Spatial Economics – Assignment 3

Gustav Pirich (h11910449) Gabriel Konecny (h11775903)

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Contents

The code that was used in compiling the assignment is available on GitHub at https://github.com/gustavpirich/spatial_econ/blob/main/03_assignment/03_assignmnet.Rmd.

Exercise A

```
## Reading layer `Sheet1' from data source
##   `/Users/gustavpirich/Desktop/GITHUB/spatial_econ/03_assignment/data/cigarettes/cigar_states.xls'
##   using driver `XLS'
```

- Estimate the demand model, and test for spatial dependence using the procedures discussed in class, and the provided weights matrix. Suppress any theoretical considerations

The demand model without spatial effects can by estimated by using the package plm:

Table 1:

	Dependent variable:
	logc
logp	−1.035***
	(0.042)
logy	0.529***
	(0.047)
Observations	1,380
R^2	0.394
Adjusted R^2	0.359
F Statistic	424.344*** (df = 2; 1303)
Note:	*p<0.1; **p<0.05; ***p<0.01

The demand for cigarettes depends negatively on the price of the cigarettes and positively on the real disposable income.

Alternatively, this could be also done by hand by including the dummies for both fixed effects and running OLS:

```
data_big <- cigarette_2var %>%
    fastDummies::dummy_cols(select_columns = "year") %>%
    fastDummies::dummy_cols(select_columns = "state") %>%
    select(c(logc, logp, logy, starts_with("year_") | starts_with("state_")))
OLS <- lm(logc ~ ., data_big)

# Get the summary of the model
summary_OLS <- summary(OLS)

# Extract coefficients and p-values
coefficients <- summary_OLS$coefficients[, 1]
p_values <- summary_OLS$coefficients[, 4]
variable_names <- rownames(summary_OLS$coefficients)

# Merge coefficients with stars denoting significance level and variable names
coefficients_with_stars <- paste0(variable_names, ": ", round(coefficients, 6),
ifelse(p_values <
    0.001, "***", ifelse(p_values < 0.01, "**", ifelse(p_values < 0.05, "*", ifelse(p_values
    <
    0.1, ".", ""))))

# Print the coefficients with stars
print(coefficients_with_stars)
```

```
##  [1] "(Intercept): 2.268869***" "logp: -1.034884***"
##  [3] "logy: 0.528543***"        "year_0: 0.193703***"
##  [5] "year_1: 0.155784***"      "year_2: 0.132723***"
##  [7] "year_3: 0.133168***"      "year_4: 0.120525***"
##  [9] "year_5: 0.120748***"      "year_6: 0.082984**"
```

```
## [11] "year_7: 0.081913**"       "year_8: 0.103591***"
## [13] "year_9: 0.11573***"       "year_10: 0.045043*"
## [15] "year_11: 0.009956"        "year_12: 0.00534"
## [17] "year_13: 0.034624"        "year_14: -0.016762"
## [19] "year_15: -0.001889"       "year_16: -0.078735**"
## [21] "year_17: -0.127222***"    "year_18: -0.17373***"
## [23] "year_19: -0.140931***"    "year_20: -0.068159**"
## [25] "year_21: -0.044138*"      "year_22: -0.04184."
## [27] "year_23: -0.030373"       "year_24: -0.037881*"
## [29] "year_25: -0.050626**"     "year_26: -0.057631***"
## [31] "year_27: -0.064987***"    "year_28: -0.067632***"
## [33] "state_1: 0.024896"        "state_2: -0.068988***"
## [35] "state_3: 0.094613***"     "state_4: -0.163163***"
## [37] "state_5: -0.065755*"      "state_6: 0.220373***"
## [39] "state_7: 0.050081*"       "state_8: 0.135211***"
## [41] "state_9: 0.018074"        "state_10: -0.143422***"
## [43] "state_11: -0.039591."     "state_12: 0.045096*"
## [45] "state_13: -0.055706**"    "state_14: -0.121456***"
## [47] "state_15: 0.23576***"     "state_16: 0.123416***"
## [49] "state_17: 0.161945***"    "state_18: -0.112063***"
## [51] "state_19: -0.009155"      "state_20: 0.046288*"
## [53] "state_21: -0.039464."     "state_22: 0.070078**"
## [55] "state_23: 0.010597"       "state_24: -0.069139***"
## [57] "state_25: -0.121315***"   "state_26: 0.29777***"
## [59] "state_27: 0.496985***"    "state_28: -0.058595*"
## [61] "state_29: -0.171643***"   "state_30: -0.02638"
## [63] "state_31: -0.119864***"   "state_32: -0.008334"
## [65] "state_33: 0.056935**"     "state_34: -0.033331."
## [67] "state_35: 0.121267***"    "state_36: -0.040171."
## [69] "state_37: -0.109306***"   "state_38: 0.042898*"
## [71] "state_39: 0.002169"       "state_40: -0.527422***"
## [73] "state_41: 0.155119***"    "state_42: -0.108599***"
## [75] "state_43: -0.171155***"   "state_44: 0.074774***"
## [77] "state_45: -0.057652**"
```

To save space, we merged coefficients with stars indicating p-values.

Where the point estimates of logp and logy are numerically identical to that of plm(). Thus we can see that using plm with “within” model and “twoways” effect is the way to go.

— which model would the classical specification search prefer?

We run all tests including individual and time fixed effects:

```
fm1 <- logc ~ logp + logy
```

```
##
##  LM test for spatial lag dependence
##
## data:  formula (within transformation)
## LM = 46.901, df = 1, p-value = 7.468e-12
## alternative hypothesis: spatial lag dependence
```

```
##
##  LM test for spatial error dependence
##
## data:  formula (within transformation)
## LM = 54.655, df = 1, p-value = 1.437e-13
## alternative hypothesis: spatial error dependence
```

```
##
##  Locally robust LM test for spatial lag dependence sub spatial error
##
## data:  formula (within transformation)
```

```
## LM = 1.1563, df = 1, p-value = 0.2822
## alternative hypothesis: spatial lag dependence

##
##  Locally robust LM test for spatial error dependence sub spatial lag
##
## data:  formula (within transformation)
## LM = 8.9106, df = 1, p-value = 0.002835
## alternative hypothesis: spatial error dependence
```

Allowing for spatial error, the spatial lag dependence is not significant. Thus this classical specification search suggests estimating SEM model.

- Estimate the model implied by the specification search and contrast the effect estimates with a SLX model specification:

Using the spml function, we can estimate SEM model by setting the spatial lag of dependent variable to false (lag=FALSE):

SLX model can be estimated by using OLS, i.e. by using plm() function from before where we include the spatially lagged values of explanatory variables:

```
slx_OLS <- function(W) {
    W_ <- kronecker(diag(nrow(unique(cigarette_2var[, "year"]))), W)
    W_logp <- W_ %*% data[, "logp"]
    W_logy <- W_ %*% data[, "logy"]
    data_slx <- cbind(data, W_logp, W_logy)
    slx <- plm(logc ~ logp + logy + W_logp + W_logy, data = data_slx, model = "within",
        effect = "twoways")
    return((slx))
}
```

Now, to compare both:

```
summary(sem)
```

Spatial panel fixed effects error model

Call: spml(formula = fm1, data = data, listw = lW, model = “within”, effect = “twoways”, lag = FALSE, spatial.error = “b”)

Residuals: Min. 1st Qu. Median 3rd Qu. Max. -0.43226537 -0.03661431 -0.00098061 0.04223100 0.50712114

Spatial error parameter: Estimate Std. Error t-value Pr(>|t|)
rho 0.24004 0.03267 7.3474 2.021e-13 ***

Coefficients: Estimate Std. Error t-value Pr(>|t|)
logp -1.004297 0.040011 -25.101 < 2.2e-16 ***logy 0.553849 0.049217 11.253 < 2.2e-16*** — Signif. codes: 0 ‘’ ***0.001*** ” *0.01* ” 0.05 ‘.’ 0.1 ’ ’ 1

```
print("SLX Model")
```

[1] “SLX Model”

```
stargazer(slx_OLS(W), type = "latex", header = FALSE)
```

The coefficient estimates above correspond to those reported in table 2 of Vega and Elhorst 2015. Both the SEM and SLX suggest on average a decrease in domestic demand for cigarettes of around 1% following an increase in domestic prices by 1%, ceteris paribus. Similarly both models suggest an increase in demand for cigarettes given an increase in domestic income. SLX coefficient estimate for this is higher. Additionally SLX model provides effects of lagged independ variables of cigarette demand. The SEM model does not provide any additional meaningful parameters since its parameter for spatial autocorrelation of errors is a nuisance parameter.

– What could be the rationale to use the SLX model as opposed to other spatial specifications?

The choice of the model should be based on theory. If we expect spillovers of exogenous variables and spillovers of endogenous variables or correlation of error terms is expected to be less pronounced or absent, it might be

Table 2:

	Dependent variable:
	logc
logp	−1.017***
	(0.042)
logy	0.608***
	(0.060)
W_logp	−0.220***
	(0.077)
W_logy	−0.219***
	(0.080)
Observations	1,380
R^2	0.400
Adjusted R^2	0.364
F Statistic	217.105*** (df = 4; 1301)
Note:	*p<0.1; **p<0.05; ***p<0.01

good idea to consider SLX. However, there are also reasons to consider SLX as a point of departure even if the theory is not clear about which model to choose. First, when using the SLX model, the estimation and interpretation of spillover effects is more straightforward. In contrast to endogenous models, SLX allows us to parametrize W and its parameters can be estimated. Strong limitation of SAR and SAC models is that the ratio between the spillover and direct effects of is same for every explanatory variable, which is unlikely to be true in many empirical studies. This is not the case for SLX model, where we get estimates of both by construction. Thus SLX can be considered more flexible in this perspective.

– Which type(s) of spillover(s) would you expect in the model for cigarette demand?

We would expect no endogenous spillover effects, since we dont believe that demand for cigarettes in a country should determine demand for cigarettes in neighboring country if there are no shortages. If however, the price of cigarettes in neighboring country is lower, people living near the border could be buying cigarettes in the neighboring country. This is called bootlegging effect. On other hand, if the income in our country increases, the opportunity costs of time of our citizens increase and they might then prefer to buy the more expensive cigarettes in home country instead. Thus we would expect spatially lagged exogenous variables to be relevant: An increase in price of cigarettes in neighboring country should have positive effect on the demand for cigarettes in home country. Increase in income of neighboring country should have negative effect on the demand for cigarettes in home.

- Re-run the analysis using a SLX model with a distance decay specification (between centers) for the weights matrix, i.e. $w_{ij} = \frac{1}{\gamma_{ij}}$ following Halleck Vega and Elhorst (2015). Use a distance decay parameter of γ = 3.

For this we first create a neighbour list, where all regions are neighbors by setting the distance threshold to big enough value and then compute distances using nbdists. They are then transformed as suggested above and scaled by maximum eigenvalue as in the paper.

```
coords <- cigar_states[, 2:3]

decay <- function(gamma) {
    distw <<- dnearneigh(coords, 0, 999999, row.names = cigar_states$Name, longlat = FALSE)
    dists <- nbdists(distw, coords, longlat = FALSE)
    ids <- lapply(dists, function(x) lapply(x, function(y) 1/(y^gamma)))
    lWd <- nb2listw(distw, glist = ids, style = "B", zero.policy = TRUE)
    Wd1 <- listw2mat(lWd)
    Wd <- Wd1/max(abs(eigen(Wd1)$values))
    return(Wd)
}
Wd <- decay(3)
```

```
colnames(Wd) <- rownames(Wd)
```

Using gamma similar to the paper, we get coefficient estimates which are very close to those in the paper:

```
stargazer(slx_OLS(Wd), type = "latex", header = FALSE)
```

Table 3:

	Dependent variable:
	logc
logp	−0.907***
	(0.038)
logy	0.652***
	(0.043)
W_logp	0.259***
	(0.061)
W_logy	−0.824***
	(0.049)
Observations	1,380
R^2	0.514
Adjusted R^2	0.484
F Statistic	343.407*** (df = 4; 1301)
Note:	*p<0.1; **p<0.05; ***p<0.01

While the interpretation is non-lagged independent variables is similar as in regressions above, we might want to focus on spatiall lagged variables. The model suggests that increase in price of cigarettes in neighboring regions increases demand for cigarettes in home region, ceteris paribus. This is consistent with the bootlegging behavior discussed in the paper, which predicts that some people might buy cigarettes in/from other regions if the price there is lower. The coefficient estimate for income in neighboring regions suggest lower demand for cigarettes if the real income in neighboring regions increases. This suggests that if people have higher income their opportunity cost of time is higher and they might prefer the convenience of buying cigarettes in their home region.

– Describe the network recovered by Halleck Vega and Elhorst (2015):

```
distw
```

```
## Neighbour list object:
## Number of regions: 46
## Number of nonzero links: 2070
## Percentage nonzero weights: 97.82609
## Average number of links: 45
```

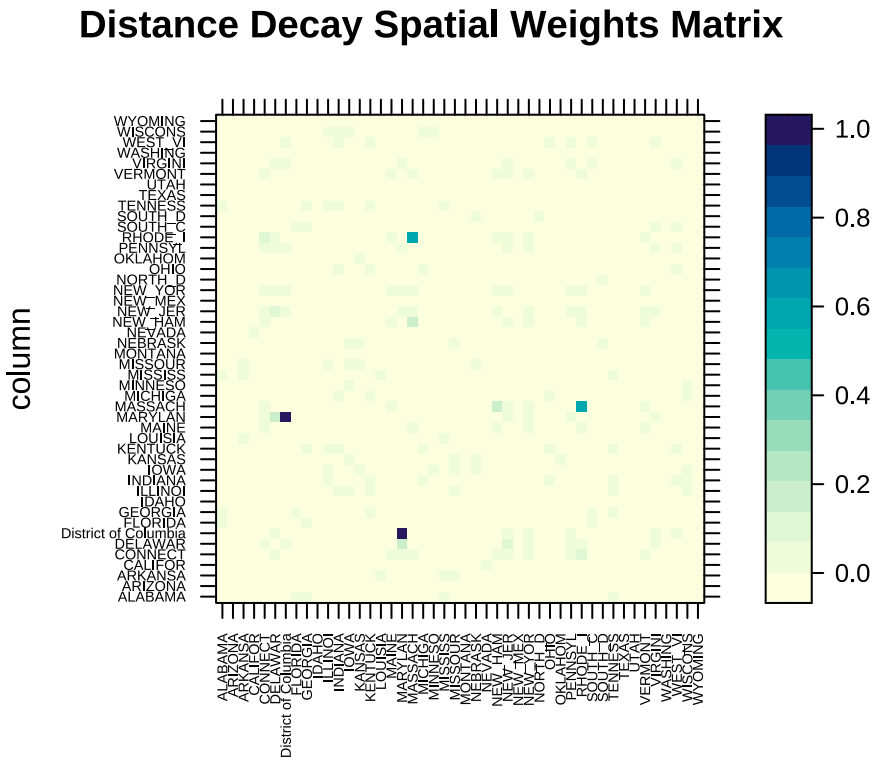
```
46 * 46 - 46
```

```
## [1] 2070
```

```
(46 * 46 - 46)/46 * 46
```

```
## [1] 2070
```

In this network all agents are connected, only the weights differ - which are given by the distance decay formula above. This corresponds to average number of links being 45 i.e. all regions except the region itself. Thus the percentage of nonzero weights simply gives the share of diagonal elements of a 46*46 matrix.



Below we visualize the spatial weights matrix:

Reminding ourself that all regions except the diagonal are neighbors, we can see that the definition of distance decay parameter used results in few strongly connected neighbors and many weaker connections.

– Who are the central agents and where do spillover effects occur?

We could define the most central agent as one, which has strongest connection to other regions overall. This could correspond to maximizing the nonlinear transformation of distance which we are using. Thus an agent is most central,if the sum of its weights to other agents is highest, i.e. if the sum of its row is highest.

```
##                          value
## MARYLAN              1.2524425
## District of Columbia 1.1439427
## MASSACH              0.7947205
## RHODE_I              0.7545230
## DELAWAR              0.3361666
## NEW_HAM              0.3198986
```

Maryland is most central agent. It could be that since multiple small regions from around Maryland are included, Maryland has very small distance to them and thus its inverse is large, contributing a lot to maximization of criterion made up above. Thus most central agent is one which has many close neighbors. From the table above we see that other important agents are District of Columbia, Massachusetts and Rhode Island.

We get similar results when using eigenvector centrality:

– What is the average partial effect of increasing income? We compute APE as follows: $\beta_{logy}+\theta_{logy}\frac{1}{N}\sum_j^J\sum_i^I w_{ij}$ where I is number of rows of W and J number of columns.

```
beta <- coef(slx_OLS(Wd))[["logy"]]
theta <- coef(slx_OLS(Wd))[["W_logy"]]
N <- nrow(Wd) * ncol(Wd)
beta + theta * sum(Wd)/N
```

```
## [1] 0.6496153
```

Thus, the average partial effect of increasing income of is very close to the effect of increasing income in home only. This is because using the specification of distance decay above, the links between the agents are mostly weak.

Table 4: Eigenvector Centrality

	state	ec
MARYLAN	MARYLAN	1.0000000
District of Columbia	District of Columbia	0.9873118
DELAWAR	DELAWAR	0.2168323
PENNSYL	PENNSYL	0.1330974
VIRGINI	VIRGINI	0.0898499
NEW_JER	NEW_JER	0.0442486
RHODE_I	RHODE_I	0.0060478
MASSACH	MASSACH	0.0057002
CONNECT	CONNECT	0.0050694
NEW_YOR	NEW_YOR	0.0049031
WEST_VI	WEST_VI	0.0034141
NEW_HAM	NEW_HAM	0.0029655
VERMONT	VERMONT	0.0018586
OHIO	OHIO	0.0016525
SOUTH_C	SOUTH_C	0.0015467
MAINE	MAINE	0.0008187
KENTUCK	KENTUCK	0.0007869
MICHIGA	MICHIGA	0.0006374
GEORGIA	GEORGIA	0.0005251
INDIANA	INDIANA	0.0005125
TENNESS	TENNESS	0.0003740
FLORIDA	FLORIDA	0.0002843
ALABAMA	ALABAMA	0.0002729
ILLINOI	ILLINOI	0.0001955
WISCONS	WISCONS	0.0001770
MISSISS	MISSISS	0.0001352
MISSOUR	MISSOUR	0.0001145
ARKANSA	ARKANSA	0.0000996
LOUISIA	LOUISIA	0.0000893
IOWA	IOWA	0.0000847
MINNESO	MINNESO	0.0000807
KANSAS	KANSAS	0.0000627
NEBRASK	NEBRASK	0.0000532
OKLAHOM	OKLAHOM	0.0000445
TEXAS	TEXAS	0.0000352
SOUTH_D	SOUTH_D	0.0000295
NORTH_D	NORTH_D	0.0000258
WYOMING	WYOMING	0.0000186
NEW_MEX	NEW_MEX	0.0000162
UTAH	UTAH	0.0000096
ARIZONA	ARIZONA	0.0000090
MONTANA	MONTANA	0.0000088
IDAHO	IDAHO	0.0000066
NEVADA	NEVADA	0.0000053
CALIFOR	CALIFOR	0.0000048
WASHING	WASHING	0.0000040

Exercise B

Unit of Observation

Cells	Stats
Min	9785713765 [m^2]
Mean	11698674731 [m^2]
Median	11904427037 [m^2]
Max	12364312149 [m^2]

The unit of observation are cells in a raster representing a 1×1 degree latitude longitude grid cover. At the equator this corresponds to a side length of 110 km. The areal extension of the cells varies with latitude. Further away from the equator, the area of a cell decreases. The table shows the area of the cells. The smallest cell has an area of about 97 857 km^2, the largest 12 364 km^2. On average the size of the cells is 11 900 km^2. Thus the gap between the smallest and largest cell amounts to about 20%. They differ because of the distortions induced by the projection of the surface of the earth, onto a 2 dimensional raster grid.

Interpretation of Coefficients

The binary contiguity matrix is *not* row normalized. This implies that the coefficients for the spatially lagged dependent and independent variables need to be interpreted as the direct impact of marginal changes in only *one* of the neighboring cells.

Replication

Dependent Variable:	ANY_EVENT_ACLED	
Model:	(1)	(2)
Variables		
SPEI4pg	0.0346***	0.0101
	(0.0053)	(0.0149)
L1_SPEI4pg	0.0026	0.0127
	(0.0049)	(0.0135)
L2_SPEI4pg	0.0104**	-0.0096
	(0.0049)	(0.0140)
GSmain_ext_SPEI4pg	-0.0338***	-0.0052
	(0.0088)	(0.0158)
L1_GSmain_ext_SPEI4pg	-0.0321***	-0.0429***
	(0.0081)	(0.0158)
L2_GSmain_ext_SPEI4pg	-0.0348***	-0.0239
	(0.0085)	(0.0156)
W_L2_GSmain_ext_SPEI4pg		-0.0027
		(0.0026)
W_L1_GSmain_ext_SPEI4pg		0.0020
		(0.0026)
W_GSmain_ext_SPEI4pg		-0.0058**
		(0.0027)
W_SPEI4pg		0.0044*
		(0.0023)
W_L1_SPEI4pg		-0.0017
		(0.0021)
W_L2_SPEI4pg		0.0036*
		(0.0021)
Fixed-effects		
as.factor(year)	Yes	Yes
Fit statistics		
Observations	35,042	35,042
R^2	0.18918	0.19568
Within R^2	0.18655	0.19307

Clustered (cell) standard-errors in parentheses
*Signif. Codes: ***: 0.01, **: 0.05, *: 0.1*

Table 5: Replication Model 3

	Estimate	Std. Error	t-value	Pr(>|t|)
lambda	0.0458	0.0010	45.3304	0
lag(ANY_EVENT_ACLED)	0.3373	0.0050	67.4562	0
SPEI4pg	−0.0017	0.0134	−0.1264	0.8994
L1_SPEI4pg	0.0138	0.0139	0.9895	0.3224
L2_SPEI4pg	−0.0160	0.0138	−1.1537	0.2486
GSmain_ext_SPEI4pg	0.0004	0.0141	0.0265	0.9789
L1_GSmain_ext_SPEI4pg	−0.0469	0.0147	−3.1841	0.0015
L2_GSmain_ext_SPEI4pg	−0.0110	0.0148	−0.7466	0.4553
W_GSmain_ext_SPEI4pg	−0.0028	0.0024	−1.1698	0.2421
W_L1_GSmain_ext_SPEI4pg	0.0066	0.0025	2.6670	0.0077
W_L2_GSmain_ext_SPEI4pg	−0.0011	0.0025	−0.4346	0.6639
W_SPEI4pg	0.0023	0.0021	1.1129	0.2658
W_L1_SPEI4pg	−0.0028	0.0021	−1.3146	0.1886
W_L2_SPEI4pg	0.0043	0.0021	2.0831	0.0372

Next, we create an adjacency matrix based on horizontal contiguity. To that end we write a function which takes

Table 6: Replication Model 4

	Estimate	Std. Error	t-value	Pr(>|t|)
lambda	0.0331	0.0011	29.7777	0
lag(ANY_EVENT_ACLED)	0.3455	0.0051	67.3891	0
SPEI4pg	−0.0003	0.0135	−0.0250	0.9800
L1_SPEI4pg	0.0182	0.0140	1.2950	0.1953
L2_SPEI4pg	−0.0139	0.0139	−0.9998	0.3174
GSmain_ext_SPEI4pg	0.0005	0.0140	0.0389	0.9690
L1_GSmain_ext_SPEI4pg	−0.0514	0.0146	−3.5081	0.0005
L2_GSmain_ext_SPEI4pg	−0.0050	0.0147	−0.3376	0.7357
W_GSmain_ext_SPEI4pg	−0.0035	0.0026	−1.3493	0.1772
W_L1_GSmain_ext_SPEI4pg	0.0071	0.0027	2.6270	0.0086
W_L2_GSmain_ext_SPEI4pg	−0.0017	0.0027	−0.6202	0.5352
W_SPEI4pg	0.0012	0.0022	0.5311	0.5953
W_L1_SPEI4pg	−0.0018	0.0023	−0.7959	0.4261
W_L2_SPEI4pg	0.0050	0.0023	2.1728	0.0298

the centroids and maximum distance for the contiguity cutoff as inputs and produces a neighboors list for each cells based on horizontal contiguity. We assign a neighboors status based on two conditions. First a cell needs to have the same latitude (or longitude in the case of vertical contiguity) and the centroid of the cell needs to be closer than a pre-specified distance cutoff. We check this condition for all cells. We then rerun specification 3 and 4 based on those contiguity matrices.

```
### Lets first create teh bianry contiguity matrix based on horizontal
### contiguity Extract centroids of the polygons if not already point data

# Calculate distances and filter horizontally contiguous points Define a
# function to find neighbors based on conditions
find_neighbors_horizontal <- function(centroids, max_distance = 150000, lat_tolerance =
0.01) {
    # Create a matrix to store distances
    distances <- st_distance(centroids)
    latitudes <- st_coordinates(centroids)[, 2]  # Extract latitudes

    # Initialize neighbors list
    neighbors_list <- vector("list", nrow(centroids))

    for (i in seq_len(nrow(centroids))) {
        # Find points within the same latitude band and within distance
        lat_condition <- abs(latitudes - latitudes[i]) < lat_tolerance
        dist_condition <- distances[i, ] < set_units(max_distance, "meters")

        # Combine conditions
        neighbor_ids <- which(lat_condition & dist_condition)

        # Exclude the point itself from its list of neighbors
        neighbor_ids <- neighbor_ids[neighbor_ids != i]

        # Store neighbor ids
        neighbors_list[[i]] <- as.integer(neighbor_ids)
    }

    return(neighbors_list)
}

find_neighbors_vertical <- function(centroids, max_distance = 150000, lon_tolerance = 0.01)
{
    # Create a matrix to store distances
```

```
    distances <- st_distance(centroids)
    longitudes <- st_coordinates(centroids)[, 1]  # Extract longitudes

    # Initialize neighbors list
    neighbors_list <- vector("list", nrow(centroids))

    for (i in seq_len(nrow(centroids))) {
        # Find points within the same longitude band and within distance
        lon_condition <- abs(longitudes - longitudes[i]) < lon_tolerance
        dist_condition <- distances[i, ] < set_units(max_distance, "meters")

        # Combine conditions
        neighbor_ids <- which(lon_condition & dist_condition)

        # Exclude the point itself from its list of neighbors
        neighbor_ids <- neighbor_ids[neighbor_ids != i]

        # Store neighbor ids
        neighbors_list[[i]] <- as.integer(neighbor_ids)
    }

    return(neighbors_list)
}

# Apply the function
neighbors_vertical <- find_neighbors_vertical(geoconflict_main_weights)

neighbors_horizontal <- find_neighbors_horizontal(geoconflict_main_weights)



# Number of elements
n <- length(neighbors_vertical)

# Initialize an adjacency matrix with 0s
adj_matrix_vertical <- matrix(0, nrow = n, ncol = n)
adj_matrix_horizontal <- matrix(0, nrow = n, ncol = n)

# Populate the adjacency matrix
for (i in seq_len(n)) {
    # Ensure indices are within the valid range
    valid_indices_vertical <- neighbors_vertical[[i]][neighbors_vertical[[i]] <=
        n]
    valid_indices_horizontal <- neighbors_horizontal[[i]][neighbors_horizontal[[i]] <=
        n]

    # Set matrix elements to 1
    adj_matrix_vertical[i, valid_indices_vertical] <- 1
    adj_matrix_horizontal[i, valid_indices_horizontal] <- 1
}

# Turn the matrices into listw objects
vertical_listw <- mat2listw(adj_matrix_vertical, style = "B", zero.policy = TRUE)
horizontal_listw <- mat2listw(adj_matrix_horizontal, style = "B", zero.policy = TRUE)
```

[[677]] Simple feature collection with 3 features and 166 fields Geometry type: POINT We can now rerun our analysis

Table 7: Model 3 Vertical

	Estimate	Std. Error	t-value	Pr(>|t|)
lambda	0.0798	0.0026	30.6366	0
lag(ANY_EVENT_ACLED)	0.3626	0.0051	71.2501	0
SPEI4pg	0.0001	0.0136	0.0090	0.9928
L1_SPEI4pg	0.0161	0.0142	1.1361	0.2559
L2_SPEI4pg	−0.0143	0.0141	−1.0190	0.3082
GSmain_ext_SPEI4pg	−0.0026	0.0144	−0.1787	0.8582
L1_GSmain_ext_SPEI4pg	−0.0434	0.0150	−2.8930	0.0038
L2_GSmain_ext_SPEI4pg	−0.0088	0.0150	−0.5852	0.5584
W_GSmain_ext_SPEI4pg	−0.0036	0.0024	−1.4839	0.1378
W_L1_GSmain_ext_SPEI4pg	0.0054	0.0025	2.1523	0.0314
W_L2_GSmain_ext_SPEI4pg	−0.0024	0.0025	−0.9317	0.3515
W_SPEI4pg	0.0029	0.0021	1.3755	0.1690
W_L1_SPEI4pg	−0.0029	0.0021	−1.3469	0.1780
W_L2_SPEI4pg	0.0047	0.0021	2.2169	0.0266

Table 8: Model 3 Horizontal

	Estimate	Std. Error	t-value	Pr(>|t|)
lambda	0.0691	0.0026	26.3343	0
lag(ANY_EVENT_ACLED)	0.3671	0.0051	71.7848	0
SPEI4pg	−0.0016	0.0137	−0.1169	0.9069
L1_SPEI4pg	0.0140	0.0142	0.9831	0.3256
L2_SPEI4pg	−0.0163	0.0141	−1.1555	0.2479
GSmain_ext_SPEI4pg	−0.0018	0.0144	−0.1273	0.8987
L1_GSmain_ext_SPEI4pg	−0.0445	0.0151	−2.9475	0.0032
L2_GSmain_ext_SPEI4pg	−0.0100	0.0151	−0.6604	0.5090
W_GSmain_ext_SPEI4pg	−0.0040	0.0024	−1.6469	0.0996
W_L1_GSmain_ext_SPEI4pg	0.0055	0.0025	2.1464	0.0318
W_L2_GSmain_ext_SPEI4pg	−0.0025	0.0025	−0.9798	0.3272
W_SPEI4pg	0.0033	0.0021	1.5491	0.1213
W_L1_SPEI4pg	−0.0026	0.0022	−1.1871	0.2352
W_L2_SPEI4pg	0.0051	0.0021	2.4010	0.0164

Table 9: Model 4 Vertical

	Estimate	Std. Error	t-value	Pr(>|t|)
lambda	0.0556	0.0027	20.8863	0
lag(ANY_EVENT_ACLED)	0.3600	0.0052	69.6567	0
SPEI4pg	0.0007	0.0136	0.0483	0.9615
L1_SPEI4pg	0.0207	0.0142	1.4605	0.1442
L2_SPEI4pg	−0.0124	0.0141	−0.8858	0.3757
GSmain_ext_SPEI4pg	−0.0017	0.0141	−0.1203	0.9043
L1_GSmain_ext_SPEI4pg	−0.0491	0.0148	−3.3294	0.0009
L2_GSmain_ext_SPEI4pg	−0.0030	0.0148	−0.2062	0.8367
W_GSmain_ext_SPEI4pg	−0.0040	0.0026	−1.5338	0.1251
W_L1_GSmain_ext_SPEI4pg	0.0061	0.0027	2.2547	0.0242
W_L2_GSmain_ext_SPEI4pg	−0.0023	0.0027	−0.8601	0.3897
W_SPEI4pg	0.0016	0.0022	0.7318	0.4643
W_L1_SPEI4pg	−0.0018	0.0023	−0.7941	0.4272
W_L2_SPEI4pg	0.0051	0.0023	2.2191	0.0265

Table 10: Model 4 Horizontal

	Estimate	Std. Error	t-value	Pr(>|t|)
lambda	0.0447	0.0026	16.9036	0
lag(ANY_EVENT_ACLED)	0.3637	0.0052	70.1381	0
SPEI4pg	−0.0003	0.0137	−0.0245	0.9805
L1_SPEI4pg	0.0192	0.0142	1.3520	0.1764
L2_SPEI4pg	−0.0138	0.0141	−0.9777	0.3282
GSmain_ext_SPEI4pg	−0.0012	0.0142	−0.0840	0.9330
L1_GSmain_ext_SPEI4pg	−0.0500	0.0148	−3.3747	0.0007
L2_GSmain_ext_SPEI4pg	−0.0037	0.0148	−0.2463	0.8055
W_GSmain_ext_SPEI4pg	−0.0044	0.0026	−1.6667	0.0956
W_L1_GSmain_ext_SPEI4pg	0.0060	0.0027	2.2128	0.0269
W_L2_GSmain_ext_SPEI4pg	−0.0025	0.0027	−0.9132	0.3612
W_SPEI4pg	0.0018	0.0022	0.8090	0.4185
W_L1_SPEI4pg	−0.0015	0.0023	−0.6483	0.5168
W_L2_SPEI4pg	0.0055	0.0023	2.3686	0.0179

Summarize and visualize the weights matrixes and briefly explain what they imply

In both the horizontal and the vertical contiguity matrix the average degree is about 1.86 and 1.9. Thus most cells have (as expected) two neighbors but some cells , at the edge, have only one neighbors. This implies that the networks are sparser and less dense than the adjacency matrix used by the authors. This is also reflected in lower 'graph density' for the horizontal and vertical contiguity matrix as compared to the baseline adjacency matrix. Visualizing the spatial weight matrices further corroborates the sparser structure

More generally, the horizontal/vertical contiguity matrix implies that only cells which share the same latitude/longitude and are directly adjacent are considered to be affecting each other directly. It is hard to make a coherent case for why this should be that way in the real world. There is no inherent reason to assume that effects only propagate in one 'direction'.

Still due to the nature of dynamic multipliers in the form of the spatial autoregressive terms, even if the network appears relatively sparse and two cells are not directly linked the cells are still connected. However, the restriction of horizontal and vertical contiguity significantly restricts the dynamic propagation of the effects in this context.

Visualization of Spillover Effects

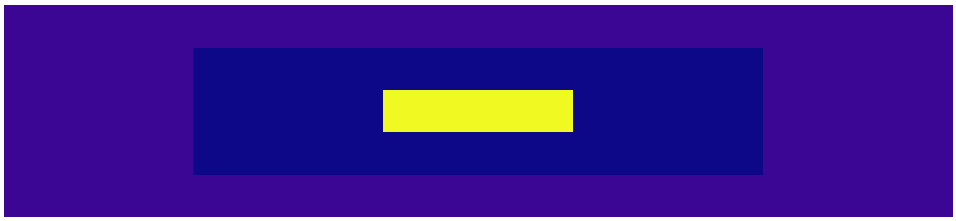
For this exercise we simulate the effect of a one-standard deviation SEPI growing season shock. We multiply the shock times -1 to obtain the negative impact and conversely the increase in conflict incidence. We exogenously shock a cell and calculate 'manually' based on the coefficients of model 4 the magnitude of the shock propagation. We take into account both the dynamic, and spatial autoregressive structure. We see that the shock reaches its peak size at t = 1, consistent with the main message of the paper. A negative one standard deviation growing season shock increases conflict incidence by about 2 percentage points. Overall the spillover effects both in absolute magnitude and visually as reflected in the figure are rather small.

```
# COMPUTATION OF SHOCKS

# first number denotes period second number denotes neighboor shock in period 0
# of standard deviation in growing season SEPI (without contemporanously
# affecting SEPI over the year)
risk_cell_0_0 <- -1 * mod4$coefficients["GSmain_ext_SPEI4pg"] *
sd(geoconflict_main_weights$GSmain_ext_SPEI4pg)

risk_cell_0_1 <- -1 * mod4$coefficients["W_GSmain_ext_SPEI4pg"] *
sd(geoconflict_main_weights$GSmain_ext_SPEI4pg) +
    risk_cell_0_0 * mod4$arcoef

risk_cell_0_2 <- -1 * mod4$coefficients["W_GSmain_ext_SPEI4pg"] *
mod4$coefficients["W_GSmain_ext_SPEI4pg"] *
    sd(geoconflict_main_weights$GSmain_ext_SPEI4pg) + mod4$arcoef * risk_cell_0_1 +
    mod4$arcoef * mod4$arcoef * risk_cell_0_0
```

```
risk_cell_1_0 <- risk_cell_0_0 * mod4$coefficients["lag(ANY_EVENT_ACLED)"] + -1 *
    mod4$coefficients["L1_GSmain_ext_SPEI4pg"] *
sd(geoconflict_main_weights$GSmain_ext_SPEI4pg)

risk_cell_1_1 <- risk_cell_0_1 * mod4$coefficients["lag(ANY_EVENT_ACLED)"] + -1 *
    mod4$coefficients["W_L1_GSmain_ext_SPEI4pg"] *
sd(geoconflict_main_weights$GSmain_ext_SPEI4pg) +
    risk_cell_1_0 * mod4$arcoef

risk_cell_1_2 <- risk_cell_0_2 * mod4$coefficients["lag(ANY_EVENT_ACLED)"] + -1 *
    mod4$coefficients["W_L1_GSmain_ext_SPEI4pg"] *
mod4$coefficients["W_L1_GSmain_ext_SPEI4pg"] *
    sd(geoconflict_main_weights$GSmain_ext_SPEI4pg) + mod4$arcoef * risk_cell_1_1 +
    mod4$arcoef * mod4$arcoef * risk_cell_1_0


risk_cell_2_0 <- risk_cell_1_0 * mod4$coefficients["lag(ANY_EVENT_ACLED)"] + risk_cell_0_0 *
    mod4$coefficients["lag(ANY_EVENT_ACLED)"] * mod4$coefficients["lag(ANY_EVENT_ACLED)"] +
    -1 * mod4$coefficients["L2_GSmain_ext_SPEI4pg"] *
    sd(geoconflict_main_weights$GSmain_ext_SPEI4pg)

risk_cell_2_1 <- risk_cell_1_1 * mod4$coefficients["lag(ANY_EVENT_ACLED)"] + risk_cell_0_1 *
    mod4$coefficients["lag(ANY_EVENT_ACLED)"] * mod4$coefficients["lag(ANY_EVENT_ACLED)"] +
    -1 * mod4$coefficients["W_L2_GSmain_ext_SPEI4pg"] *
    sd(geoconflict_main_weights$GSmain_ext_SPEI4pg) +
    risk_cell_2_0 * mod4$arcoef

risk_cell_2_2 <- -1 * mod4$coefficients["W_L2_GSmain_ext_SPEI4pg"] *
mod4$coefficients["W_L2_GSmain_ext_SPEI4pg"] *
    sd(geoconflict_main_weights$GSmain_ext_SPEI4pg) + mod4$arcoef * risk_cell_2_1 +
    mod4$arcoef * mod4$arcoef * risk_cell_2_0 + risk_cell_2_1 * mod4$arcoef + risk_cell_2_0
*
    mod4$arcoef * mod4$arcoef
```

SEPI Growing Season Shock at t = 0

t = 1



t = 2