

Confounding of Selection and Influence

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- There are important challenges in conducting causal inference on contagion effects in observational data.
- (Fyfe and Desmarais 2024) show how we can use the “split-haves” test, robust to confounding, and apply it to studies of contagion effects.
- In this tutorial I will go over the method and several replication examples.

The split halves test

- Observational data are subject to confounding when identifying contagion/influence effects because of the co-existence of homophily and influence.
- The SH test isolates the impact of contagion by assuming the pre-existence of a network in the data without conditioning on it.
 1. Test data and adjust it for non-stationarity.
 2. Randomly split observational time-series cross-section data into two halves based on node (country in country-year data).
 3. Calculate mean values for each half for every time period.
 4. Run regression setting time t means as the dependent variable and $t - 1$ means of each half as independent variables.
 5. Perform steps 1 – 3 N times to recover a mean and p -value that indicate whether contagion is present or not.
 6. Contagion signal is the average value of the estimated relationship between the mean value of the first half at time t with the mean value of the second half at time $t - 1$, conditional on the mean value of the first half at $t - 1$. In a way, it is the relationship between both halves at different time points.
 7. The p -value is calculated as the minimum of two proportions, the proportion of times the contagion signal is > 0 and the proportion of times when it is < 0 . We obtain the p -value by multiplying the minimum proportion by 2 for a two-tailed test of whether there is contagion in the data.
 8. The estimate of general contagion tells us the average effect of a one-unit increase in the outcome value of any other node in the following year.

Applying the split-halves test

- I will use three of the replication examples in (Fyfe and Desmarais 2024) to illustrate the use of the split-halves test and its impact on results of previous studies.

Confirmation of Contagion: Conflict Onset

- The first replication is of (Buhaug and Gleditsch 2008), who find that there is a neighborhood effect of armed conflict.
- DV is binary indicator of conflict onset and a three-level ordinal variable indicating the type of conflict.
- This is an example where the SH test confirms the author's main results of contagion.

```
## Buhaug and Gleditsch
# Packages
library(dplyr)
library(maditr)
library(ggplot2)
library(tidyr)
library(ggplot2)
library(haven)
library(ContagionTest) # can download from GitHub
# devtools::install_github("rebekahfyfe/ContagionTest")

# Data
d <- read.table("conflict.tab", header = T)

# Removing duplicate rows
d <- d[-4439,]
d <- d[-5014,]

# Selecting necessary columns (country, date, DV)
d1 <- d %>%
  select("abbrev", "year", "allons3") %>%
  pivot_wider(names_from = year, values_from = allons3)
d1 <- as.data.frame(d1)

# Changing null values to NAs (treating as missing data)
d1[d1 == 'NULL'] <- NA

# Formatting for split-halves test
d1 <- STFormat(d1)

# Running split-halves contagion test
simmodels <- lag_pc_test(d1, 1000, 3, T, 0.1,
  lagWin = 1, missingData = T)
```

```
## [1] "Did not take 1st difference"
```

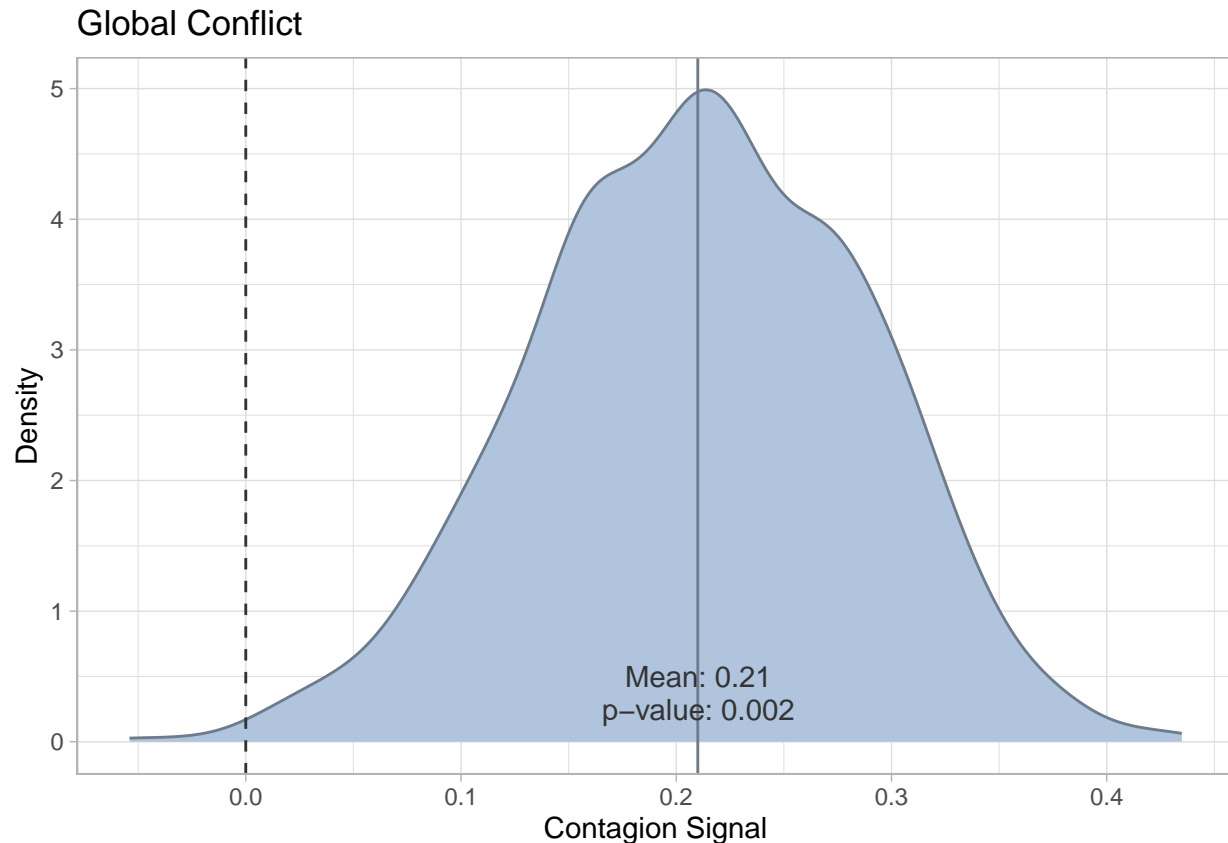
```
# Summary of models
simmodels <- as.data.frame(simmodels)
names(simmodels) <- c("intercept", "t-1coef", "counterpart")

# Calculate contagion signal
xmean <- mean(simmodels$counterpart) ## input this in the plot
xmean <- round(xmean, digits = 4)

# P-value of the signal (proportion of results < 0)
```

```
pval <- sum(simmodels$counterpart < 0) / 1000 ## pvalue
pval <- round(pval, digits = 3)

# Density graph of results
density_graph(simmodels, 1000, xmean, 0.5, xmean, 0.25,
              title = "Global Conflict")
```



- Using 1,000 random splits, the SH returns a positive contagion signal and a p -value < 0.01 .
- The expected prevalence of civil conflict onset in one country increases by approximately 0.02 for every 0.1 increase in lagged civil war prevalence among the other countries.
- The results support the findings of Buhaug and Gleditsch.

Challenging Non-Contagion: Pro-Democracy Protests

- The second replication is of (Brancati and Lucardi 2018), who find that, contrary to some strands of literature, pro-democracy protests do not diffuse to other countries.
- DV is protest onset.
- This is an example where the SH test challenges the author's results of no contagion.

```
## Brancati and Lucardi
# Violence in the Netherlands data, from Braun 2011
d <- as.data.frame(read.delim("violneth.tab"))

# Selecting only the necessary columns
d <- d[, c(3, 5, 21:55)]

# Creating a single variable for dates
T1 <- c(rep(1, 30), rep(0, 1065))
d$T1 <- rep(T1, 474)
date <- seq(as.Date("2001-01-01"),
            as.Date("2003-12-31"), by = "days")
d$date <- rep(date, 474)

# Changing to wide format
d <- d %>% select(number, date, countinc) %>%
  pivot_wider(names_from = date, values_from = countinc)

# Formatting to be used with contagion test
d <- STFormat(d)

# Running parallel contagion test
simmodNVio <- lag_pc_test(d, 1000, 1, T, 0.05)
```

```
## [1] "Took 1st difference"
```

```
summary(simmodNVio)
```

```
##      (Intercept)      c(j1mean.tm1, j2mean.tm1) c(j2mean.tm1, j1mean.tm1)
## Min.   :0.0003944 Min.   :0.0979           Min.   :0.05258
## 1st Qu.:0.0004100 1st Qu.:0.1482           1st Qu.:0.11515
## Median :0.0004112 Median :0.1638           Median :0.13170
## Mean   :0.0004115 Mean   :0.1646           Mean   :0.13075
## 3rd Qu.:0.0004129 3rd Qu.:0.1802           3rd Qu.:0.14722
## Max.   :0.0004291 Max.   :0.2437           Max.   :0.19829
```

```
# Creating a dataframe with results
simmodNVio <- as.data.frame(simmodNVio)
names(simmodNVio) <- c("intercept", "t-1coef", "counterpart")

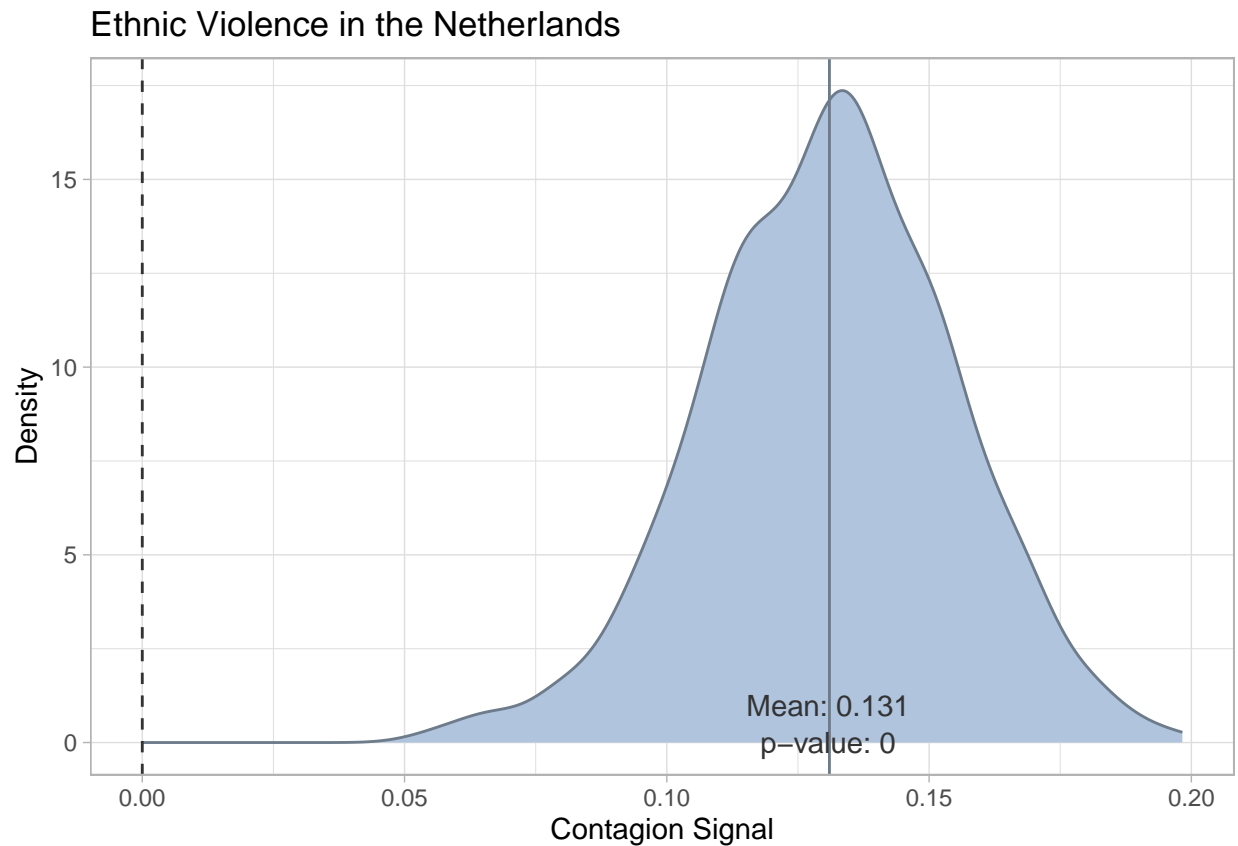
# Calculating mean (contagion signal)
mean <- mean(simmodNVio$counterpart) ## input this in the plot below
(mean <- round(mean, digits = 10))
```

```
## [1] 0.1307509
```

```
# Significance of the signal, proportion of means less than 0
pval <- sum(simmodNVio$counterpart < 0) / 1000 ## pvalue
(pval <- round(pval, digits = 3))
```

```
## [1] 0
```

```
# Density graph of results
density_graph(simmodNVio, 1000, mean, 1, mean, 0,
              title = "Ethnic Violence in the Netherlands")
```



- While Brancati and Lucardi find no contagion effect, the SH test shows that there is indeed statistical evidence of contagion.

Challenging Contagion: Civilian Targeting

- The third replication is of (Lis, Spagat, and Lee 2021), who find that there is a spillover effect that results in the spreading of violence against civilians by armed actors.
- DV is the Civilian Targeting Index, which ranges from 0 to 1000.
- This is an example where the SH test challenges the author's results of contagion.

```
# Data from Lis, Spagat, and Lee
d <- read_dta("civiliantargeting.dta")

# Selecting necessary columns (country, date, DV)
d <- d %>%
  select(actor_id, year, cti)

# Changing to wide format
d <- dcast(d, actor_id ~ year, value.var = "cti")
```

```

# Formatting to be used with split-halves contagion test
d <- STFormat(d)

# running contagion test
lslres <- lag_pc_test(d, 1000, 1, T, 0.05, 1, F)

## [1] "Took 1st difference"

summary(lslres)

##      (Intercept)      c(j1mean.tm1, j2mean.tm1) c(j2mean.tm1, j1mean.tm1)
##   Min.      :-1.7334   Min.      :-0.58013      Min.      :-0.28110
##   1st Qu.   :-1.2921   1st Qu.   :-0.40000      1st Qu.   :-0.02571
##   Median    :-1.2359   Median    :-0.34034      Median    : 0.04104
##   Mean      :-1.2402   Mean      :-0.33310      Mean      : 0.03548
##   3rd Qu.   :-1.1891   3rd Qu.   :-0.26924      3rd Qu.   : 0.10138
##   Max.      :-0.6576   Max.      :-0.02375      Max.      : 0.27829

# Creating a data frame with results
lslres <- as.data.frame(lslres)
names(lslres) <- c("intercept", "t-1coef", "counterpart")

# Calculating mean (contagion signal)
lslresmean <- mean(lslres$counterpart) ## input this in the plot below
(lslresmean <- round(lslresmean, digits = 10))

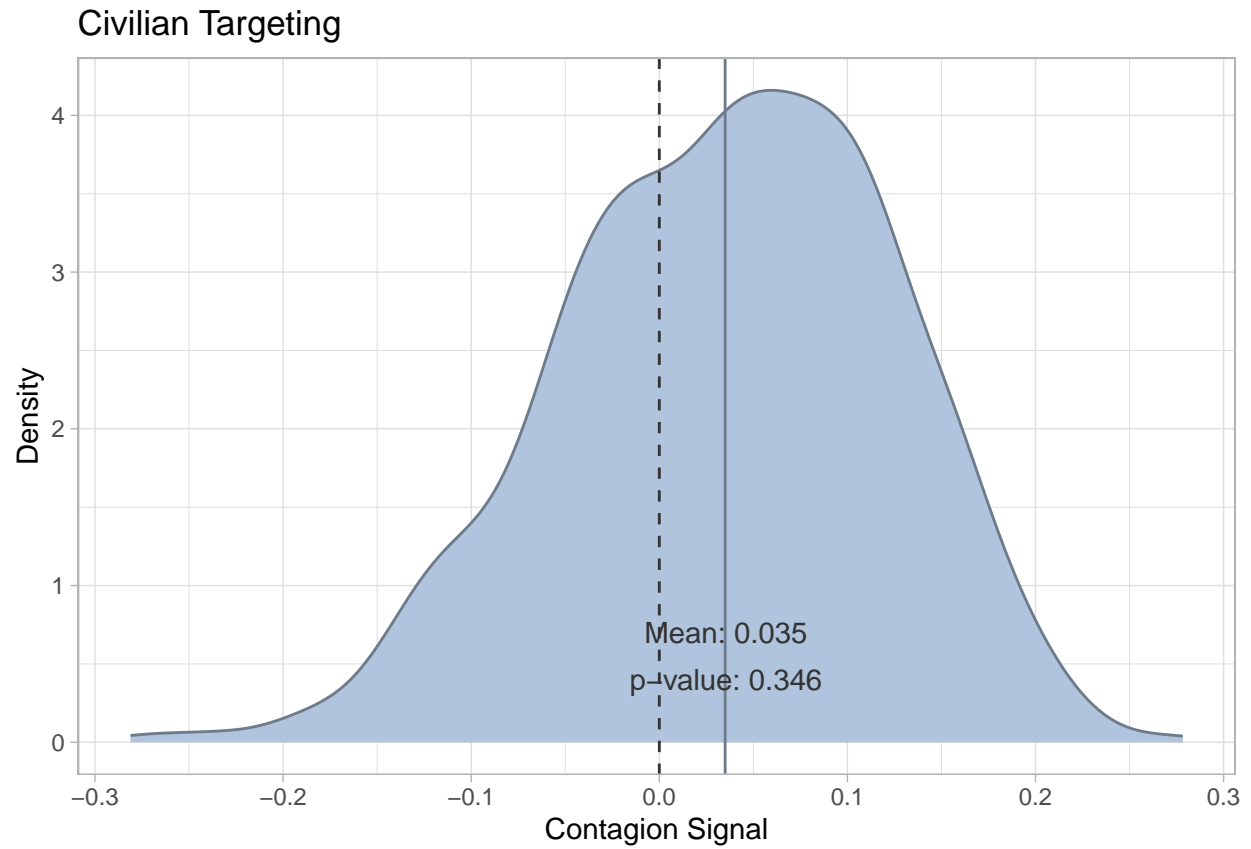
## [1] 0.03548029

# Significance of the signal, proportion of means less than 0
lslrespval <- sum(lslres$counterpart < 0) / 1000 ## pvalue
(lslrespval <- round(lslrespval, digits = 3))

## [1] 0.346

# Density graph of results
density_graph(lslres, 1000, lslresmean, 0.7, lslresmean, 0.4,
              title = "Civilian Targeting")

```



- While Lis, Spagat and Lee find a contagion effect, the SH test shows that there is no statistical evidence of contagion.

References

- Brancati, Dawn, and Adrián Lucardi. 2018. “Why Democracy Protests Do Not Diffuse.” *Journal of Conflict Resolution* 63 (10): 2354–89. <https://doi.org/10.1177/0022002718815957>.
- Buhaug, Halvard, and Kristian Skrede Gleditsch. 2008. “Contagion or Confusion? Why Conflicts Cluster in Space.” *International Studies Quarterly* 52 (2): 215–33. <https://doi.org/10.1111/j.1468-2478.2008.00499.x>.
- Fyfe, Rebekah, and Bruce Desmarais. 2024. “Causal Evidence for Theories of Contagious Civil Unrest.” *International Studies Quarterly* 68 (4). <https://doi.org/10.1093/isq/sqae124>.
- Lis, Piotr, Michael Spagat, and Uih Ran Lee. 2021. “Civilian Targeting in African Conflicts: A Poor Actor’s Game That Spreads Through Space.” *Journal of Peace Research* 58 (5): 900–914. <https://doi.org/10.1177/0022343320961150>.