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 pints.
 - 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.
- First, here is the replication of the original study.

```
## Buhaug and Gleditsch
rm(list = ls())
# Libraries
library(ggplot2)
library(dplyr)
library(broom)
library(nnet)
library(dplyr)
library(haven)
# Original data
rep <- read_dta("hb_ksg_replication.dta")</pre>
# Model 1
model1 <- glm(allons3 ~ neighall + neighpol + I(neighpol^2) +</pre>
                neighlgdp + peaceall,
              data = rep, family = binomial(link = "logit"))
# Model 2
model2 <- glm(allons3 ~ ncivwar + neighpol + I(neighpol^2) +</pre>
                neighlgdp + peaceall,
              data = rep, family = binomial(link = "logit"))
# Model 3
model3 <- glm(allons3 ~ neighall + neighpol + I(neighpol^2) +</pre>
                neighlgdp + polity2l + I(polity21^2) +
              lgdp961 + lnpop + peaceall,
              data = rep, family = binomial(link = "logit"))
# Model 4
model4 <- glm(allons3 ~ ncivwar + neighpol + I(neighpol^2) +</pre>
                neighlgdp + polity21 + I(polity21^2) +
              lgdp961 + lnpop + peaceall,
              data = rep, family = binomial(link = "logit"))
# Table 2, conflict neighbors only
# Model 5
model5 <- glm(allons3 ~ lnblength + lndist + ethlink2 +</pre>
                lneighbref + pop_nc + nterr + lbd_cum +
                polity21 + I(polity21^2) + lgdp961 + lnpop +
                peaceall,
              data = subset(rep, ncivwar == 1),
              family = binomial(link = "logit"))
# Model 6
model6 <- glm(allons3 ~ lnblength + confbord + ethlink2 +</pre>
```

```
lneighbref + pop_nc + nterr + lbd_cum +
              polity21 + I(polity21^2) + lgdp961 + lnpop +
               peaceall,
              data = subset(rep, ncivwar == 1),
              family = binomial(link = "logit"))
# Model 7, multinomial logit of terr and gov conflicts
model7 <- multinom(mons3 ~ ethlink2 + lneighbref + nterr +</pre>
                   polity21 + I(polity21^2) +
                    lgdp961 + lnpop + peaceall,
                   data = subset(rep, ncivwar == 1))
## # weights: 30 (18 variable)
## initial value 2906.928116
## iter 10 value 747.699419
## iter 20 value 582.810436
## iter 30 value 550.835875
## final value 550.816856
## converged
# Summary
library(texreg)
screenreg(list(model1, model2, model3, model4))
```

# # ==========		.=======	.=======	
# #	Model 1	Model 2	Model 3	Model 4
# (Intercept)			-3.11 **	-3.31 **
#	(0.88)	(0.87)	(1.07)	(1.04)
# neighall	0.59 *		0.32	
#	(0.28)		(0.30)	
# neighpol	0.01	0.01	-0.00	0.00
#	(0.02)	(0.02)	(0.02)	(0.02)
# neighpol^2	-0.00	-0.00	0.00	0.00
#	(0.00)	(0.00)	(0.00)	(0.00)
# neighlgdp	-0.39 ***	-0.37 ***	-0.03	-0.02
#	(0.11)	(0.11)	(0.15)	(0.15)
# peaceall	-0.02 **	-0.02 **	-0.01 *	-0.01 *
#	(0.01)	(0.01)	(0.01)	(0.01)
# ncivwar		0.67 ***		0.48 **
#		(0.15)		(0.15)
# polity2l			0.00	0.01
#			(0.01)	(0.01)
<pre># polity21^2</pre>			-0.01 ***	-0.01 ***
#			(0.00)	(0.00)
# lgdp961			-0.30 *	-0.27 *
#			(0.12)	(0.12)
# lnpop			0.30 ***	0.28 ***
#			(0.05)	(0.05)
# # AIC	 1849.17	1832.01	1795.70	1786.60
# BIC	1889.93	1872.77	1863.64	1854.54

screenreg(list(model5, model6, model7))

	Model 1	Model 2	Model 3
(Intercept)	-3.83 **	-3.85 **	
	(1.37)	(1.38)	
lnblength	-0.04	-0.04	
	(0.10)	(0.10)	
lndist	-0.01		
	(0.04)		
ethlink2	0.61 *	0.62 *	
	(0.26)	(0.26)	
lneighbref	0.04	0.04	
	(0.02)	(0.02)	
pop_nc	-0.03	-0.03	
	(0.08)	(0.08)	
nterr	0.61 **	0.61 **	
1h.J	(0.20)	(0.20)	
lbd_cum	-0.05	-0.05	
nol:+::01	(0.05) 0.02	(0.05)	
polity21		0.02 (0.02)	
noli++21^0	(0.02) -0.01	-0.01	
polity21^2	(0.00)	(0.00)	
lgdp961	-0.23	-0.23	
1gap301	(0.12)	(0.12)	
lnpop	0.32 ***		
тпрор	(0.06)	(0.06)	
peaceall	-0.01	-0.01	
podobali	(0.01)	(0.01)	
confbord	(0.01)	0.01	
00111 D 0 1 Q		(0.26)	
1: (Intercept)		(0.20)	-6.13 **
1. (1mooloopo)			(1.80)
1: ethlink2			0.63
			(0.40)
1: lneighbref			-0.02
G			(0.04)
1: nterr			1.18 **
			(0.31)
1: polity2l			0.04
- •			(0.02)
1: polity21^2			-0.00
- •			(0.01)
1: lgdp961			-0.49 **
			(0.19)

```
## 1: lnpop
                                               0.54 ***
##
                                               (0.09)
## 1: peaceall
                                               -0.03 *
                                               (0.02)
##
## 2: (Intercept)
                                               -4.84 ***
                                               (1.45)
##
## 2: ethlink2
                                               0.68 *
                                               (0.32)
##
## 2: lneighbref
                                               0.06 *
##
                                               (0.02)
## 2: nterr
                                               0.14
##
                                               (0.25)
## 2: polity21
                                               -0.01
                                               (0.02)
##
## 2: polity21^2
                                               -0.01 *
##
                                               (0.00)
                                               0.05
## 2: lgdp961
##
                                               (0.15)
## 2: lnpop
                                               0.06
##
                                               (0.08)
## 2: peaceall
                                               -0.00
                                               (0.01)
## -----
## AIC
                   991.09
                                991.12
                                            1137.63
## BIC
                  1067.47
                               1067.50
                                            1243.49
## Log Likelihood -482.54
                               -482.56
                                            -550.82
## Deviance
                   965.09
                                965.12
                                             1101.63
## Num. obs.
                                             2646
                  2632
                                2632
## K
                                               3
## *** p < 0.001; ** p < 0.01; * p < 0.05
```

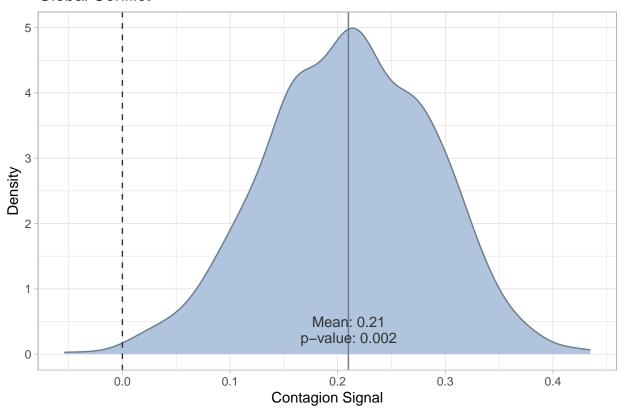
- The results from models 1 4 (full sample analysis) show that the presence of neighboring conflicts (neighall) and the number of neighboring civil wars (ncivwar) significantly increases the likelihood of conflict onset.
- The results from models 5 7 show that ethnic links to neighboring conflicts (ethlink2) and the number of territorial conflicts in neighboring countries (nterr) significantly increases the risk of conflict spillover.
- Below, by applying the split halves test, we see that this is an example where the SH test confirms the author's main results of contagion.

```
## Replication
# Libraries
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)</pre>
```

[1] "Did not take 1st difference"

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

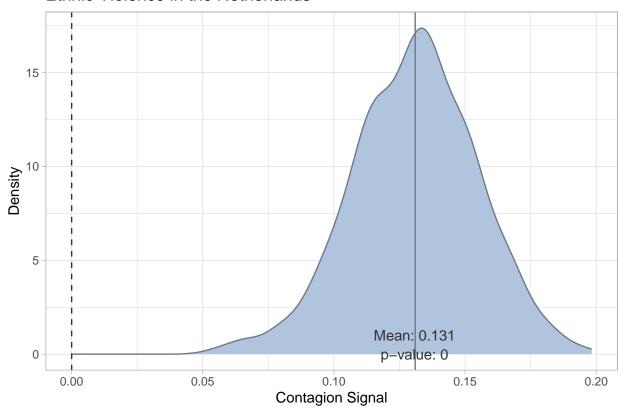
```
## Brancati and Lucardi
# Violence in the Netherlands data, from Braun 2011
d <- as.data.frame(read.delim(unzip("violneth.zip")))

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

# Creating a singe variable for dates
T1 <- c(rep(1, 30), rep(0, 1065))
d$TT1 <- rep(T1, 474)</pre>
```

```
date <- seq(as.Date("2001-01-01"),
            as.Date("2003-12-31"), by = "days")
d$date <- rep(date, 474)
# Changing to wide format
d <- d %>% select(number, date, countinc) %>%
 pivot_wider(names_from = date, values_from = countinc)
# Formatting to be used with congation test
d <- STFormat(d)</pre>
# Running parallel contagion test
simmodNVio <- lag_pc_test(d, 1000, 1, T, 0.05)
## [1] "Took 1st difference"
summary(simmodNVio)
##
     (Intercept)
                        c(j1mean.tm1, j2mean.tm1) c(j2mean.tm1, j1mean.tm1)
## Min.
           :0.0003944
                        Min. :0.0979
                                                  Min.
                                                        :0.05258
## 1st Qu.:0.0004100 1st Qu.:0.1482
                                                  1st Qu.:0.11515
## Median :0.0004112 Median :0.1638
                                                 Median: 0.13170
## Mean :0.0004115 Mean :0.1646
                                                 Mean :0.13075
## 3rd Qu.:0.0004129 3rd Qu.:0.1802
                                                  3rd Qu.:0.14722
## Max. :0.0004291
                              :0.2437
                                                  Max. :0.19829
                        Max.
# Creating a dataframe with results
simmodNVio <- as.data.frame(simmodNVio)</pre>
names(simmodNVio) <- c("intercept","t-1coef","counterpart")</pre>
# Calculating mean (contagion signal)
mean <- mean(simmodNVio$counterpart) ## input this in the plot below
(mean <- round(mean, digits = 10))</pre>
## [1] 0.1307509
\# Significance of the signal, proportion of means less than 0
pval <- sum(simmodNVio$counterpart < 0) / 1000 ## pvalue</pre>
(pval <- round(pval, digits = 3))</pre>
## [1] 0
# Density graph of results
density_graph(simmodNVio, 1000, mean, 1, mean, 0,
              title = "Ethnic Violence in the Netherlands")
```

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 authors find that there is a spillover effect that results in the spreading of violence against civilians by armed actors (Lis, Spagat, and Lee 2021).
- 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.

```
## Replication data
d <- read_dta("civiliantargeting.dta")

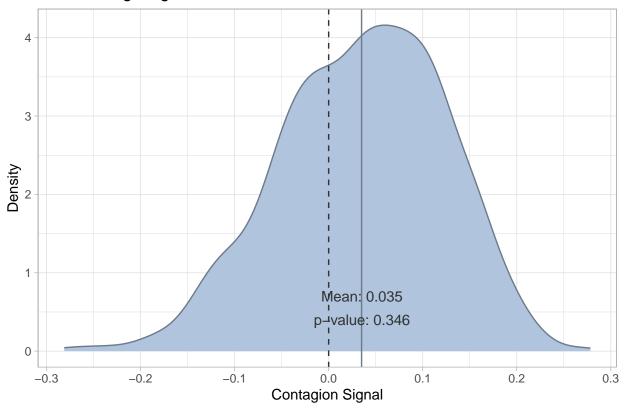
# Selecting necessary columns (country, date, DV)
d <- d %>%
    select(actor_id, year, cti)

# 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)</pre>
```

```
# running contagion test
lslres <- lag_pc_test(d, 1000, 1, T, 0.05, 1, F)</pre>
## [1] "Took 1st difference"
summary(lslres)
##
     (Intercept)
                     c(j1mean.tm1, j2mean.tm1) c(j2mean.tm1, j1mean.tm1)
          :-1.7334 Min. :-0.58013
## Min.
                                                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)</pre>
names(lslres) <- c("intercept","t-1coef","counterpart")</pre>
# Calculating mean (contagion signal)
lslresmean <- mean(lslres$counterpart) ## input this in the plot below
(lslresmean <- round(lslresmean, digits = 10))</pre>
## [1] 0.03548029
# Significance of the signal, proportion of means less than O
lslrespval <- sum(lslres$counterpart < 0) / 1000 ## pvalue</pre>
(lslrespval <- round(lslrespval, digits = 3))</pre>
## [1] 0.346
# Density graph of results
density_graph(lslres, 1000, lslresmean, 0.7, lslresmean, 0.4,
             title = "Civilian Targeting")
```

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.
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