



INLA and inlabru with spatial patterns

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Overzicht

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- Pearson residuals
- Variogram

2 Prepare the model

- Creating a mesh
- Creating an SPDE model

3 Fitting the model

- Only the data
- Predictions





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Checking spatial auto- correlation



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Checking spatial auto- correlation

Pearson residuals

Definition

► components?



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Definition

- ▶ components?
- ▶ observed value (y_i), fitted value (\hat{y}_i), mean squared error (MSE)



Definition

- ▶ components?
- ▶ observed value (y_i), fitted value (\hat{y}_i), mean squared error (MSE)
- ▶ formula



Definition

- ▶ components?
- ▶ observed value (y_i), fitted value (\hat{y}_i), mean squared error (MSE)
- ▶ formula
- ▶

$$pr_i = \frac{y_i - \hat{y}_i}{\sqrt{MSE}}$$



Definition

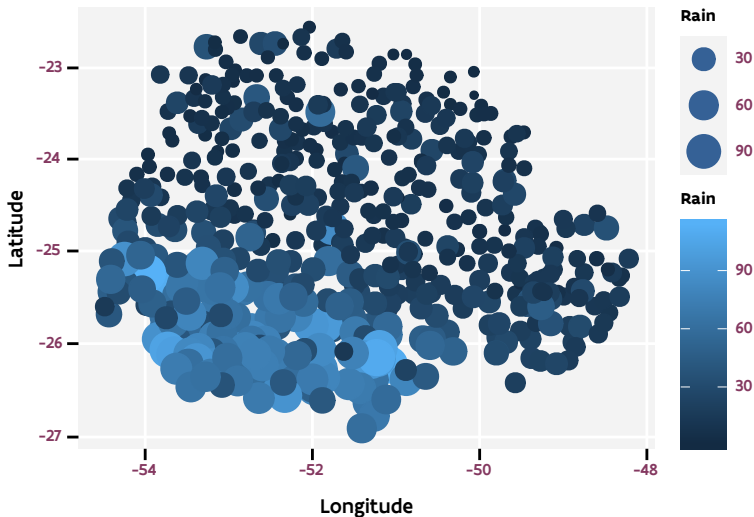
- ▶ components?
- ▶ observed value (y_i), fitted value (\hat{y}_i), mean squared error (MSE)
- ▶ formula
- ▶

$$pr_i = \frac{y_i - \hat{y}_i}{\sqrt{MSE}}$$

- ▶ MSE (variance) depends on distribution! Check it using `inla.doc("name_of_your_distribution")`.



Example data: rainfall in Parana state, Brazil



Calculate Pearson residuals

```
model_iid <- inla(Rain ~ Xc + Yc, family = "gamma", data = dataset,  
                 control.compute = list(waic = TRUE))  
dataset %>%  
  mutate(  
    mu = model_iid$summary.fitted.values$mean,  
    sigma2 = mu ^ 2 / model_iid$summary.hyperpar[1, "mean"],  
    Pearson_iid = (Rain - mu) / sqrt(sigma2)  
  ) -> dataset
```



Challenge 1

- ▶ What is the mean for your model?
- ▶ What is the variance for your model? Hint: `inla.doc("your distribution")`
- ▶ Calculate the Pearson residuals for your model





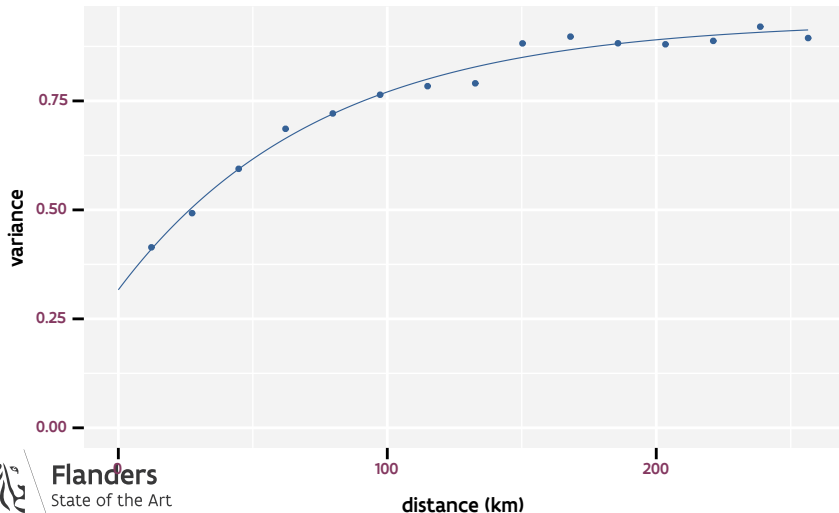
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Checking spatial auto- correlation

Variogram

Definition

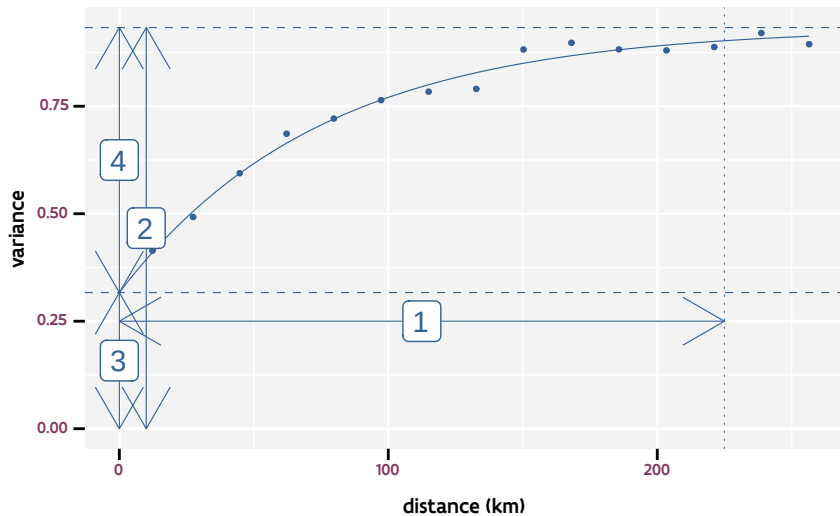
```
vg_default <- variogram(Pearson_iid ~ 1, locations = ~X + Y,  
                        data = as.data.frame(dataset), cressie = TRUE)
```



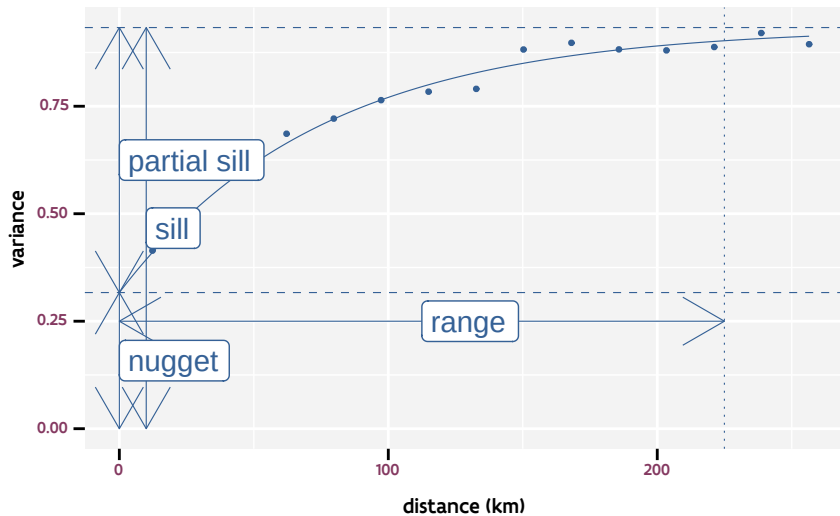
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distance (km)

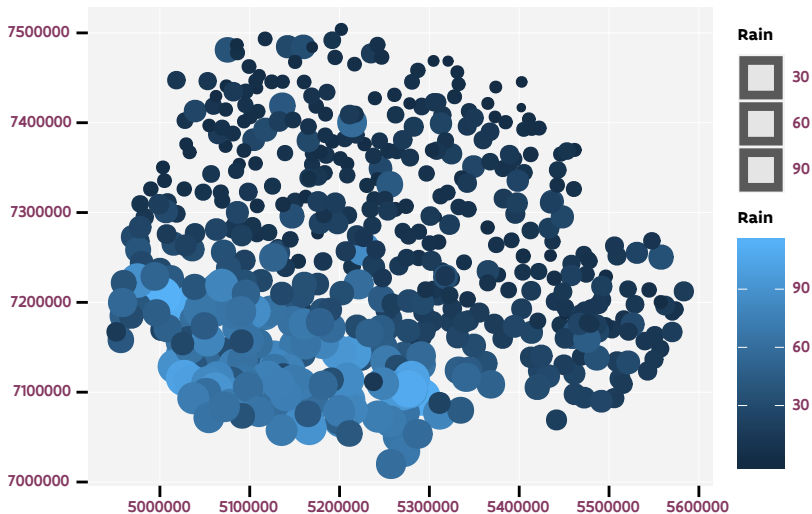
Important characteristics



Important characteristics



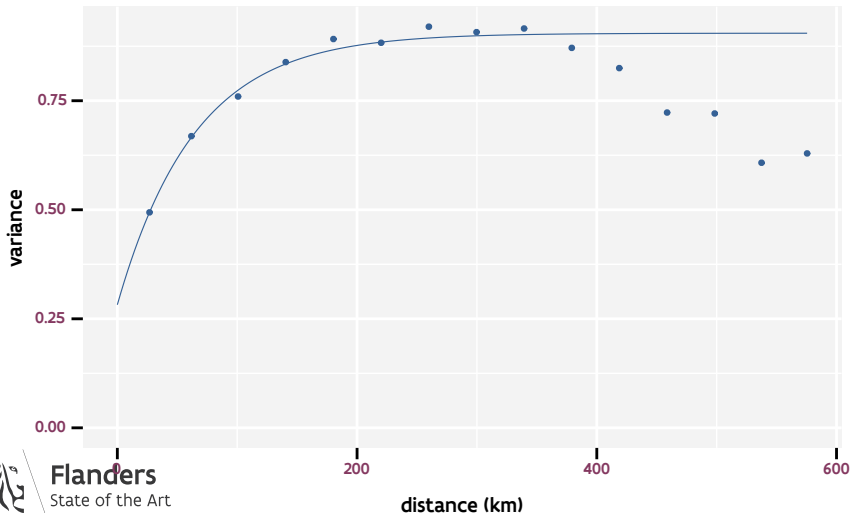
Projected example data



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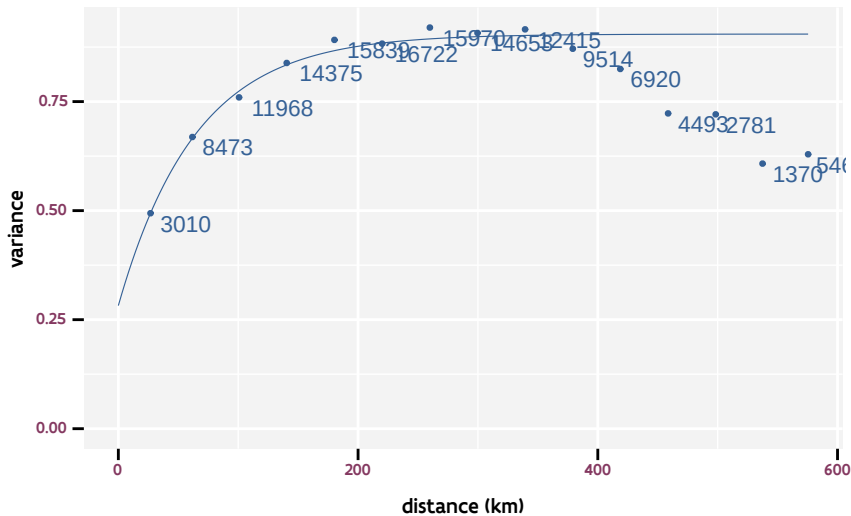
Increased cutoff

```
vg_large <- variogram(Pearson_iid ~ 1, locations = ~X + Y, cressie = TRUE,  
  data = as.data.frame(dataset), cutoff = 600e3)
```



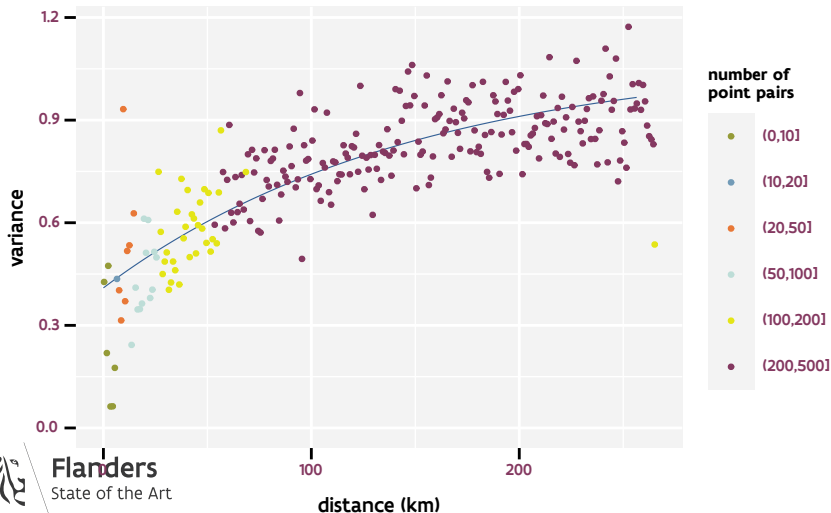
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Number of point pairs is important



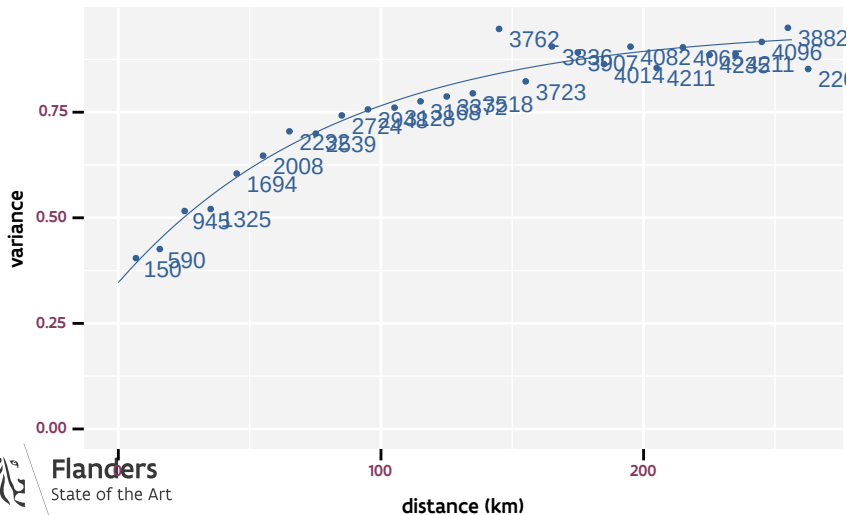
Too small width leads to unstable variograms

```
vg_small <- variogram(Pearson_iid ~ 1, locations = ~X + Y, cressie = TRUE,  
                      data = as.data.frame(dataset), width = 1e3)
```



Sensible small width yields the most informative variogram

```
vg_final <- variogram(Pearson_iid ~ 1, locations = ~X + Y, cressie = TRUE,  
  data = as.data.frame(dataset), width = 10e3)
```



Challenge 2

- ▶ What is the minimum binwidth for your data?
- ▶ Calculate the variogram for your model
- ▶ What is the approximate range of of the variogram?
- ▶ What is the nugget, sill and partial sill?





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Prepare the model

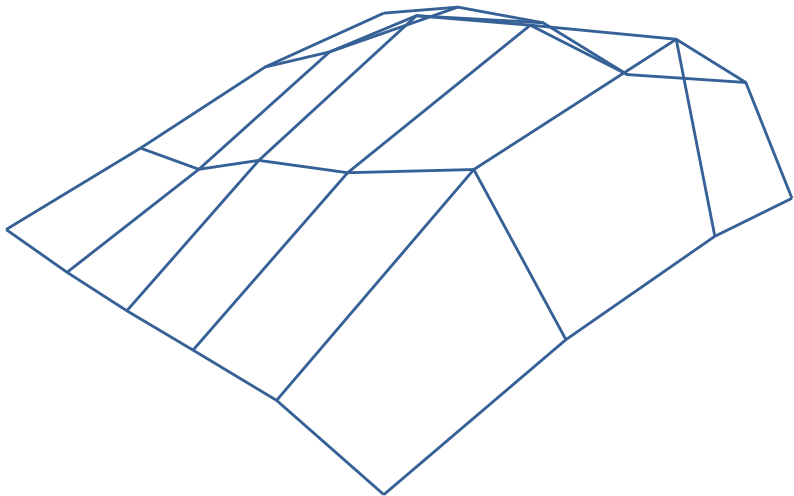


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Prepare the model

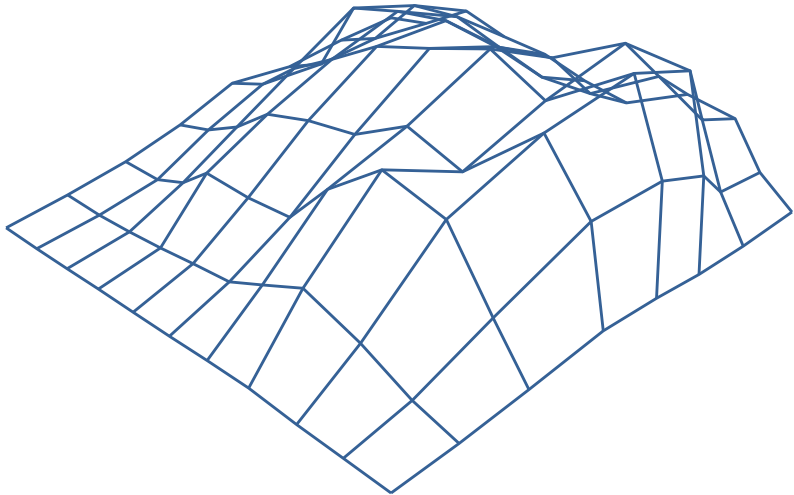
Creating a mesh

Size of a mesh I

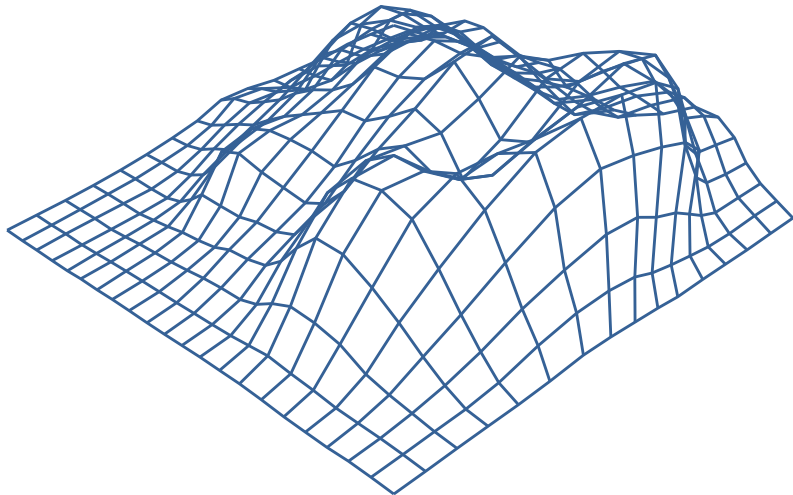


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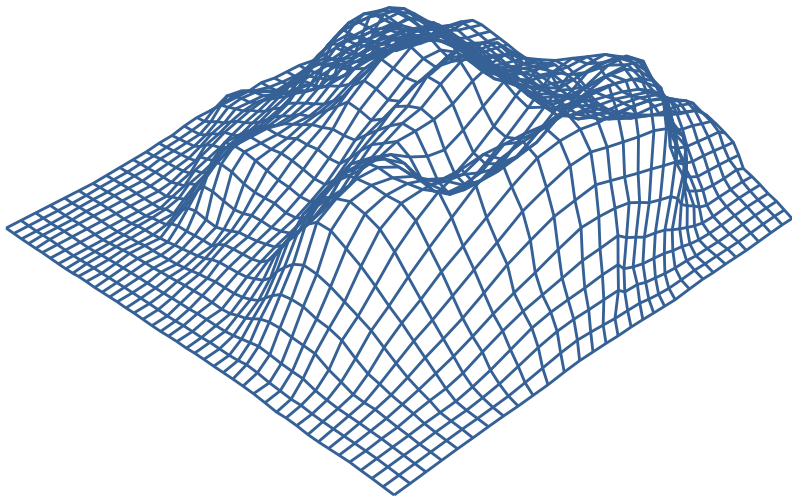
Size of a mesh II



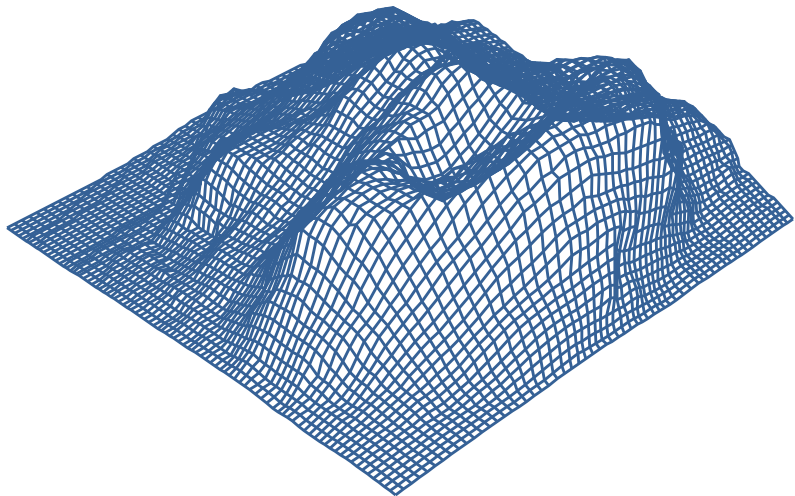
Size of a mesh III



Size of a mesh IV



Size of a mesh V



Guidelines

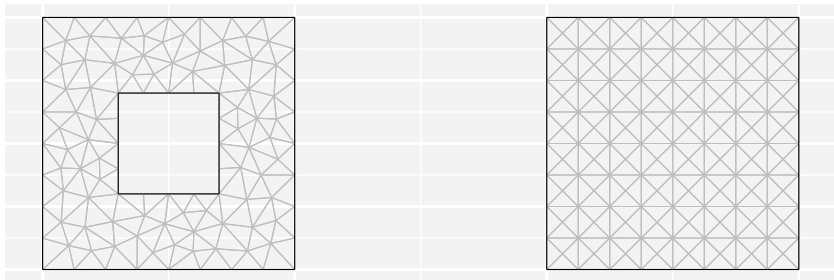
- ▶ equilateral triangles work best
- ▶ edge length should be around a third to a tenth of the range
- ▶ avoid narrow triangles
- ▶ avoid small edges
- ▶ add extra, larger triangles around the border
- ▶ simplify the border



Mesh only within the border

```
mesh <- inla.mesh.2d(boundary = border, max.edge = 0.15)  
ggplot() + gg(mesh) + coord_fixed() + theme_map() +  
  ggtitle(paste("Vertices: ", mesh$n))
```

Vertices: 261

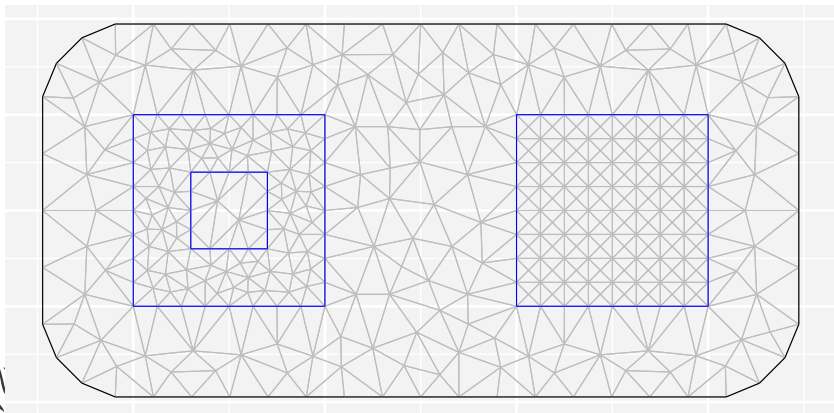


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Mesh going outside the border

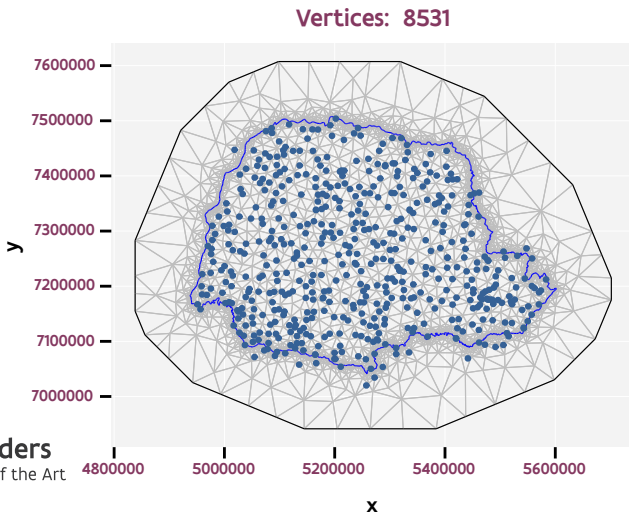
```
mesh <- inla.mesh.2d(boundary = border, max.edge = c(0.15, 0.3))  
ggplot() + gg(mesh) + coord_fixed() + theme_map() +  
  ggtitle(paste("Vertices: ", mesh$n))
```

Vertices: 417



Mesh for rainfall data

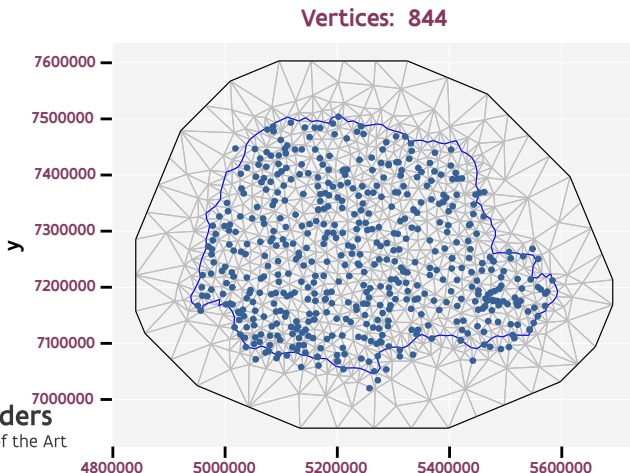
```
mesh <- inla.mesh.2d(boundary = boundary, max.edge = c(30e3, 100e3))  
ggplot(dataset) + gg(mesh) + geom_sf() + ggtitle(paste("Vertices: ", mesh$n)) +  
  coord_sf(datum = st_crs(5880))
```



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Use cutoff to simplify mesh

```
mesh1 <- inla.mesh.2d(boundary = boundary, max.edge = c(30e3, 100e3),  
                      cutoff = 10e3)  
ggplot(dataset) + gg(mesh1) + geom_sf() +  
  ggtitle(paste("Vertices: ", mesh1$n)) + coord_sf(datum = st_crs(5880))
```

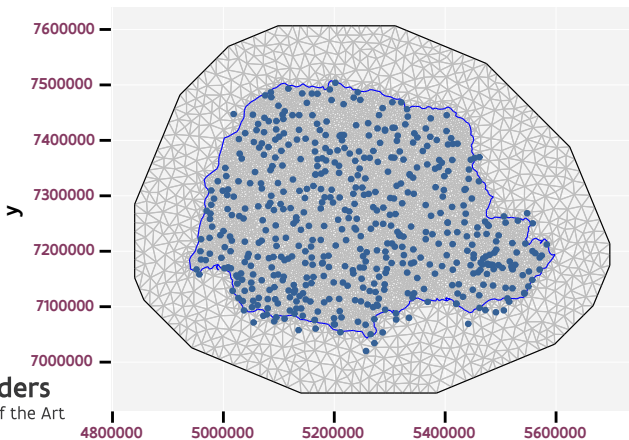


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Finer mesh for final model run

```
mesh2 <- inla.mesh.2d(boundary = boundary, max.edge = c(10e3, 30e3),  
                      cutoff = 5e3)  
ggplot(dataset) + gg(mesh2) + geom_sf() +  
  ggtitle(paste("Vertices: ", mesh2$n)) + coord_sf(datum = st_crs(5880))
```

Vertices: 5920



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Challenge 3

- ▶ What are the relevant max.edge and cutoff for a course mesh?
- ▶ What are the relevant max.edge and cutoff for a smooth mesh?
- ▶ Create a course and a smooth mesh for your data





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Prepare the model

Creating an SPDE model

SPDE using penalised complexity priors

Stochastic Partial Differential Equations

- ▶ $\text{prior.range} = c(r, \alpha_r): P(\rho < r) < \alpha_r$
- ▶ $\text{prior.sigma} = c(s, \alpha_s): P(\sigma > s) < \alpha_s$

```
spde1 <- inla.spde2.pcmatern(mesh1, prior.range = c(100e3, 0.5),  
                             prior.sigma = c(0.9, 0.05))  
spde2 <- inla.spde2.pcmatern(mesh2, prior.range = c(100e3, 0.5),  
                             prior.sigma = c(0.9, 0.05))
```



Challenge 4

- ▶ What are relevant priors for the range and sigma for your data
 - ▶ Hint: see challenge 2
- ▶ Make the SPDE models for your data





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Fitting the model



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Fitting the model

Only the data

The stack for the observed data

```
A1 <- inla.spde.make.A(mesh = mesh1, loc = st_coordinates(dataset))
stack1 <- inla.stack(
  tag = "estimation", ## tag
  data = list(Rain = dataset$Rain), ## response
  A = list(A1, 1), ## projector matrices (SPDE and fixed effects)
  effects = list(
    list(site = seq_len(spde1$n.spde)), ## random field index
    dataset %>%
      as.data.frame() %>%
      transmute(Intercept = 1, Xc, Yc) ## fixed effect covariates
  )
)
```



Model fit

INLA

```
model_spde1 <- inla(Rain ~ 0 + Intercept + Xc + Yc + f(site, model = spde1),  
  family = "gamma", data = inla.stack.data(stack1),  
  control.predictor = list(A = inla.stack.A(stack1)),  
  control.compute = list(waic = TRUE)  
)
```

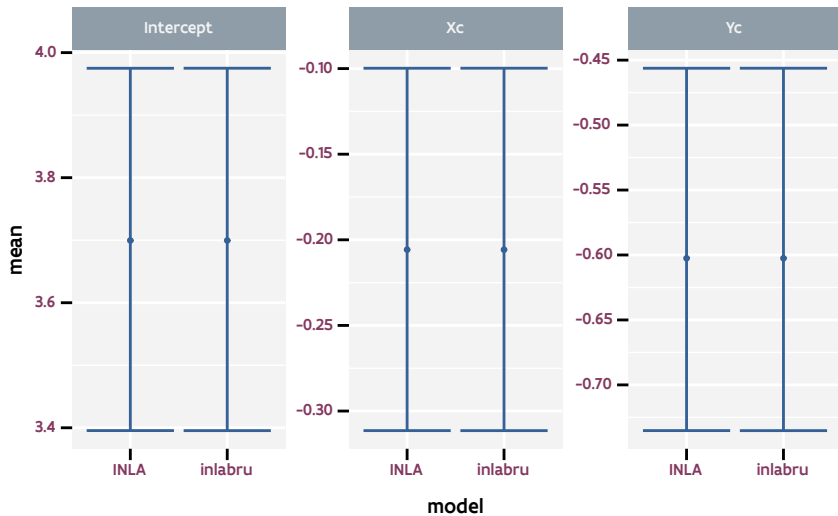
inlabru

```
bru_spde1 <- bru(Rain ~ Xc + Yc + site(map = st_coordinates, model = spde1),  
  family = "gamma", data = dataset)
```

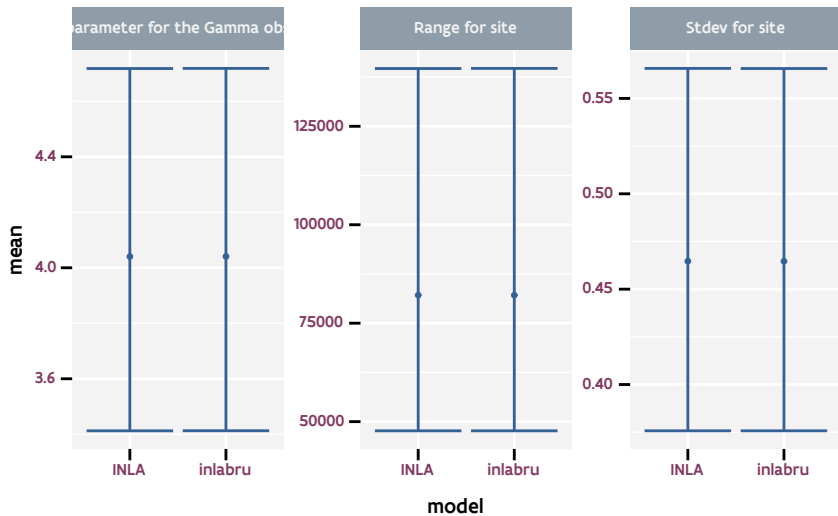
```
bru_spde1 <- bru(Rain ~ Xc + Yc + site(map = coordinates, model = spde1),  
  family = "gamma", data = as_Spatial(dataset))
```



Comparison of fixed effect parameters



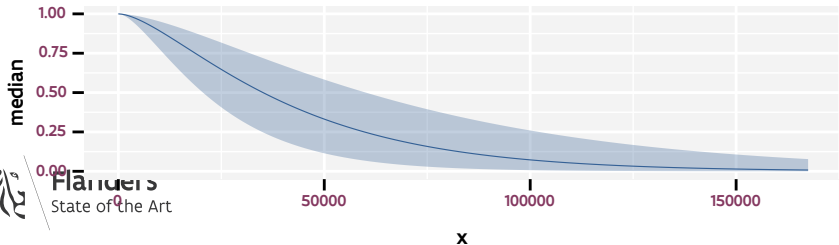
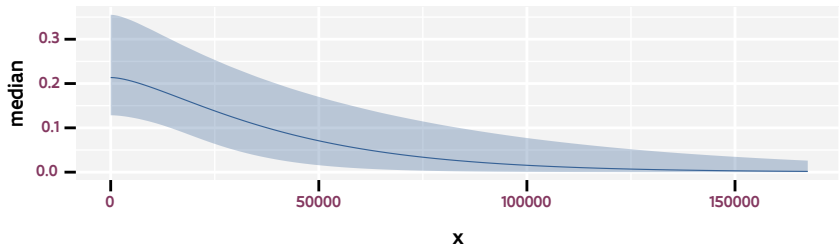
Comparing hyperparameters



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Correlation structure

```
spde.posterior(bru_spde1, "site", what = "matern.covariance") -> covplot  
spde.posterior(bru_spde1, "site", what = "matern.correlation") -> corplot  
multiplot(plot(covplot), plot(corplot))
```



Calculate Pearson residuals

```
dataset %>%  
  mutate(  
    mu = model_spde1$summary.fitted.values$mean,  
    sigma2 = mu ^ 2 / model_spde1$summary.hyperpar[1, "mean"],  
    Pearson_iid = (Rain - mu) / sqrt(sigma2)  
  ) -> dataset
```

Error: Column `mu` must be length 528 (the number of rows) or one, not 1664



Using the stack index

```
si <- inla.stack.index(stack1, "estimation")$data
dataset %>%
  mutate(
    mu = model_spde1$summary.fitted.values$mean[si],
    sigma2 = mu ^ 2 / model_spde1$summary.hyperpar[1, "mean"],
    Pearson_spde = (Rain - mu) / sqrt(sigma2)
  ) -> dataset
```



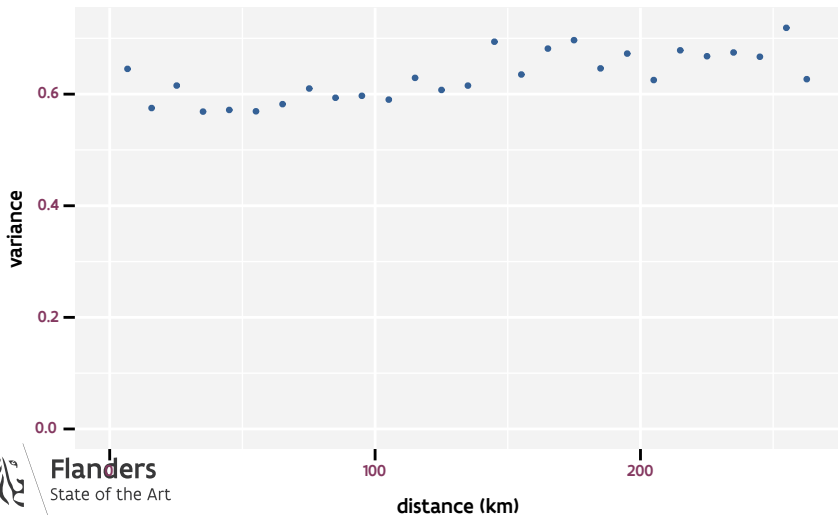
Using inlabru

```
fit <- predict(bru_spde1, as_Spatial(dataset), ~exp(Intercept + Xc + Yc + site))
dataset %>%
  mutate(
    mu = fit$mean,
    sigma2 = mu ^ 2 / model_spde1$summary.hyperpar[1, "mean"],
    Pearson_spde = (Rain - mu) / sqrt(sigma2)
  ) -> dataset
```



Variogram

```
vg_fit <- variogram(Pearson_spde ~ 1, cressie = TRUE,  
  data = as_Spatial(dataset), width = 10e3)
```



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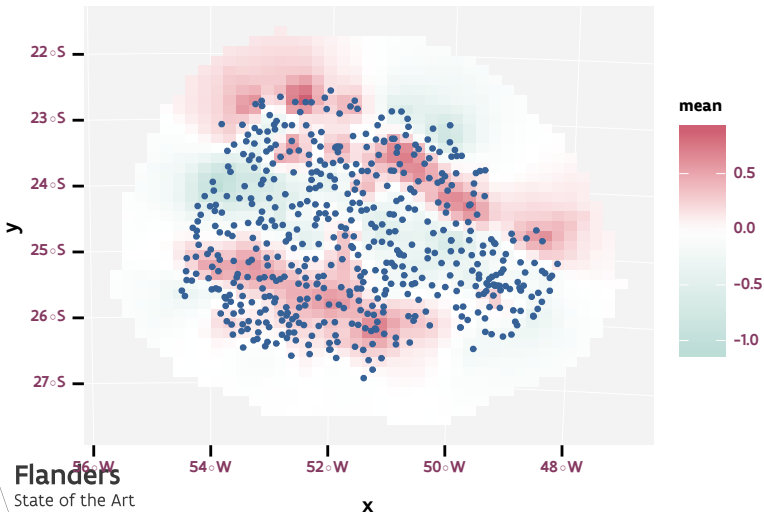
Interpolate GMRF

```
A1.grid <- inla.mesh.projector(mesh1, dims = c(41, 41))
inla.mesh.project(A1.grid, model_spde1$summary.random$site) %>%
  as.matrix() %>%
  as.data.frame() %>%
  bind_cols(
    expand.grid(x = A1.grid$x, y = A1.grid$y)
  ) %>%
  filter(!is.na(ID)) -> eta_spde
```



Plot GMRF

```
ggplot(dataset) + geom_tile(data = eta_spde, aes(x = x, y = y, fill = mean)) +  
  geom_sf() + scale_fill_gradient2()
```



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Fitting the model

Predictions

Prediction stack for SPDE grid + fixed effects

```
expand.grid(X = A1.grid$x, Y = A1.grid$y) %>%  
  mutate(Intercept = 1, Xc = X / 1e5 - 53, Yc = Y / 1e5 - 71) -> grid_data  
stack1_grid <- inla.stack(  
  tag = "grid", ## tag  
  data = list(Rain = NA), ## response  
  A = list(A1.grid$proj$A, 1), ## projector matrices (SPDE and fixed effects)  
  effects = list(  
    list(site = seq_len(spde1$n.spde)), ## random field index  
    grid_data ## covariates at grid locations  
  )  
)
```



Refit the model with the combined stack

```
stack_all <- inla.stack(stack1, stack1_grid)
model_grid <- inla(Rain ~ 0 + Intercept + Xc + Yc + f(site, model = spde1),
  family = "gamma", data = inla.stack.data(stack_all),
  control.predictor = list(A = inla.stack.A(stack_all),
    link = 1),
  control.compute = list(waic = TRUE),
  control.mode = list(theta = model_spde1$mode$theta,
    restart = FALSE),
  control.results = list(return.marginals.random = FALSE,
    return.marginals.predictor = FALSE)
)
```

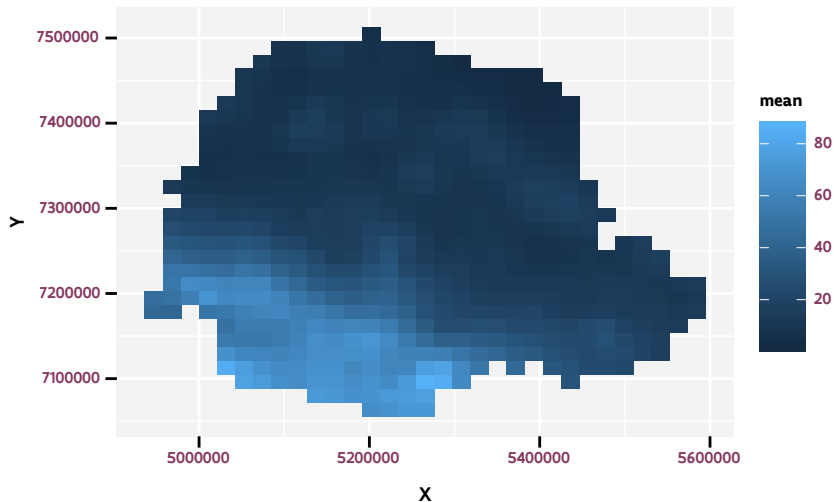


Plot grid I

```
si <- inla.stack.index(stack_all, "grid")$data
grid_data %>%
  bind_cols(model_grid$summary.fitted.values[si, ]) %>%
  `coordinates<-`(~X + Y) %>%
  `proj4string<-`(CRS("+init=epsg:5880")) -> gd
gd[!is.na(over(gd, boundary)), ] %>%
  as.data.frame() %>%
  ggplot() + geom_tile(aes(x = X, y = Y, fill = mean)) + coord_fixed()
```

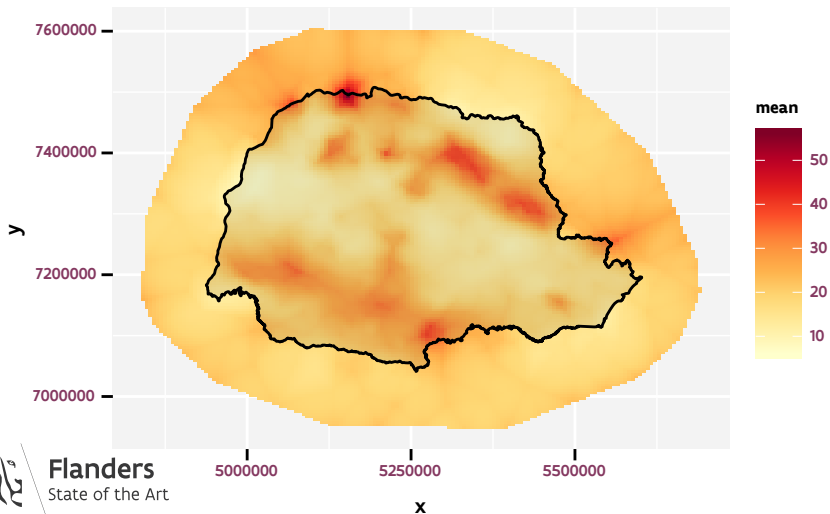


Plot grid II



Using inlabru

```
pred_mesh <- predict(bru_spde1, pixels(mesh1), ~exp(Intercept + Xc + Yc + site))  
ggplot() + gg(pred_mesh) + gg(boundary)
```



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Challenge 5

- ▶ Fit the model using the SPDE
- ▶ Plot a map of the GMRF
- ▶ Plot a map of the predictions and their credible interval

