

# How to use RandomForestsGLS

We will start by installing and loading the `RandomForestsGLS` R package. This package fits non-linear regression models on dependent data with Generalised Least Square (GLS) based Random Forest (RF-GLS) detailed in Saha, Basu and Datta (2020) <https://arxiv.org/abs/2007.15421>. We start by loading the package,

```
library(devtools)
install_github("ArkajyotiSaha/RandomForestsGLS", ref = "HEAD")
library(RandomForestsGLS)
```

Next, we discuss how the ‘RandomForestsGLS’ package can be used for estimation and prediction in a non-linear regression setup under correlated errors in different scenarios.

## 1. Spatial Data

We consider spatial point referenced data with the following model:

$$y_i = m(\mathbf{x}_i) + w(\mathbf{s}_i) + \epsilon_i;$$

where,  $y_i, \mathbf{x}_i$  denotes the response and the covariate corresponding to the  $i^{th}$  observed location  $\mathbf{s}_i$ .  $m(\mathbf{x}_i)$  denotes the covariate effect, spatial random effect,  $w(\mathbf{s})$  accounts for spatial dependence beyond covariates, and  $\epsilon$  accounts for the independent and identically distributed random Gaussian noise.

In the spatial mixture model setting, the package ‘RandomForestsGLS’ allows for fitting  $m(\cdot)$  using RF-GLS. Spatial random effects are modelled using Gaussian Process as is the practice. For model fitting, we use the computationally convenient Nearest Neighbor Gaussian Process (NNGP) (Datta, Banerjee, Finley, and Gelfand (2016)). Along with prediction of the mean function  $m(\cdot)$  we also offer kriging based prediction of spatial responses at new location.

## Simulation

Next we simulate a data from the following model:

$$y_i = 10 \sin(\pi x_i) + w(\mathbf{s}_i) + \epsilon_i; \quad \epsilon \sim N(\mathbf{0}, \tau^2 \mathbf{I}), \tau^2 = 0.1; \quad w \sim \text{exponential GP}; \quad \sigma^2 = 10; \phi = 1.$$

Here, the mean function is  $E(Y) = 10 \sin(\pi X)$ ;  $w$  accounts for the spatial correlation, which is modelled by a exponential Gaussian process with spatial variance  $\sigma^2 = 10$  and spatial correlation decay  $\phi = 1$ ; and  $\epsilon$  is the i.i.d random noise with variance  $\tau^2$ , which is also called the nugget in spatial literature.

For illustration purposes, we simulate with  $n = 200$ :

```
rmvn <- function(n, mu = 0, V = matrix(1)){
  p <- length(mu)
  if(any(is.na(match(dim(V),p))))
    stop("Dimension not right!")
  D <- chol(V)
  t(matrix(rnorm(n*p), ncol=p)%*%D + rep(mu,rep(n,p)))
}

set.seed(5)
```

```

n <- 200
coords <- cbind(runif(n,0,1), runif(n,0,1))
set.seed(2)
x <- as.matrix(runif(n),n,1)
sigma.sq = 10
phi = 1
tau.sq = 0.1
D <- as.matrix(dist(coords))
R <- exp(-phi*D)
w <- rmvn(1, rep(0,n), sigma.sq*R)
y <- rnorm(n, 10*sin(pi * x) + w, sqrt(tau.sq))

```

## Model fitting

In the package `RandomForestsGLS`, the working precision matrix used in the GLS-loss are NNGP approximations of precision matrices corresponding to Matérn covariance function.

In order to fit the model, the code requires:

- Coordinates ('coords'): an  $n \times 2$  matrix of 2-dimensional locations.
- Response ('y'): an  $n$  length vector of response at the observed coordinates.
- Covariates ('X'): an  $n \times p$  matrix of the covariates in the observation coordinates.
- Covariates for estimation ('Xest'): an  $n_{test} \times p$  matrix of the covariates where we want to estimate the function. Must have identical variables as that of 'X'. Default is 'X'.
- Minimum size of leaf nodes ('nthsize'): We recommend not setting this value too small, as that will lead to very deep trees that takes a lot of time to be built and can produce unstable estimates. Default value is 20.
- The parameters corresponding to the covariance function (detailed afterwards).

For the details on choice of other parameters, please refer to the help file of the code '`RFGLS_estimate_spatial`', which can be accessed with '`?RFGLS_estimate_spatial`'.

## Known Covariance Parameters

If the covariance parameters are known we set '`param_estimate = FALSE` (default value)'; the code additionally requires the following:

- Covariance Model ('cov.model'): Supported keywords are: "exponential", "matern", "spherical", and "gaussian" for exponential, Matérn, spherical and Gaussian covariance function respectively. Default value is "exponential".
- $\sigma^2$  ('sigma.sq'): The spatial variance. Default value is 1.
- $\tau^2$  ('tau.sq'): The nugget. Default value is 0.01.
- $\phi$  ('phi'): The spatial correlation decay parameter. Default value is 5.
- $\nu$  ('nu'): The smoothing parameter corresponding to the Matérn covariance function. Default value is 0.5.

We can fit the model as follows:

```

set.seed(1)
est_known <- RFGLS_estimate_spatial(coords, y, x, ntree = 50, cov.model = "exponential",
                                     nthsize = 20, sigma.sq = sigma.sq, tau.sq = tau.sq,
                                     phi = phi)

```

The estimate of the function at the covariates 'Xtest' is given in '`estimation_reult$predicted`'. For interpretation of the rest of the outputs, please see the help file of the code '`RFGLS_estimate_spatial`'. As far as choice of covariance models are concerned, the code for covariance models other than exponential model are in beta testing stage.

## Unknown Covariance Parameters

If the covariance parameters are not known we set 'param\_estimate = TRUE'; the code additionally requires the covariance model ('cov.model') to be used for parameter estimation prior to RF-GLS fitting. We fit the model with unknown covariance parameters as follows.

```
set.seed(1)
est_unknown <- RFGLS_estimate_spatial(coords, y, x, ntree = 50, cov.model = "exponential",
                                       nthsize = 20, param_estimate = TRUE)
```

## Prediction of mean function

Given a fitted model using 'RFGLS\_estimate\_spatial', we can estimate the mean function at new covariate values as follows:

```
Xtest <- matrix(seq(0,1, by = 1/10000), 10001, 1)
RFGLS_predict_known <- RFGLS_predict(est_known, Xtest)
```

## Performance comparison

We obtain the Mean Integrated Squared Error (MISE) of the estimates  $\hat{m}$  from RF-GLS on  $[0,1]$  and compare it with that corresponding to the classical Random Forest (RF) obtained using package 'randomForest' (with similar minimum nodesize, 'nodesize' = 20, as default 'nodesize' performs worse). We see that our method has a significantly smaller MISE. Additionally, we show that the MISE obtained with unknown parameters is comparable to that of the MISE obtained with known covariance parameters.

```
library(randomForest)
set.seed(1)
RF_est <- randomForest(x, y, nodesize = 20)

RF_predict <- predict(RF_est, Xtest)
#RF MISE
mean((RF_predict - 10*sin(pi * Xtest))^2)
#> [1] 8.36778

#RF-GLS MISE
mean((RFGLS_predict_known$predicted - 10*sin(pi * Xtest))^2)
#> [1] 0.150152

RFGLS_predict_unknown <- RFGLS_predict(est_unknown, Xtest)
#RF-GLS unknown MISE
mean((RFGLS_predict_unknown$predicted - 10*sin(pi * Xtest))^2)
#> [1] 0.1851947
```

We plot the true  $m(x) = 10\sin(\pi x)$  along with the loess-smoothed version of estimated  $\hat{m}(\cdot)$  obtained from RF-GLS and RF where we show that RF-GLS estimate approximates  $m(x)$  better than that corresponding to RF.

```
rfgls_loess_10 <- loess(RFGLS_predict_known$predicted ~ c(1:length(Xtest)), span=0.1)
rfgls_smoothed10 <- predict(rfgls_loess_10)

rf_loess_10 <- loess(RF_predict ~ c(1:length(RF_predict)), span=0.1)
rf_smoothed10 <- predict(rf_loess_10)

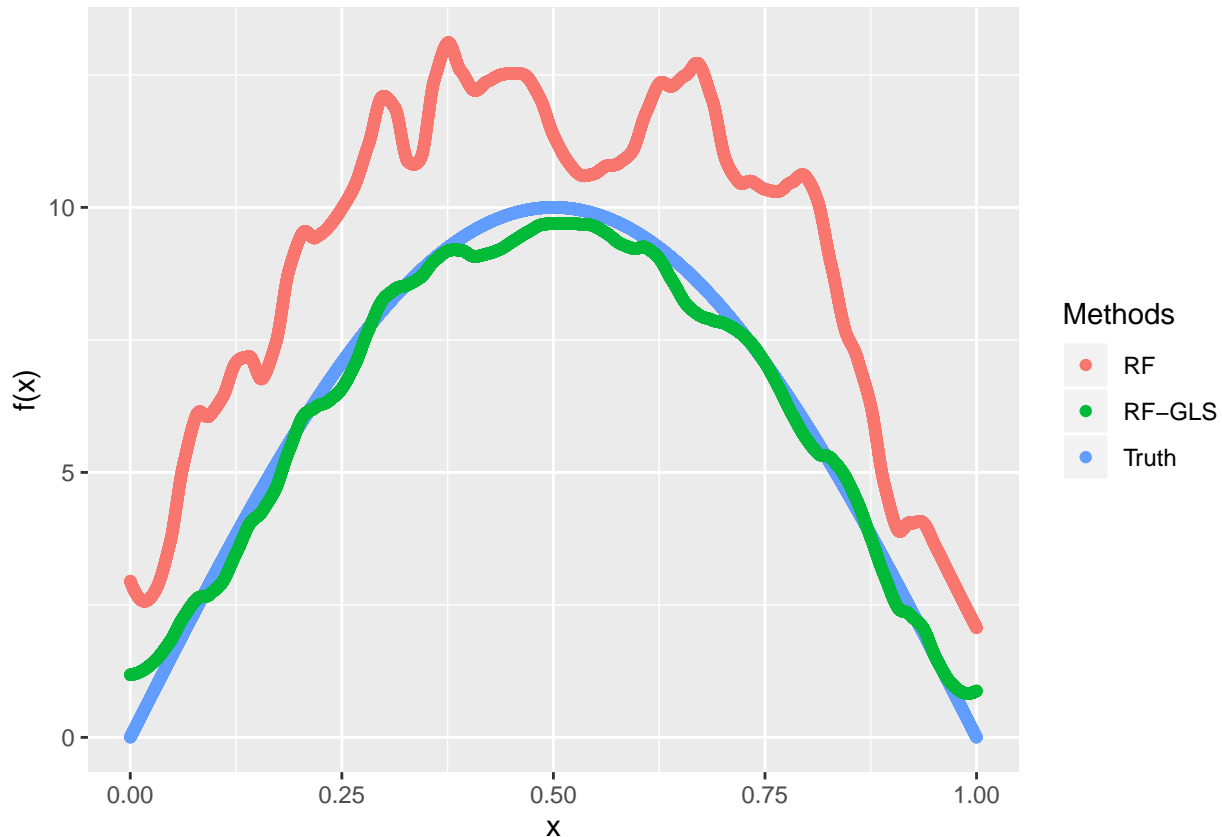
xval <- c(10*sin(pi * Xtest), rf_smoothed10, rfgls_smoothed10)
xval_tag <- c(rep("Truth", length(10*sin(pi * Xtest))), rep("RF", length(rf_smoothed10)),
```

```

      rep("RF-GLS",length(rfgls_smoothed10)))
plot_data <- as.data.frame(xval)
plot_data$Methods <- xval_tag
coval <- c(rep(seq(0,1, by = 1/10000), 3))
plot_data$Covariate <- coval

library(ggplot2)
ggplot(plot_data, aes(x=Covariate, y=xval, color=Methods)) +
  geom_point() + labs( x = "x") + labs( y = "f(x)")

```



## Prediction of spatial response

Given a fitted model using 'RFGLS\_estimate\_spatial', we can predict the spatial response/outcome at new locations provided the covariates at that location. This approach performs kriging at a new location using the mean function estimates at the corresponding covariate values. Here we partition the simulated data into training and test sets in 4:1 ratios. Next we perform prediction on the test set using a model fitted on the training set.

```

est_known_short <- RFGLS_estimate_spatial(coords[1:160,], y[1:160],
      matrix(x[1:160,],160,1), ntree = 50, cov.model = "exponential",
      nthsize = 20, param_estimate = TRUE)
RFGLS_predict_spatial <- RFGLS_predict_spatial(est_known_short, coords[161:200,],
      matrix(x[161:200,],40,1))

```

## Misspecification in covariance model

When the covariance parameters are not known, we show that RF-GLS can outperform classical RF even when the parameters are estimated from a misspecified covariance model. In order to show this, we simulate the spatial correlation from a Matérn covariance function with smoothing parameter  $\nu = 1.5$ . While fitting the RF-GLS, we estimate the covariance parameters using an exponential covariance model ( $\nu = 0.5$ ) and show that the obtained MISE can compare favorably to that of classical RF.

```
#Data simulation from matern with nu = 1.5
nu = 3/2
R1 <- (D*phi)^nu/(2^(nu-1)*gamma(nu))*besselK(x=D*phi, nu=nu)
diag(R1) <- 1
set.seed(2)
w <- rmvn(1, rep(0,n), sigma.sq*R1)
y <- rnorm(n, 10*sin(pi * x) + w, sqrt(tau.sq))

#RF-GLS with exponential covariance
set.seed(3)
est_misspec <- RFGLS_estimate_spatial(coords, y, x, ntree = 50, cov.model = "exponential",
                                     nthsize = 20, param_estimate = TRUE)
RFGLS_predict_misspec <- RFGLS_predict(est_misspec, Xtest)

set.seed(4)
RF_est <- randomForest(x, y, nodesize = 20)
RF_predict <- predict(RF_est, Xtest)

#RF-GLS MISE
mean((RFGLS_predict_misspec$predicted - 10*sin(pi * Xtest))^2)
#> [1] 0.1380615
#RF MISE
mean((RF_predict - 10*sin(pi * Xtest))^2)
#> [1] 2.295639
```

## Parallelization

For both ‘RFGLS\_estimate\_spatial’ and ‘RFGLS\_predict’, one can also take the advantage of parallelization, contingent upon the availability of multiple cores. The component ‘h’ in both the codes determines the number of cores to be used. Here we demonstrate an example with ‘h = 4’.

```
#simulation from exponential distribution
set.seed(5)
n <- 200
coords <- cbind(runif(n,0,1), runif(n,0,1))
set.seed(2)
x <- as.matrix(runif(n),n,1)
sigma.sq = 10
phi = 1
tau.sq = 0.1
nu = 0.5
D <- as.matrix(dist(coords))
R <- exp(-phi*D)
w <- rmvn(1, rep(0,n), sigma.sq*R)
y <- rnorm(n, 10*sin(pi * x) + w, sqrt(tau.sq))
```

```

set.seed(1)
est_known_pl <- RFGLS_estimate_spatial(coords, y, x, ntree = 50, cov.model = "exponential",
                                       nthsize = 20, sigma.sq = sigma.sq, tau.sq = tau.sq,
                                       phi = phi, h = 4)
RFGLS_predict_known_pl <- RFGLS_predict(est_known_pl, Xtest, h = 4)
#MISE from single core
mean((RFGLS_predict_known_pl$predicted - 10*sin(pi * Xtest))^2)
#> [1] 0.150152
#MISE from parallel computation
mean((RFGLS_predict_known_pl$predicted - 10*sin(pi * Xtest))^2)
#> [1] 0.150152

```

For ‘RFGLS\_estimate\_spatial’ very small dataset (‘n’) and small number of trees (‘ntree’), communication overhead between the nodes for parallelization outweighs the benefits of the parallel computing hence it is recommended to parallelize moderately high ‘n’ and/or ‘ntree’. It is strongly recommended that the max value of ‘h’ is kept strictly less than the number of total cores available. For ‘RFGLS\_predict’, even for large dataset, single core performance is very fast, hence unless ‘ntest’ and ‘ntree’ are very high, we do not recommend using parallelization for ‘RFGLS\_predict’.