

# Package ‘scgwr’

May 9, 2026

**Type** Package

**Title** Scalable Geographically Weighted Regression

**Version** 0.1.2-21

**Date** 2021-11-11

**Author** Daisuke Murakami[cre,aut], Narumasa Tsutsumida[ctb], Takahiro Yoshida[ctb], Tomoki Nakaya[ctb], Lu Binbin[ctb]

**Maintainer** Daisuke Murakami <dmuraka@ism.ac.jp>

**Description** Fast and regularized version of GWR for large dataset, detailed in Murakami, Tsutsumida, Yoshida, Nakaya, and Lu (2019) <[doi:10.48550/arXiv.1905.00266](https://doi.org/10.48550/arXiv.1905.00266)>.

**License** GPL (>= 2)

**Encoding** UTF-8

**Imports** FNN, spData, sp, dplyr, parallel, optimParallel

**NeedsCompilation** no

**Repository** CRAN

**Date/Publication** 2021-11-11 07:40:02 UTC

## Contents

predict0	1
scgwr	3
scgwr_p	4

<b>Index</b>	<b>7</b>
--------------	----------

---

predict0	<i>Spatial prediction using the scalable GWR model</i>
----------	--

---

## Description

This function predicts explained variables and spatially varying coefficients at unobserved sites using the scalable GWR model.

**Usage**

```
predict0( mod, coords0, x0 = NULL )
```

**Arguments**

mod	Output from the scgwr function
coords0	Matrix of spatial point coordinates at predicted sites (N0 x 2)
x0	Matrix of explanatory variables at predicted sites (N0 x K). If NULL, explained variables are not predicted (only spatially varying coefficients are predicted). Default is NULL

**Value**

pred	Vector of predicted values (N0 x 1)
b	Matrix of estimated coefficients (N0 x K)
bse	Matrix of the standard errors for the coefficients (N0 x k)
t	Matrix of the t-values for the coefficients (N0 x K)
p	Matrix of the p-values for the coefficients (N0 x K)

**Examples**

```
require(spData)
data(boston)

id_obs <- sample(dim(boston.c)[1],400)

##### data at observed sites
y <- log(boston.c[id_obs,"MEDV"])
x <- boston.c[id_obs, c("CRIM", "INDUS","ZN","NOX","AGE")]
coords <- boston.c[id_obs , c("LON", "LAT") ]

##### data at predicted sites
x0 <- boston.c[-id_obs, c("CRIM", "INDUS","ZN","NOX", "AGE")]
coords0 <- boston.c[-id_obs , c("LON", "LAT") ]

mod <- scgwr( coords = coords, y = y, x = x )
pred0 <- predict0( mod=mod, coords0=coords0, x0=x0)

pred <- pred0$pred # predicted value
b <- pred0$b # spatially varying coefficients
b[1:5,]

bse <- pred0$bse # standard error of the coefficients
bt <- pred0$t # t-values
bp <- pred0$p # p-values
```

**Description**

This function estimates a scalable geographically weighted regression (GWR) model. See [scgwr\\_p](#) for parallel implementation of the model for very large samples.

**Usage**

```
scgwr( coords, y, x = NULL, knn = 100, kernel = "gau",
       p = 4, approach = "CV", nsamp = NULL)
```

**Arguments**

coords	Matrix of spatial point coordinates (N x 2)
y	Vector of explained variables (N x 1)
x	Matrix of explanatory variables (N x K). Default is NULL
knn	Number of nearest-neighbors being geographically weighted. Default is 100. Larger knn is better for larger samples (see Murakami et al., 2019)
kernel	Kernel to model spatial heterogeneity. Gaussian kernel ("gau") and exponential kernel ("exp") are available
p	Degree of the polynomial to approximate the kernel function. Default is 4
approach	If "CV", leave-one-out cross-validation is used for the model calibration. If "AICc", the corrected Akaike Information Criterion is minimized for the calibration. Default is "CV"
nsamp	Number of samples used to approximate the cross-validation. The samples are randomly selected. If the value is large enough (e.g., 10,000), error due to the random sampling is quite small owing to the central limit theorem. The value must be smaller than the sample size. Default is NULL

**Value**

b	Matrix of estimated coefficients (N x K)
bse	Matrix of the standard errors for the coefficients (N x k)
t	Matrix of the t-values for the coefficients (N x K)
p	Matrix of the p-values for the coefficients (N x K)
par	Estimated model parameters including a scale parameter and a shrinkage parameter if <code>penalty = TRUE</code> (see Murakami et al., 2018)
e	Error statistics. It includes sum of squared errors (SSE), residual standard error (resid_SE), R-squared (R2), adjusted R2 (adjR2), log-likelihood (logLik), corrected Akaike information criterion (AICc), and the cross-validation (CV) score measured by root mean squared error (RMSE) (CV_score(RMSE))

pred	Vector of predicted values (N x 1)
resid	Vector of residuals (N x 1)
other	Other objects internally used

## References

Murakami, D., Tsutsumida, N., Yoshida, T., Nakaya, T., and Lu, B. (2019) Scalable GWR: A linear-time algorithm for large-scale geographically weighted regression with polynomial kernels. <arXiv:1905.00266>.

## See Also

[scgwr\\_p](#), [predict0](#)

## Examples

```
require( spData )
data( boston )
coords <- boston.c[, c("LON", "LAT") ]
y <- log(boston.c[, "MEDV"])
x <- boston.c[, c("CRIM", "ZN", "INDUS", "CHAS", "AGE")]
res <- scgwr( coords = coords, y = y, x )
res
```

---

scgwr_p	<i>Parallel implementation of scalable geographically weighted regression</i>
---------	---

---

## Description

Parallel implementation of scalable geographically weighted regression for large samples

## Usage

```
scgwr_p( coords, y, x = NULL, knn = 100, kernel = "gau",
         p = 4, approach = "CV", nsamp = NULL, cl = NULL)
```

## Arguments

coords	Matrix of spatial point coordinates (N x 2)
y	Vector of explained variables (N x 1)
x	Matrix of explanatory variables (N x K). Default is NULL
knn	Number of nearest-neighbors being geographically weighted. Default is 100. Larger knn is better for larger samples (see Murakami et al., 2019)
kernel	Kernel to model spatial heterogeneity. Gaussian kernel ("gau") and exponential kernel ("exp") are available

p	Degree of the polynomial to approximate the kernel function. Default is 4
approach	If "CV", leave-one-out cross-validation is used for the model calibration. If "AICc", the corrected Akaike Information Criterion is minimized for the calibration. Default is "CV"
nsamp	Number of samples used to approximate the cross-validation. The samples are randomly selected. If the value is large enough (e.g., 10,000), error due to the sampling is quite small owing to the central limit theorem. The value must be smaller than the sample size. Default is NULL
cl	Number of cores used for the parallel computation. If cl = NULL, which is the default, the number of available cores is detected and used

### Value

b	Matrix of estimated coefficients (N x K)
bse	Matrix of the standard errors for the coefficients (N x k)
t	Matrix of the t-values for the coefficients (N x K)
p	Matrix of the p-values for the coefficients (N x K)
par	Estimated model parameters including a scale parameter and a shrinkage parameter if penalty = TRUE (see Murakami et al., 2018)
e	Error statistics. It includes sum of squared errors (SSE), residual standard error (resid_SE), R-squared (R2), adjusted R2 (adjR2), log-likelihood (logLik), corrected Akaike information criterion (AICc), and the cross-validation (CV) score measured by root mean squared error (RMSE) (CV_score(RMSE))
pred	Vector of predicted values (N x 1)
resid	Vector of residuals (N x 1)
other	Other objects internally used

### References

Murakami, D., Tsutsumida, N., Yoshida, T., Nakaya, T., and Lu, B. (2019) Scalable GWR: A linear-time algorithm for large-scale geographically weighted regression with polynomial kernels. <arXiv:1905.00266>.

### See Also

[scgwr](#), [predict0](#)

### Examples

```
# require(spData);require(sp)
# data(house)
# dat <- data.frame(coordinates(house), house@data[,c("price", "age", "rooms", "beds", "syear")])
# coords<- dat[,c("long", "lat")]
# y <- log(dat[, "price"])
# x <- dat[,c("age", "rooms", "beds", "syear")]

# Parallel estimation
```

```
# res1 <- scgwr_p( coords = coords, y = y, x = x )
# res1

# Parallel estimation + Approximate cross-validation using 10000 samples
# res2 <- scgwr_p( coords = coords, y = y, x = x, nsamp = 10000 )
# res2
```

# Index

predict0, 1, 4, 5

scgwr, 3, 5

scgwr\_p, 3, 4, 4