Exercise: Exploratory Data Analysis with GeoDa

D G Rossiter Cornell University, Section of Soil & Crop Sciences

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Note: This exercise is an adaptation and extension of one prepared by Dr. Diana Sinton for Cornell course PLSCS/NTRES 6200 in 2016.

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1 Introduction

GeoDa is an open-source program, cross-platform program designed as a simple tool for exploratory spatial data analysis (ESDA) and some spatial modelling of **spatial polygon** data, that is, maps of polygon units such as census tracts or political divisions with a set of **attributes** measured on each one.

GeoDa was first developed at Arizona State University and is now hosted at the University of Chicago¹. The GeoDa program, documentation and sample data is freely available for download from the Geodata Center's GitHub².

GeoDa allows users to experiment with visualization functionality such as linking and brushing across windows. This can be very helpful both for interpretation of and communication about these spatial patterns. It also incorporates several spatial statistical models.

TASK 1: Download and install GeoDa.

Таѕк 2: Start GeoDa.

2 Dataset

We will analyze a small dataset, part of the larger "New York leukemia dataset" developed by Waller and Gotway [2] and adapted by Bivand et al. [1]. This is information on the census tracts in an eight county area including Syracuse (NY) city, relating possible causes to the incidence of leukemia, in particular, exposure to the industrial chemical TCE³.

I have reduced this to just Syracuse city to reduce the size of maps and graphs.

Note: In the USA census tracts have 1 500–8 000 people (optimum size 4 000). They are designed to be socio-economically and demographically fairly homogeneous. Each tract has several block groups; these are made up of 20–40 individual blocks. The tract is usually large enough to compile reliable statistics.

TASK 3: Load the Syracuse leukemia incidence dataset into GeoDa, using the File | New ... menu item or the file open icon. This is a shapefile with base name Syr, so select Syr.shp •

You will see a plain map of the polygons (Fig. 1).

¹ https://spatial.uchicago.edu/geoda

² http://geodacenter.github.io

³ Trichloroethylene, an industrial solvent often found in groundwater



Figure 1: Base map and GeoDa tool bar

TASK 4 : Open the data table by clicking the Table icon. Examine the rows and columns.

We will work with these variables (fields):

These are all reported on the basis of 1980 census tracts. First, the **response** (dependent) variables:

- Cases : the number of leukemia cases 1978–1982; some cases had insufficient georeference, these were added proportionally to tracts, so some "counts" are not integers.
 - Z : log-transformed rate, i.e., normalized by census tract population: $Z_i = \log(1000[\text{Cases} + 1]/n)$, where *n* is the population of the tract.

Second, possible predictors:

- PEXPOSURE : potential exposure, computed as the logarithm of 100 times the inverse of the distance between a census tract centroid and the nearest TCE-producing site;
- PCTAGE65P : percent older than 65 years; this could represent long-term exposure to any environmental factor;
- PCTOWNHOME : percent home ownership; this could indicate lifestyle or economic level.

TASK 5 : Rearrange the Table and the Map so that you can view both. •

The basic GIS "linking" functionality is in place; you can click on polygons in the map and their associated records in the table will highlight, and vice versa. To unselect objects, click anywhere in the white area surrounding the map or at the upper-left of the table. You can select multiple polygons (on the map) or tracts (in the attribute table) with Shift-click for a set or Ctrl-click to add one-by-one. You can also "brush" over the map by holding down the left mouse button, to select in a window,

Q1: Click on the northeaternermost census tract. What is its AREAKEY? What is its population? What percent of its homes are owned rather than rented?



3 Exploratory Data Analysis

3.1 Univariate

TASK 6: Display some themed maps in the Map menu, for one or more of the variables, for example PCTSGE65P.

Compare quantile, percentile, box, and natural breaks maps. Examine how they present the same theme in different ways (Fig. 2).

Q2 : Look at the southeasternmost census tract in these four maps. How do they describe its proportion of older residents, compared to the entire City? Which map(s) best show(s) whether it is unusual?

Q3: Which map is best for assessing spatial autocorrelation of this variable? Why? Does there appear to be autocorrelation? Across how many neighbouring census tracts?

3.2 Bivariate

Now we explore some feature-space plots.

TASK 7: Under the Explore menu, create a Histogram of PCTAGE65P.



Figure 2: Thematic maps

Also create a Scatter Plot of the proportion of residents over the age of 65 PCTAGE65P (Y variable) and the proportion of homes that are owned rather than rented. PCTOWNHOME (X variable)



Figure 3: Histogram and scatterplot

Q4 : *Which tract has the highest proportion of older residents?*

4

Q5 : *Describe the relation between these two attributes.*

TASK 8 : Find an unusual tract (not fitting the overall pattern for the city) and click on its point in the scatterplot.

See Figure 4.



Figure 4: An unusual tract

Because all of the individual graphic elements for each of the 63 polygons are linked, any time that one or more are selected in one window or in one of the exploration plots, their linked highlighted display will activate in all other windows.

Q6: Which tract did you select? Where is it located?

TASK 9: Brush over the southern few tracts by holding down the left mouse button as you define a rectangular window.

Q7: What is the overall relation between home ownership and proportion of over-65 residents? What is this relation for the four southernmost tracts? How do you explain this?

Another interesting plot is the Cartogram.

TASK 10: Make a cartogram (Map | Cartogram) of the proportion over 65 years old PCTAGE65P as the circle *size*, with the disease incidence Z as the circle *colour*.

See Figure 5.

Q8 : *How are the circles placed in the plot? What insight does this give you into the relation between disease incidence and older residents?* •



Figure 5: Cartogram of leukemia incidence vs. older residents

3.3 Multivariate

TASK 11: Open a scatterplot matrix (Explore | Scatter Plot Matrix of the three PCTOWNHOME, PCTAGE65P, and PEXPOSURE, as well as the response variable Z.

See Figure 6.

Q9: Describe the feature-space distributions of the four variables. Looking at the proposed bivariate linear regressions, which have tracts with high leverage, i.e., that greatly influence the line? •

TASK 12 : In the matrix, select the histogram bar for the lowest proportion of home ownership, i.e., where more households rent.

Note how the linked maps highlight these tracts. See Figure 7.

TASK 13: With the scatterplot matrix displayed, select menu option Options | View | Regime Regression. This will then show the separate regression lines and statistics for the overall, selected, and non-selected census tracts.

Q10: Look at the red proposed regression lines – which bivariate correlations are substantially different from the overall correlations if we only consider these tracts?



Figure 6: Scatterplot matrix

TASK 14: Open a Parallel Coordinate Plot (PCP) and Include the three possible predictors PCTOWNHOME, PCTAGE65P, and PEXPOSURE, as well as the response variable Z.

Q11: What is their overall inter-relation?

TASK 15: Click on the line to the highest response.

See Figure <mark>8</mark>.

Q12 : *Which tract is this? How is this response related to the three predictors?* •



Figure 7: Scatterplot matrix with low home ownership tracts selected





4 Neighbors and Distances

For spatial models, we must impose some **spatial structure** on the 63 polygons, that is, how they are related in space. Then we can assess this statistically.

One way is by **distance between polygon centroids**, as in point geostatistics; the spatial weights are based on separation, typically as inverse distance. This considers that distance is the only factor driving any spatial correlation.

However, there are other ways to build a **weights matrix** that relates neighbours; these relate to different hypotheses about how space affects the response. For example, a binary neighbours weighting considers that all first-order neighbours contribute equally to any spatial effect, i.e., it is averaged across the neighbours. With this weighting every tract is influenced equally (on average) by neighbours, and this influence is divded among the neighbours.

TASK 16 : Generate two weights files: (1) Distance Weights and indicate X and Y coordinates, (2) order-1 Contiguity with Queen neighbours (i.e., tracts meeting only at a point are also considered to be neighbours).

To generate a Weights File, choose Tools | Weights Manager | Create. Every shape must have its own unique ID, so check the Add ID Variable and use the existing AREAKEY variable. By default, the new ID variable will be named POLY_ID, or you can choose otherwise. You can then select a type of weighting methods

For the distance weighting, the Threshold distance will automatically be calculated at the minimum distance to ensure that every polygon has at least one neighbor, but you can set any distance you desire. The distance units will be the units associated with the shapefile. For example, if we think that the phenomenon might be spatially-correlated (after accounting for the feature-space regression) to 2.4 km, set the threshold distance to 2400 m.

See Figure 9.

Q13: *What is the maximum number of neighbours considered for any tract in this distance weighting?* •

Note: You could also use the k-Nearest Neighbors option to specific a set number of its closest neighbors that you wish each polygon to use.

When your choices are set, Create the file and name the file with a label indicating the approach used to calculate the neighbors. For example, Queen1.gal would indicate a Queen directionality with 1 order of contiguity; Dist24.gwt would indicate a 2400 m radius inverse-distance weighting.

🏶 Weights Manager — 🗆 🗙	Weights File Creation	X 🦑 Connectivity Histogram - SyrDist24k — 🗆 X
Create Load Remove	Weights File ID Variable AREAKEY Add ID Variable Contiguity Weight	۳)
Weights Name SyrDist SyrDist24k	Queen contiguity Order of contiguity 	
Property Value type threshold symmetry symmetric file SyrDist24K.gwt id variable AREAKEY distance metric Euclidean distance vars centroids distance unit Meter threshold value 2400	Distance Weight Distance metric X-coordinate variable X-coordinate variable X-coordinate variable X-centroids> Ok-Nearest Neighbors Number of neighbors	
Histogram Connectivity Map	Create Close	3 6 9 12 15 18 21 24 27 Number of Neighbors

Figure 9: Creating a distance weighting

As you are deciding which method derives the most valid weights file for your question of interest, you can visually compare the results by using the Connectivity Histogram button in the Weights Manager window, each time after you select your different weights tables. See Figure 10.



Figure 10: Connectivity map and histogram, Queen lag-1 neigbours

Q14: *What is the most common number of neighbours using the Queen lag-1 neighbours?* •

5 Assessing Global Spatial Autocorrelation

Here we evaluate whether the rates of leukemia across the study area may be spatially auto-correlated, without considering any predictors. **TASK 17**: Use Space | Univariate Moran's I, with Z as the variable, to produce a global Moran's I plot. Do this for all the weighting schemes you defined.

See Figure 11.



Figure 11: Global Moran's I

Q15 : Do the weighting schemes all give the same value of global Moran's I? If not, which implies stronger spatial correlation? Why? •

TASK 18: Open two themed maps: decile (10-quantile) of Z (incidence) and PEXPOSURE (exposure).

In the Moran's I scatterplot, click on the highest positive Z (incidence) and highest weighted lag Z.

See Figure 12.

Q16: *Where is this tract located? Does it also have a high exposure? Do its neighbours have high incidences? Do they have high exposures?* •

6 Assessing Local Spatial Autocorrelation

TASK 19 : Examine where in the map are the hotspots of local autocorrelation.

First, make sure that your desired Weights file is set as the default, i.e., highlighted in the Weights Manager window.

TASK 20: Use Space | Univariate Local Moran's I, with Z as the variable, to produce a local Moran's I plot. Do this for all the weighting schemes you defined. Generate two output windows: the Significance Map and the Cluster Map.

The Significance map shows where there are leukemia values that are statistically significantly higher **or** lower than the neighboring values would



Figure 12: Global Moran's I

have predicted. With the Cluster Map, you can see where the higherthan-expected and lower-then-expected values vary.

See Figure 13.



Figure 13: Local Moran's I significance and clusters

Q17: Which areas of the city are clusters of high leukemia incidence? Are there any tracts that have high incidence, but are surrounded by tracts with low incidence?

Another way to find hot spots is with Geary's G or G*; if you want you can experiment with these.

7 Spatial Regression

Here we try to find the covariates ("predictors") correlated (which maybe cause) leukemia. Of course, we can do this non-spatially, i.e., all in attribute space, not taking spatial relations into account.

TASK 21: Compute a multivariate linear regression model of leukemia incidence (response) as predicted by the three possible causitive factors (predictors). This is with the Regression menu item. Select Z as the dependent variable, and PCTOWNHOME, PCTAGE65P, and PEXPOSURE as the covariates. This is the Classic linear model, i.e., not taking spatial correlation into account.

Non-spatial linear model: $\mathbf{Y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}$

Figure 14 shows how to specify the regression; Figure 15 shows the results.

	Regression	n	
Variables		Dependent Variable	
х	>	Z	
Y		Covariates	
POP8		PEXPOSURE	
TRACTCAS		PCTOWNHOME	
PROPCAS	>	PCTAGE65P	
AVGIDIST	<		
Cases	>>		
Xm	<<		
Ym			
Xshift			
Yshift			
/eights File Syr24 Models			0 14
Classic	O Spatial Lag	g OSpatial Error	
🔽 Pred. Val. and F	les. 🔽 Coeff.	Var. Mat. 🛛 White Test	

Figure 14: Specifying a "classic" linear regression

Q18 : What is the adjusted R² of this model? What are the signs of the slopes for each predictor? What is the interpretation? Which (if any) predictors are significantly different from zero?

The model summary shows many problems with the linear model:

		Re	gression Report		
2					
>>03/07/2019	17:18:29				
REGRESSION					
		DV TENCE C	QUARES ESTIMAT	101	
	: Sy		QUARES ESTIMAT	ION	
Dependent Var	iable :	z ·	Number of Obse	rvations: 63	
Mean depender	t var : 0	.0377522	Number of Vari	ables : 4	
S.D. depender	t var :	0.996518	Degrees of Fre	edom : 59	
R-squared	:	0.185475	F-statistic	: 4	.47829
Adjusted R-so	uared :	0.144059	Prob(F-statist	ic) : 0.00	671609
Sum squared r	esidual:	50.9583	Log likelihood	: -8	2.7112
Sigma-square	:	0.863701	Akaike info cr	: 4 ic) : 0.00 : -8 iterion : 1 ion : 1	73.422
S.E. of regre	ession :	0.929355	scnwarz criter	10n : 1	81.995
Sigma-square S.E of regres	ru :	0.800369			
5.5 Of regres	STOIL HE:	0.077300			
				t-Statistic	
CON	ISTANT -	3.15559	2.16024	-1.46076 1.24206 -0.649259	0.14939
PEXE	OSURE	2.64063	2.12602	1.24206	0.21913
PCTOW	NHOME -0	.307937	0.47429	-0.649259	0.51869
PCTA	GE65P	4.24105	1.22995	3.44815	0.00105
REGRESSION DI	ACNOSTICS				
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Jarque-Bera					
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DIAGNOSTICS F RANDOM COEFFI TEST	CIENTS		VALUE	PROB	
DIAGNOSTICS F RANDOM COEFFI TEST	CIENTS		VALUE 15.6910	0.00131	
DIAGNOSTICS F RANDOM COEFFI TEST Breusch-Pagar Koenker-Basse	CIENTS DF a test 3 ett test 3		VALUE		
DIAGNOSTICS F RANDOM COEFFI TEST	CIENTS DF a test 3 ett test 3		VALUE 15.6910 5.2255	0.00131	
DIAGNOSTICS F RANDOM COEFFI TEST Breusch-Pagar Koenker-Basse SPECIFICATION	CIENTS DF a test 3 ett test 3 N ROBUST TEST		VALUE 15.6910 5.2255 VALUE	0.00131 0.15601	
DIAGNOSTICS F RANDOM COEFFI TEST Breusch-Pagar Koenker-Basse SPECIFICATION TEST	CCIENTS DF a test 3 ett test 3 I ROBUST TEST DF		VALUE 15.6910 5.2255	0.00131 0.15601 PROB	
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A CONSTRUCT OF A CONS	CIENTS DF a test 3 att test 3 ROBUST TEST DF 9 VARIANCE MAT PEXPOSURE -4.556146	RIX PCTOWNHOME 0.015900	VALUE 15.6910 5.2255 VALUE 13.2268 PCTAGE65P -0.238955	0.00131 0.15601 PROB	
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DAGNOSTICS F RANDOM COEFFI TEST Breusch-Pagar Koenker-Basse SPECIFICATION TEST White CONSTANT 4.666623 -4.556146 0.015900	CIENTS DF a test 3 att test 3 ROBUST TEST DF 9 VARIANCE MAT PEXPOSURE -4.556146	RIX PCTOWNHOME 0.015900 -0.092970 0.224951	VALUE 15.6910 5.2255 VALUE 13.2268 PCTAGE65P -0.238955 0.028862 -0.039303	0.00131 0.15601 PROB	

Figure 15: Multiple linear regression results

- 1. The **multicolilinearity** (or multiple) **condition number** represents the sensitivity of the model to small changes in the design matrix, i.e., the values of the covariables. A high value (often > 30) indicates high colinearity in one or more predictors; here we see that is the case.
- 2. The Jarque-Bera test is whether the residuals have the skewness and kurtosis matching a normal distribution. Here we see a high value, quite unlikely to be normal.

However we will not fix up this model, we proceed to compare it to models which do take into account spatial correlation.

7.1 Spatial Error model

The first model with a spatial component we will consider is the **spatial error model**. This allows **resduals** of the linear model to be spatially-correlated, and quantifies to what extent they are included in the model.

This typically occurs when there is some spatially-correlated covariate that (1) affects the response and (2) we do not know, or maybe even

suspect – otherwise we would identify it, measure it, and include in the linear model. However, we may suspect a factor that we have not, or can not, measure, and this factor has spatial correlation.

For example, this database does not report the proportion of different ethnic groups, nor of different occupational groups (factory workers, office workers, service workers ...). These may be (1) related to leukemia (genetic susceptibility, occupational exposure), (2) spatially-correlated. If such factors influence leukemia, they will be represented in the residuals, and thus the spatial error model will be provably better than the feature space-only model.

The spatial error model is:

- formula: $\mathbf{Y} = \mathbf{X}\boldsymbol{\beta} + \lambda \mathbf{W}\mathbf{u} + \boldsymbol{\varepsilon}$
- W is a matrix representing the spatial structure (e.g., neighbour weights)
- $\mathbf{u} = (\mathbf{Y} \mathbf{X}\boldsymbol{\beta})$ are the spatially-correlated **residuals**
- λ is the strength of autoregression of the errors
- *ε* is the independent error (not autoregressive)

TASK 22: Compute a multivariate linear regression model of leukemia incidence (response) as predicted by the three possible causitive factors (predictors). This time (1) select a Weights file (one you created above), and then you can specify the SAR **Spatial Error** linear model. This takes spatial correlation of the **linear model residuals** into account, considering the values of the model **residual** in each tract's neighbourhood, as defined by the weights.

Figure 16 shows the results.

Q19: What is the adjusted R² of this model? Is it higher or lower than that for the feature-space only model? Is this expected? How can it be explained?

Q20: What are the signs of the slopes for each predictor? What is the interpretation? Which (if any) predictors are significantly different from zero? What changes in this model compared to the feature-space multiple regression? I.e., which predictors become more or less important and/or significant?

Q21 : What is the strength of the autocorrelation parameter λ ?

The **likelihood ratio test** gives the probability that the SAR spatial error model is *not* better than the feature-space-only multiple regression.

		Re	gression Repor	t	
2					
>>03/07/2019	18:28:29				
REGRESSION					
			DEL - MAXIMUM	LIKELIHOOD P	STIMATION
Data set Spatial Weigl	: S	yr m24			
Dependent Way	riable : D	7 7	Number of Obs	orvations	63
Mean depender	t var :	0.037752	Number of Var	iables :	4
S.D. depender	nt var :	0.996518	Number of Obs Number of Var Degrees of Fr	eedom :	59
Lag coeff. (1	Lambda) :	0.466941	,		
R-squared	:	0.241243	R-squared (BU Log likelihoo	SE) :-	
Sq. Correlat:	ion : -		Log likelihoo Akaike info o	d : -	-81.094551
Sigma-square		0.753483	Akaike info c	riterion :	170.189
5.E of regres	ssion :	0.868034	Schwarz crite	rion :	178.762
		pefficient		z-value	e Probability
					0.25983
PEXI	OSURE	2.87842	3.04238	0.94610	58 0.25983 07 0.34409 17 0.97769
PCTO	VNHOME -	0.0135155	0.483305	-0.027964	0.97769
					15 0.00058
	LAMBDA		0.221434		0.03497
REGRESSION D	ACNOSTICS				
DIAGNOSTICS 1		EDASTICITY			
RANDOM COEFF		abhbiiciii			
TEST			DF	VALUE	PROB
Breusch-Paga	n test			11.6116	0.00884
DIAGNOSTICS 1			-	~ /	
SPATIAL ERRON	C DEPENDENCI	S FOR WEIGHT	MATRIX : Syr		PROB
rest Likelihood Ra	tio Test		DF 1	3.2332	
Treiinood K	itto iest		1	5.2332	0.0/210
COEFFICIENTS	VARIANCE M	ATRIX	D00000000000	TAMORE	
CONSTANT	PEXPOSURE	PCTOWNHOME	PCTAGE65P	LAMBDA	
9.303402	-9.354/8/	0.029032	-0.28/450	0.000000	
0.029032	-0.119136	0 233584	PCTAGE65P -0.287450 0.079041 -0.018171	0.000000	
-0.287450	0.079041	-0.018171	1.402760	0.000000	
			0.000000		
0.000000					

Figure 16: SAR spatial error model regression results

Q22 : What is the probability that the SAR spatial error model is not better than the feature-space-only multiple regression? What does this imply about the possible causes of leukemia?

7.2 Spatial Lag model

Another possible effect of spatial autocorrelation is in the response, that is, the values of the response in a tract's neighbours directly influence the response in the tract, after taking into account the feature-space prediction. This measures "contagion", which seems unlikely for human leukemia⁴, however we still evaluate this.

TASK 23: Compute a multivariate linear regression model of leukemia incidence (response) as predicted by the three possible causitive factors (predictors). This time (1) select a Weights file (one you created above), and then you can specify the SAR **Spatial Lag** linear model. This takes

⁴ although quite likely for feline leukemia, if infected cats travel across tract boundaries

spatial correlation into account, considering the values of the **response** variable in each tract's neighbourhood, as defined by the weights.

The spatial lag model is: $\mathbf{Y} = \rho \mathbf{W} \mathbf{Y} + \mathbf{X} \boldsymbol{\beta} + \boldsymbol{\varepsilon}$, where ρ is the strength of autoregression of the response; this multiplies the weights matrix times the response **WY** on the right-hand (predictor) side of the equation.

Figure 17 shows the results.

	Re	egression Report	t	
•				
>>03/07/2019 18:27:5 REGRESSION	3			
SUMMARY OF OUTPUT: S Data set		EL - MAXIMUM L	IKELIHOOD ESTIN	MATION
matial Maight	. Craw24			
Dependent Variable	· 59124	Number of Obs	ervations: 6	3
lean dependent var	: 0.03//522	Numper of var.	lapies : :	5
J.D. dependent var	: 0.996518	Degrees of Fr	eedom : 5	3
S.D. dependent var Lag coeff. (Rho)	: 0.435129	-		
R-squared	: 0.240639	Log likelihoo	d : .	-81.0283
Sq. Correlation	: -	Akaike info c	riterion :	172.057
Sigma-square S.E of regression	: 0.754082	Schwarz crite	rion :	182.772
.E of regression	: 0.868379			
	Coofficient	Ctd Enviro	1	Duchahilit
variabie	Coefficient			
WZ	0.435129	0.213419	2.03885	0.04147
CONSTANT	-2.38973	2.07463		0.24937
PEXPOSURE	1.86442	2.0381		0.36031
PCTOWNHOME		0.443173		
PCTAGE65P	3.99453	1.15859	3.44//5	0.00057
REGRESSION DIAGNOSTI				
DIAGNOSTICS FOR HETE				
RANDOM COEFFICIENTS				
TEST			VALUE	
Breusch-Pagan test		3	12.0911	0.00708
DIAGNOSTICS FOR SPAT	IAL DEPENDENCE			
SPATIAL LAG DEPENDEN				
TEST				PROB
Likelihood Ratio Tes	st	1	3.3656	0.06657
COEFFICIENTS VARIANC	E MATRIX			
CONSTANT PEXPOS	SURE PCTOWNHOM	E PCTAGE65P	WZ	
	0.01442	6 -0.278994	0.102293	
4.304079 -4.196			-0.097233	
4.304079 -4.196 -4.196260 4.153	-0.08168	8 0.092085		
4.304079 -4.196 -4.196260 4.153 0.014426 -0.081	856 -0.08168 688 0.19640	2 -0.034482	0.000243	
4.304079 -4.196 -4.196260 4.153 0.014426 -0.081	856 -0.08168 688 0.19640	2 -0.034482	0.000243	
4.304079 -4.196 -4.196260 4.153	856 -0.08168 688 0.19640	2 -0.034482	0.000243	

Figure 17: SAR spatial lag model regression results

Q23 : Is the lag coefficient ρ significant in the regression? What is the probability that the SAR spatial lag model is not better than the feature-space-only multiple regression? What does this imply about the possible causes of leukemia?

8 Finishing with GeoDa

GeoDa provides several opportunities to save images or export newly derived data in tabular form. If while using GeoDa you have derived values and added variables to the attribute table of your shapefile, you will be prompted to Save these as you Exit. Otherwise, the program can simply be closed.

References

- Roger S. Bivand, Edzer J. Pebesma, and V. Gómez-Rubio. Applied Spatial Data Analysis with R. Springer, 2nd edition, 2013. ISBN 978-1-4614-7617-7; 978-1-4614-7618-4 (e-book). URL http://www. asdar-book.org/.
- [2] L. A. Waller and C. A. Gotway. *Applied spatial statistics for public health data*. Wiley-Interscience, Hoboken, N.J., 2004.