

## Example 10-14

September 11, 2020

```
[ ]: # install the following packages and libraries
```

```
install.packages("pder")
install.packages("plm")
install.packages("splm")
```

```
library(spData)
install.packages("spDataLarge")
library("pder")
library("plm")
library("splm")
library("spdep")
library("lmtest")
```

```
[2]: ##-----Block 1-----
```

```
#### Example 10-14
```

```
## -----
```

```
data("EvapoTransp", package = "pder")
data("etw", package = "pder")
```

```
# panel model with spatial and serial correlation
```

```
evapo <- et ~ prec + meansmd + potet + infil + biomass + plantcover +  
  softforbs + tallgrass + diversity + matgram + dwarfshrubs + legumes
```

```
semsr.evapo <- sprem1(evapo, data=EvapoTransp, w=etw,  
  lag=FALSE, errors="semsr")
```

```
summary(semsr.evapo)
```

ML panel with , AR(1) serial correlation, spatial error correlation

Call:

```
sprem1(formula = evapo, data = EvapoTransp, w = etw, lag = FALSE,  
  errors = "semsr")
```

Residuals:

	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
	-2.2598	-0.5000	0.0209	-0.0469	0.4198	2.3730

Error variance parameters:

	Estimate	Std. Error	t-value	Pr(> t )
psi	0.166538	0.048267	3.4503	0.0005599 ***
rho	0.866527	0.024556	35.2877	< 2.2e-16 ***

Coefficients:

	Estimate	Std. Error	t-value	Pr(> t )
(Intercept)	0.86604110	0.56232635	1.5401	0.123535
prec	-0.12963576	0.15433822	-0.8399	0.400939
meansmd	0.01896769	0.00445226	4.2602	2.042e-05 ***
potet	0.55114397	0.33582766	1.6412	0.100766
infil	0.02351347	0.02187619	1.0748	0.282445
biomass	0.00233540	0.00030511	7.6542	1.945e-14 ***
plantcover	0.01917420	0.11033208	0.1738	0.862033
softforbs	0.13235937	0.04146285	3.1922	0.001412 **
tallgrass	0.17454017	0.05409913	3.2263	0.001254 **
diversity	0.04077520	0.03578975	1.1393	0.254579
matgram	-0.02981440	0.03304014	-0.9024	0.366861
dwarfshrubs	0.09840506	0.05412693	1.8180	0.069058 .
legumes	-0.01630359	0.00559113	-2.9160	0.003546 **

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

```
[3]: ##-----Block 2-----
# t test
coeftest(plm(evapo, EvapoTransp, model="pooling"))
```

t test of coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	1.07699252	0.16858645	6.3884	4.485e-10 ***
prec	-0.20795429	0.03756344	-5.5361	5.480e-08 ***
meansmd	0.02274891	0.00648884	3.5058	0.0005045 ***
potet	0.76794143	0.08837988	8.6891	< 2.2e-16 ***
infil	0.05564773	0.03103026	1.7933	0.0736434 .
biomass	0.00010371	0.00038920	0.2665	0.7900058
plantcover	0.04465691	0.15780076	0.2830	0.7773207
softforbs	0.10430509	0.05829972	1.7891	0.0743214 .
tallgrass	0.17301265	0.07866758	2.1993	0.0284057 *
diversity	0.01621414	0.05133289	0.3159	0.7522648
matgram	-0.06953728	0.04900063	-1.4191	0.1566140
dwarfshrubs	0.07145072	0.07713510	0.9263	0.3548227
legumes	-0.01944683	0.00811509	-2.3964	0.0169976 *

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

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[4]: ##-----Block 3-----
# t test omitting the spatial error term
coefest(spreml(evapo, EvapoTransp, w=etw, errors="sem"))
```

z test of coefficients:

	Estimate	Std. Error	z value	Pr(> z )	
(Intercept)	1.0625398	0.5660491	1.8771	0.0605022	.
prec	-0.1556424	0.1533498	-1.0150	0.3101294	
meansmd	0.0179289	0.0039623	4.5249	6.043e-06	***
potet	0.5325736	0.3274930	1.6262	0.1039042	
infil	0.0221105	0.0193959	1.1400	0.2543036	
biomass	0.0023116	0.0002860	8.0824	6.348e-16	***
plantcover	0.0163067	0.0972885	0.1676	0.8668886	
softforbs	0.1319485	0.0365432	3.6108	0.0003053	***
tallgrass	0.1766059	0.0476921	3.7030	0.0002130	***
diversity	0.0383890	0.0315495	1.2168	0.2236855	
matgram	-0.0310060	0.0291469	-1.0638	0.2874258	
dwarfshrubs	0.1044045	0.0478209	2.1832	0.0290180	*
legumes	-0.0166536	0.0049294	-3.3784	0.0007290	***

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