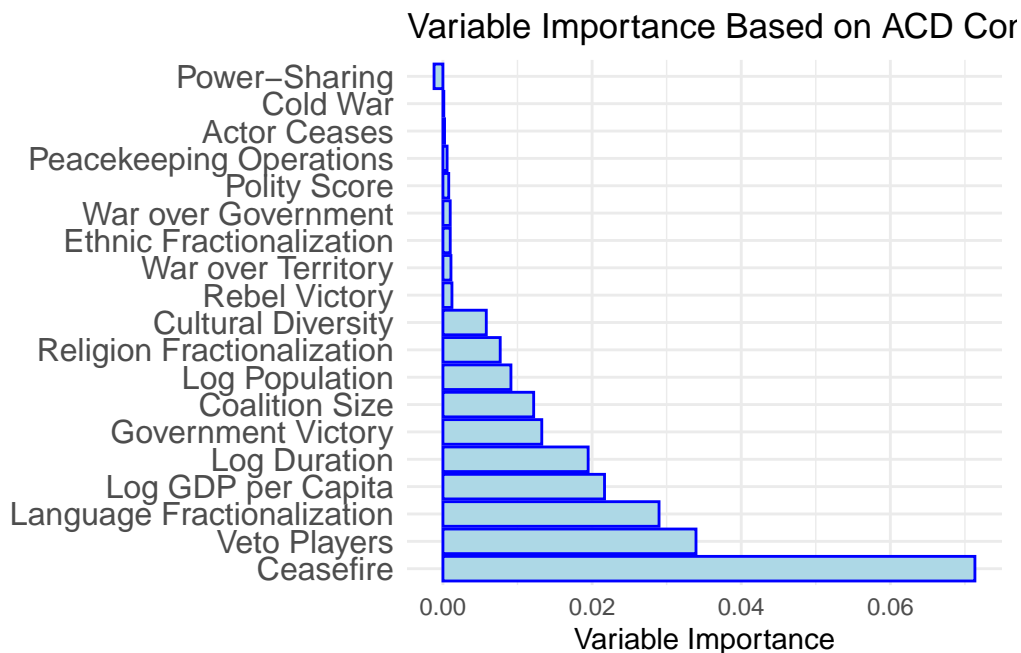
# Relabeling variables for plot
var_labels <- c("Actor Ceases", "Ceasefire", "Government Victory", "Rebel Victory", "Peaceke
names(var_labels) <- c("dis", "cease", "govvic", "rebvic", "pko_u", "log_dur", "cold_war", ".

```

```
# Adding variable labels
importance_df$Variable <- factor(importance_df$Variable, levels = names(var_labels), labels = var_labels)

# Order by importance
importance_df <- importance_df %>% arrange(desc(Importance))
importance_df$Variable <- factor(importance_df$Variable, levels = importance_df$Variable)

# Plotting using ggplot2
ggplot(importance_df, aes(x = Importance, y = Variable)) +
  geom_bar(stat = "identity", fill = "lightblue", color = "blue") +
  theme_minimal() +
  labs(title = "Variable Importance Based on ACD Conflict ID", x = "Variable Importance", y = "Variable") +
  theme(axis.text.y = element_text(size = 12))
```



RSF - Var Im - Link

```
# Install and load necessary packages
# install.packages("randomForestSRC")
# install.packages("timeROC")
# install.packages("caret")
```

```

library(randomForestSRC)
library(readxl)
library(pec)
library(timeROC)
library(survivalROC)
library(ggplot2)
library(expss)
library(survival)
library(dplyr)
library(caret)

# Assuming 'episodes' is already loaded

# Applying labels
episodes_a <- apply_labels(episodes,
                           time_to_recur = "Time to Recur",
                           recur_any = "War Recurrence",
                           dis = "Actor Ceases",
                           cease = "Ceasefire",
                           govvic = "Government Victory",
                           rebvic = "Rebel Victory",
                           W4 = "Coalition Size",
                           pko_u = "Peacekeeping Operations",
                           log_dur = "Log Duration",
                           cold_war = "Cold War",
                           p_polity2 = "Polity Score",
                           fe_etfra = "Ethnic Fractionalization",
                           log_gdp = "Log GDP per Capita",
                           log_pop = "Log Population",
                           al_language2000 = "Language Fractionalization",
                           ps_original = "Power-Sharing",
                           al_religion2000 = "Religion Fractionalization",
                           fe_cultdiv = "Cultural Diversity",
                           veto_u = "Veto Players",
                           gov_war = "War over Government",
                           ter_war = "War over Territory")

# Subsetting data
X <- subset(episodes_a, select = c(time_to_recur, recur_any, dis, cease, govvic, rebvic, pko_u, log_dur, cold_war, p_polity2, fe_etfra, log_gdp, log_pop, al_language2000, ps_original, al_religion2000, fe_cultdiv, veto_u, gov_war, ter_war))
X <- na.omit(X)

# Ensure numeric variables are correctly formatted

```

```

X <- X %>%
  mutate(across(everything(), as.numeric))

# Define cross-validation folds
set.seed(123) # For reproducibility
folds <- createFolds(X$recur_any, k = 5, list = TRUE)

# Initialize a list to store variable importance scores
importance_list <- vector("list", length(folds))

# Perform cross-validation
for (i in seq_along(folds)) {
  # Define training and validation sets
  train_indices <- unlist(folds[-i])
  test_indices <- unlist(folds[i])

  train_data <- X[train_indices, ]
  test_data <- X[test_indices, ]

  # Train the Random Survival Forest model
  rsf <- rfsrc(Surv(time_to_recur, recur_any) ~ ., data = train_data, ntree = 1000, nodesize

  # Store the variable importance scores
  importance_list[[i]] <- rsf$importance
}

# Aggregate variable importance scores
importance_df <- as.data.frame(do.call(cbind, importance_list))
importance_df$Variable <- rownames(importance_df)
rownames(importance_df) <- NULL
importance_df$Importance <- rowMeans(importance_df[, -ncol(importance_df)])
importance_df <- importance_df[, c("Variable", "Importance")]

# Relabeling variables for plot
var_labels <- c("Actor Ceases", "Ceasefire", "Government Victory", "Rebel Victory", "Peaceke
names(var_labels) <- c("dis", "cease", "govvic", "rebvic", "pko_u", "log_dur", "cold_war", "

# Adding variable labels
importance_df$Variable <- factor(importance_df$Variable, levels = names(var_labels), labels =

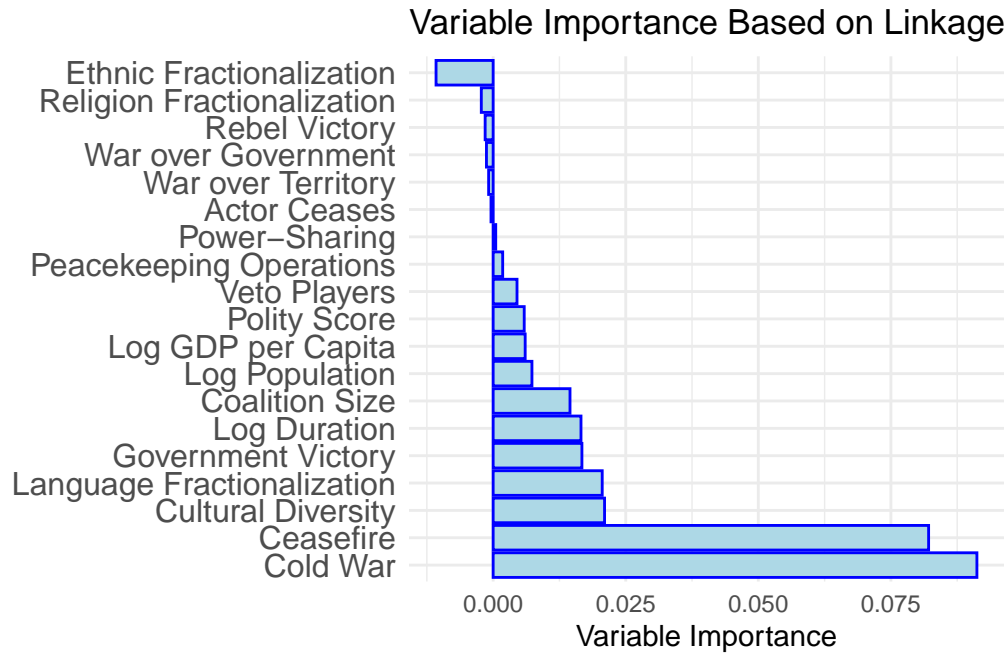
# Order by importance
importance_df <- importance_df %>% arrange(desc(Importance))

```



```
importance_df$Variable <- factor(importance_df$Variable, levels = importance_df$Variable)

# Plotting using ggplot2
ggplot(importance_df, aes(x = Importance, y = Variable)) +
  geom_bar(stat = "identity", fill = "lightblue", color = "blue") +
  theme_minimal() +
  labs(title = "Variable Importance Based on Linkage", x = "Variable Importance", y = "") +
  theme(axis.text.y = element_text(size = 12))
```



RSF -Sur Probs - Any

```
# Install and load necessary packages
# install.packages("randomForestSRC")
# install.packages("ggplot2")
library(randomForestSRC)
library(ggplot2)
library(dplyr)
library(tidyr)

# Assuming 'episodes' is your dataset
```

```

# Subsetting and cleaning data
X <- subset(episodes, select = c(time_to_recur_any, recur_any, dis, cease, govvic, rebvic, p
                                cold_war, p_polity2, fe_etfra, log_gdp, log_pop, al_language,
                                al_religion2000, fe_cultdiv, veto_u, gov_war, ter_war))

X <- na.omit(X)
X <- X %>% mutate(across(everything(), as.numeric))

# Train the Random Survival Forest model on the entire dataset
fit <- rfsrc(Surv(time_to_recur_any, recur_any) ~ ., data = X, ntree = 1000, nodesize = 5, n

# Predict survival probabilities for the entire dataset
pred <- predict(fit, X, OOB = TRUE, type = "response")
survival_probs <- as.data.frame(pred$survival)

# Extract time points
time_points <- fit$time.interest
colnames(survival_probs) <- paste("Time", time_points, sep = "_")
survival_probs$id <- 1:nrow(survival_probs)
X$id <- 1:nrow(X)

# Convert survival probabilities to long format for plotting
survival_probs_long <- survival_probs %>%
  pivot_longer(cols = -id, names_to = "Time", values_to = "Probability") %>%
  mutate(Time = as.numeric(gsub("Time_", "", Time)))

# Combine with the original data to get the binary outcome variables
survival_probs_long <- left_join(survival_probs_long, X, by = "id")

# Function to calculate mean and confidence intervals
calculate_summary <- function(data) {
  data %>%
    group_by(Time) %>%
    summarise(
      Mean_Probability = mean(Probability, na.rm = TRUE),
      SD = sd(Probability, na.rm = TRUE),
      Lower_CI = Mean_Probability - qt(0.975, length(Probability)-1) * SD / sqrt(length(Probab
      Upper_CI = Mean_Probability + qt(0.975, length(Probability)-1) * SD / sqrt(length(Probab
    )
}

# Calculate summary statistics for each binary outcome variable
dis_summary <- calculate_summary(filter(survival_probs_long, dis == 1))

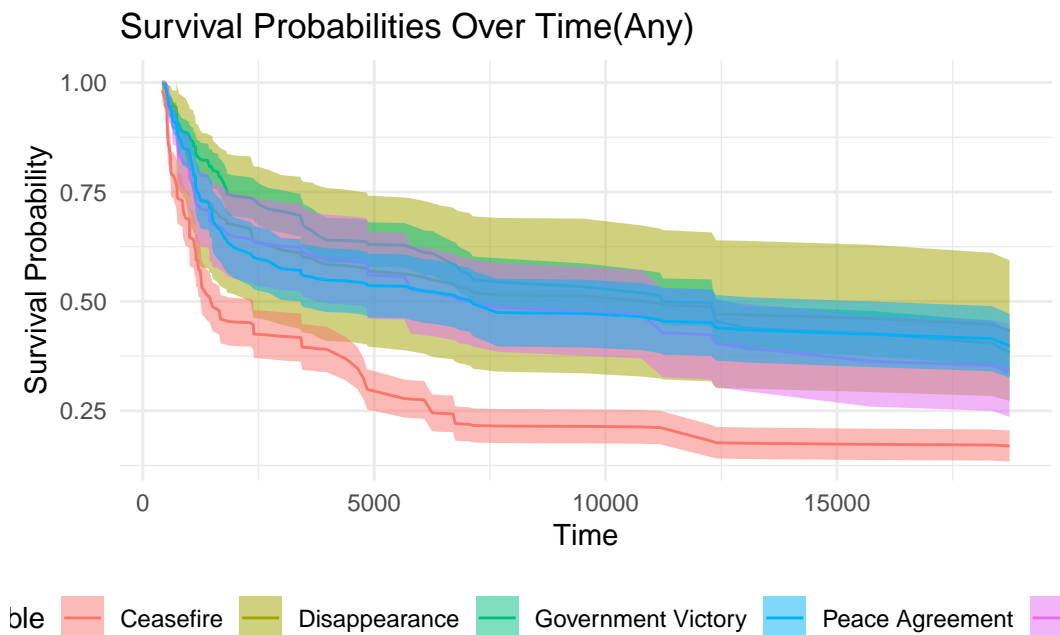
```

```

cease_summary <- calculate_summary(filter(survival_probs_long, cease == 1))
govvic_summary <- calculate_summary(filter(survival_probs_long, govvic == 1))
rebvic_summary <- calculate_summary(filter(survival_probs_long, rebvic == 1))
peace_summary <- calculate_summary(filter(survival_probs_long, peace == 1))

# Plot using ggplot2, one line per variable of interest with confidence intervals
ggplot() +
  geom_line(data = dis_summary, aes(x = Time, y = Mean_Probability, color = "Disappearance")) +
  geom_ribbon(data = dis_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Disappearance")) +
  geom_line(data = cease_summary, aes(x = Time, y = Mean_Probability, color = "Ceasefire")) +
  geom_ribbon(data = cease_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Ceasefire")) +
  geom_line(data = govvic_summary, aes(x = Time, y = Mean_Probability, color = "Government Victory")) +
  geom_ribbon(data = govvic_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Government Victory")) +
  geom_line(data = rebvic_summary, aes(x = Time, y = Mean_Probability, color = "Rebel Victory")) +
  geom_ribbon(data = rebvic_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Rebel Victory")) +
  geom_line(data = peace_summary, aes(x = Time, y = Mean_Probability, color = "Peace Agreement")) +
  geom_ribbon(data = peace_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Peace Agreement")) +
  labs(title = "Survival Probabilities Over Time(Any)", x = "Time", y = "Survival Probability") +
  theme_minimal() +
  theme(legend.position = "bottom")

```



RSF - Sur Probs- Side

```
# Install and load necessary packages
# install.packages("randomForestSRC")
# install.packages("ggplot2")
library(randomForestSRC)
library(ggplot2)
library(dplyr)
library(tidyr)

# Assuming 'episodes' is your dataset

# Subsetting and cleaning data
X <- subset(episodes, select = c(time_to_recur, recur_side, dis, cease, govvic, rebvic, peace,
                                cold_war, p_polity2, fe_etfra, log_gdp, log_pop, al_language,
                                al_religion2000, fe_cultdiv, veto_u, gov_war, ter_war))

X <- na.omit(X)
X <- X %>% mutate(across(everything(), as.numeric))

# Train the Random Survival Forest model on the entire dataset
fit <- rfsrc(Surv(time_to_recur, recur_side) ~ ., data = X, ntree = 1000, nodesize = 5, nspl = 10)

# Predict survival probabilities for the entire dataset
pred <- predict(fit, X, OOB = TRUE, type = "response")
survival_probs <- as.data.frame(pred$survival)

# Extract time points
time_points <- fit$time.interest
colnames(survival_probs) <- paste("Time", time_points, sep = "_")
survival_probs$id <- 1:nrow(survival_probs)
X$id <- 1:nrow(X)

# Convert survival probabilities to long format for plotting
survival_probs_long <- survival_probs %>%
  pivot_longer(cols = -id, names_to = "Time", values_to = "Probability") %>%
  mutate(Time = as.numeric(gsub("Time_", "", Time)))

# Combine with the original data to get the binary outcome variables
survival_probs_long <- left_join(survival_probs_long, X, by = "id")

# Function to calculate mean and confidence intervals
calculate_summary <- function(data) {
```

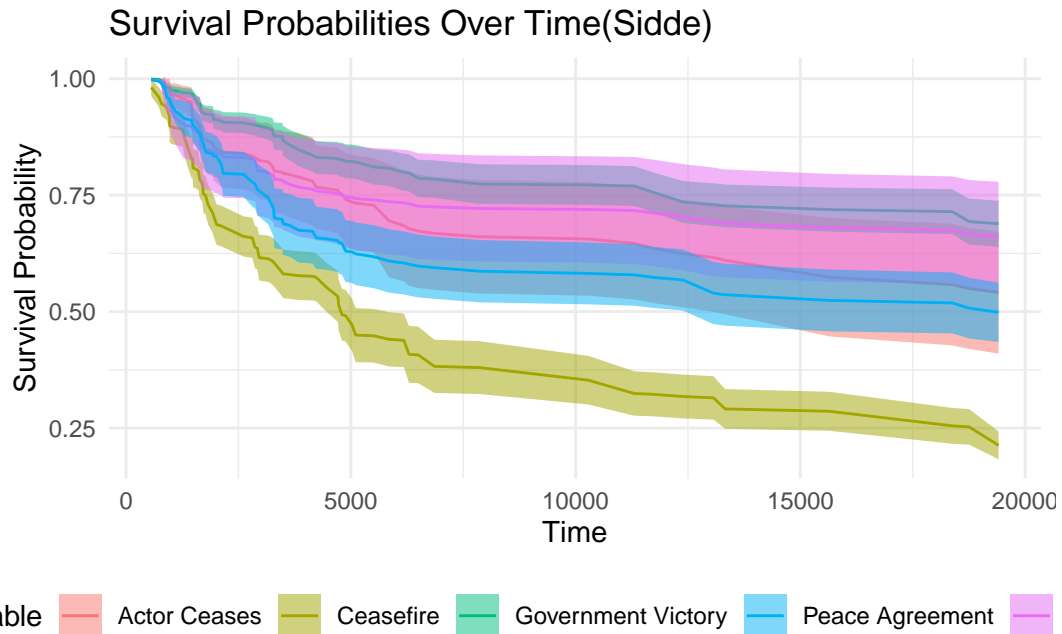
```

data %>%
  group_by(Time) %>%
  summarise(
    Mean_Probability = mean(Probability, na.rm = TRUE),
    SD = sd(Probability, na.rm = TRUE),
    Lower_CI = Mean_Probability - qt(0.975, length(Probability)-1) * SD / sqrt(length(Probability)),
    Upper_CI = Mean_Probability + qt(0.975, length(Probability)-1) * SD / sqrt(length(Probability))
  )
}

# Calculate summary statistics for each binary outcome variable
dis_summary <- calculate_summary(filter(survival_probs_long, dis == 1))
cease_summary <- calculate_summary(filter(survival_probs_long, cease == 1))
govvic_summary <- calculate_summary(filter(survival_probs_long, govvic == 1))
rebvic_summary <- calculate_summary(filter(survival_probs_long, rebvic == 1))
peace_summary <- calculate_summary(filter(survival_probs_long, peace == 1))

# Plot using ggplot2, one line per variable of interest with confidence intervals
ggplot() +
  geom_line(data = dis_summary, aes(x = Time, y = Mean_Probability, color = "Actor Ceases"))
  geom_ribbon(data = dis_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Actor Ceases"))
  geom_line(data = cease_summary, aes(x = Time, y = Mean_Probability, color = "Ceasefire"))
  geom_ribbon(data = cease_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Ceasefire"))
  geom_line(data = govvic_summary, aes(x = Time, y = Mean_Probability, color = "Government Victory"))
  geom_ribbon(data = govvic_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Government Victory"))
  geom_line(data = rebvic_summary, aes(x = Time, y = Mean_Probability, color = "Rebel Victory"))
  geom_ribbon(data = rebvic_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Rebel Victory"))
  geom_line(data = peace_summary, aes(x = Time, y = Mean_Probability, color = "Peace Agreement"))
  geom_ribbon(data = peace_summary, aes(x = Time, ymin = Lower_CI, ymax = Upper_CI, fill = "Peace Agreement"))
  labs(title = "Survival Probabilities Over Time(Sidde)", x = "Time", y = "Survival Probabilities")
  theme_minimal() +
  theme(legend.position = "bottom")

```



Causal Identification

Synthetic Control

```
# Create a subset with only the relevant variables
episodes_control <- subset(episodes, select = c("episode_id", "year", "recur_any",
                                                "log_gdp", "log_pop", "p_polity2",
                                                "fe_etfra", "log_dur", "cease"))
```

```
# Remove rows with missing values
episodes_control <- na.omit(episodes_control)
```

```
# Check the structure of the dataset
str(episodes_control)
```

```
tibble [184 x 9] (S3: tbl_df/tbl/data.frame)
 $ episode_id: num [1:184] 4 17 18 27 31 34 40 48 51 59 ...
 $ year      : num [1:184] 1968 1996 1998 1990 2012 ...
 $ recur_any : num [1:184] 0 1 1 0 1 1 1 0 1 1 ...
 $ log_gdp   : num [1:184] 5.28 7.2 6.91 7.27 7.06 ...
```

```
$ log_pop      : num [1:184] 15.3 18.1 18.1 15.2 17.7 ...
$ p_polity2    : num [1:184] -4 8 8 2 -3 -4 -6 -5 8 -5 ...
$ fe_etfra     : num [1:184] 0.743 0.161 0.161 0.132 0.522 ...
$ log_dur      : num [1:184] 5.29 9.16 3.61 0 8.37 ...
$ cease        : num [1:184] 0 1 1 0 1 0 0 0 0 1 ...
- attr(*, "na.action")= 'omit' Named int [1:70] 1 2 3 5 6 7 8 9 12 13 ...
..- attr(*, "names")= chr [1:70] "1" "2" "3" "5" ...
```

```
# Check for missing values again to ensure they've been removed
sum(is.na(episodes_control))
```

```
[1] 0
```

Matching