

Moran Eigenvectors

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1 Introduction

The Moran eigenvector approach (Dray et al., 2006; Griffith and Peres-Neto, 2006) involved the spatial patterns represented by maps of eigenvectors; by choosing suitable orthogonal patterns and adding them to a linear or generalised linear model, the spatial dependence present in the residuals can be moved into the model.

It uses brute force to search the set of eigenvectors of the matrix \mathbf{MWM} , where

$$\mathbf{M} = \mathbf{I} - \mathbf{X}(\mathbf{X}^T\mathbf{X})^{-1}\mathbf{X}^T$$

is a symmetric and idempotent projection matrix and \mathbf{W} are the spatial weights. In the spatial lag form of `SpatialFiltering` and in the GLM ME form below, \mathbf{X} is an n -vector of ones, that is the intercept only.

In its general form, `SpatialFiltering` chooses the subset of the n eigenvectors that reduce the residual spatial autocorrelation in the error of the model with covariates. The lag form adds the covariates in assessment of which eigenvectors to choose, but does not use them in constructing the eigenvectors. `SpatialFiltering` was implemented and contributed by Yongwan Chun and Michael Tiefelsdorf, and is presented in Tiefelsdorf and Griffith (2007); ME is based on Matlab code by Pedro Peres-Neto and is discussed in Dray et al. (2006) and Griffith and Peres-Neto (2006).

```
> library(maptools)
> owd <- getwd()
> setwd(system.file("etc/shapes", package = "spdep"))
> NY8 <- readShapeSpatial("NY8_utm18")
> library(spdep)
> setwd(system.file("etc/weights", package = "spdep"))
> NY_nb <- read.gal("NY_nb.gal", region.id = row.names(NY8))
> setwd(owd)

> nySFE <- SpatialFiltering(Z ~ PEXPOSURE + PCTAGE65P + PCTOWNHOME, data = NY8,
+   nb = NY_nb, style = "W", verbose = FALSE)
> nylmSFE <- lm(Z ~ PEXPOSURE + PCTAGE65P + PCTOWNHOME + fitted(nySFE),
+   data = NY8)
> summary(nylmSFE)
```

Call:

```
lm(formula = Z ~ PEXPOSURE + PCTAGE65P + PCTOWNHOME + fitted(nySFE),
    data = NY8)
```

Residuals:

Min	1Q	Median	3Q	Max
-1.5184	-0.3523	-0.0105	0.3221	3.1964

Coefficients:

```

              Estimate Std. Error t value Pr(>|t|)
(Intercept)   -0.51728    0.14606   -3.542 0.000469 ***
PEXPOSURE       0.04884    0.03230    1.512 0.131717
PCTAGE65P      3.95089    0.55776    7.083 1.25e-11 ***
PCTOWNHOME     -0.56004    0.15688   -3.570 0.000423 ***
fitted(nySFE)vec13 -2.09397  0.60534   -3.459 0.000630 ***
fitted(nySFE)vec44 -2.24003  0.60534   -3.700 0.000261 ***
fitted(nySFE)vec6  1.02979  0.60534    1.701 0.090072 .
fitted(nySFE)vec38 1.29282  0.60534    2.136 0.033613 *
fitted(nySFE)vec20 1.10064  0.60534    1.818 0.070150 .
fitted(nySFE)vec14 -1.05105  0.60534   -1.736 0.083662 .
fitted(nySFE)vec75  1.90600  0.60534    3.149 0.001826 **
fitted(nySFE)vec21 -1.06331  0.60534   -1.757 0.080138 .
fitted(nySFE)vec36 -1.17861  0.60534   -1.947 0.052578 .
fitted(nySFE)vec61 -1.08582  0.60534   -1.794 0.073986 .
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

Residual standard error: 0.6053 on 267 degrees of freedom
Multiple R-squared:  0.3401,    Adjusted R-squared:  0.308
F-statistic: 10.58 on 13 and 267 DF,  p-value: < 2.2e-16

> nylm <- lm(Z ~ PEXPOSURE + PCTAGE65P + PCTOWNHOME, data = NY8)
> anova(nylm, nylmSFE)

```

Analysis of Variance Table

```

Model 1: Z ~ PEXPOSURE + PCTAGE65P + PCTOWNHOME
Model 2: Z ~ PEXPOSURE + PCTAGE65P + PCTOWNHOME + fitted(nySFE)
      Res.Df    RSS Df Sum of Sq    F    Pr(>F)
1       277 119.619
2       267  97.837 10    21.782 5.9444 3.988e-08 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

Since the SpatialFiltering approach does not allow weights to be used, we see that the residual autocorrelation of the original linear model is absorbed, or ‘whitened’ by the inclusion of selected eigenvectors in the model, but that the covariate coefficients change little. The addition of these eigenvectors – each representing an independent spatial pattern – relieves the residual autocorrelation, but otherwise makes few changes in the substantive coefficient values.

The ME function also searches for eigenvectors from the spatial lag variant of the underlying model, but in a GLM framework. The criterion is a permutation bootstrap test on Moran’s I for regression residuals, and in this case, because of the very limited remaining spatial autocorrelation, is set at $\alpha = 0.5$. Even with this very generous stopping rule, only two eigenvectors are chosen; their combined contribution just improves only the fit of the GLM model.

```

> NYlistwW <- nb2listw(NY_nb, style = "W")
> set.seed(111)
> nyME <- ME(Cases ~ PEXPOSURE + PCTAGE65P + PCTOWNHOME, data = NY8, offset = log(POP8),
+   family = "poisson", listw = NYlistwW, alpha = 0.5)
> nyME

Eigenvector ZI pr(ZI)
0          NA NA   0.35
1          24 NA   0.40
2         223 NA   0.45
3         206 NA   0.56

> NY8$eigen_24 <- fitted(nyME)[, 1]
> NY8$eigen_223 <- fitted(nyME)[, 2]

```

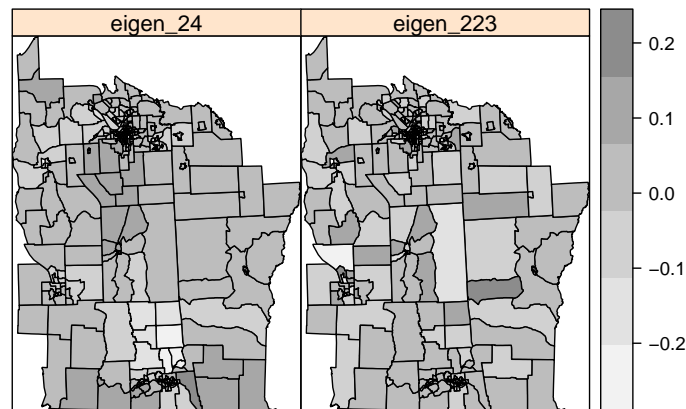


Figure 1: Maps of the two eigenvalues selected for inclusion in the Poisson regression model

```
> nyglmME <- glm(Cases ~ PEXPOSURE + PCTAGE65P + PCTOWNHOME + offset(log(POP8)) +
+   fitted(nyME), data = NY8, family = "poisson")
> summary(nyglmME)
```

Call:

```
glm(formula = Cases ~ PEXPOSURE + PCTAGE65P + PCTOWNHOME + offset(log(POP8)) +
    fitted(nyME), family = "poisson", data = NY8)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-2.5697	-1.0681	-0.2121	0.6094	3.1629

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	-8.12711	0.18362	-44.261	< 2e-16 ***
PEXPOSURE	0.14236	0.03143	4.530	5.90e-06 ***
PCTAGE65P	4.11021	0.59970	6.854	7.19e-12 ***
PCTOWNHOME	-0.38249	0.19265	-1.985	0.0471 *
fitted(nyME)vec24	1.52690	0.72280	2.112	0.0346 *
fitted(nyME)vec223	0.81347	0.70114	1.160	0.2460
fitted(nyME)vec206	0.01330	0.69180	0.019	0.9847

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for poisson family taken to be 1)

Null deviance: 428.25 on 280 degrees of freedom
 Residual deviance: 347.34 on 274 degrees of freedom
 AIC: Inf

Number of Fisher Scoring iterations: 5

```
> nyGLMp <- glm(Cases ~ PEXPOSURE + PCTAGE65P + PCTOWNHOME + offset(log(POP8)),
+   data = NY8, family = "poisson")
> anova(nyGLMp, nyglmME, test = "Chisq")
```

Analysis of Deviance Table

Model 1: Cases ~ PEXPOSURE + PCTAGE65P + PCTOWNHOME + offset(log(POP8))

Model 2: Cases ~ PEXPOSURE + PCTAGE65P + PCTOWNHOME + offset(log(POP8)) +

```
fitted(nyME)
  Resid. Df Resid. Dev Df Deviance Pr(>Chi)
1      277      353.35
2      274      347.34  3    6.011    0.1111
```

Figure 1 shows the spatial patterns chosen to match the very small amount of spatial autocorrelation remaining in the model. As with the other Poisson regressions, the closeness to TCE sites is highly significant. Since, however, many TCE sites are also in or close to more densely populated urban areas with the possible presence of both point-source and non-point-source pollution, it would be premature to take such results simply at their face value. There is, however, a potentially useful contrast between the cities of Binghamton in the south of the study area with several sites in its vicinity, and Syracuse in the north without TCE sites in this data set.

References

- Dray, S., Legendre, P., and Peres-Neto, P. R. (2006). Spatial modeling: A comprehensive framework for principle coordinate analysis of neighbor matrices (PCNM). *Ecological Modelling*, 196:483–493.
- Griffith, D. A. and Peres-Neto, P. R. (2006). Spatial modeling in ecology: The flexibility of eigenfunction spatial analyses. *Ecology*, 87:2603–2613.
- Tiefelsdorf, M. and Griffith, D. A. (2007). Semiparametric filtering of spatial autocorrelation: The eigenvector approach. *Environment and Planning A*, 39:1193–1221.