## Spatial lag model for spatial effects

#### **Parametrization**

The slm model is defined as

$$\mathbf{x} = (I_n - \rho W)^{-1} (X\beta + \varepsilon)$$

where  $I_n$  is the identity matrix of dimension  $n \times n$ , W is a spatial weights matrix, X is a matrix of covariates,  $\rho$  is a spatial autocorrelation parameter,  $\beta$  are coefficients of the covariates and  $\varepsilon$  is zero mean Gaussian noise with precision  $\tau$ .

#### Hyperparameters

This model has two hyperparameters  $\theta = (\theta_1, \theta_2)$ . The precision parameter  $\tau$  is represented as

$$\theta_1 = \log \tau$$

and the prior is defined on  $\theta_1$ . The spatial autocorrelation parameter  $\rho$  is represented as

$$\rho^* = \frac{\rho - \rho_{\min}}{\rho_{\max} - \rho_{\min}}$$

and then

$$\theta_2 = \log(\rho^*/(1 - \rho^*))$$

and the prior is defined on  $\theta_2$ . Here,  $\rho_{\min}$  and  $\rho_{\max}$  are lower and upper limits of the legal range for  $\rho$ .

#### **Specification**

The slm model is specified inside the f() function as

args.slm is used to define the slm-spesific parameters in the model.

**rho.min and rho.max** define the range in which  $\rho$  can take values. Note that,  $\rho^*$  is in the interval (0,1) and that it is re-scaled to the interval (rho.min, rho.max) when computing  $I_n - \rho W$ . Initial values on  $\rho$  need to be re-scaled to the (0,1) interval.

**X** defines the matrix of covariates.

**W** defines the adjacency matrix.

**Q.beta** defines the precision of the vector of coefficients  $\beta$  in the model.

# Hyperparameter spesification and default values hyper

```
theta1
         name log precision
         short.name prec
         initial 4
         fixed FALSE
         prior loggamma
         param 1 5e-05
         to.theta function(x) log(x)
         from.theta function(x) exp(x)
     theta2
         name rho
         short.name rho
         initial 0
         fixed FALSE
         prior normal
         param 0 10
         to.theta function(x) log(x/(1-x))
         from.theta function(x) 1/(1+exp(-x))
constr FALSE
nrow.ncol FALSE
augmented FALSE
aug.factor 1
aug.constr
n.div.by
n.required TRUE
set.default.values TRUE
pdf slm
status experimental
Example
## Example using the Boston dataset from package spdep
require(INLA)
require(spdep)
data(boston)
## Index for the latent model
n <- nrow(boston.c)</pre>
boston.c$idx <- 1:n
```

```
## Define adjacency using a row-standardised matrix
lw <- nb2listw(boston.soi)</pre>
W <- as(as_dgRMatrix_listw(lw), "CsparseMatrix")</pre>
## Model definition
f1 <- log(CMEDV) ~ CRIM + ZN + INDUS + CHAS + I(NOX^2)+ I(RM^2) + AGE +
  log(DIS) + log(RAD) + TAX + PTRATIO + B + log(LSTAT)
mmatrix <- model.matrix(f1, boston.c)</pre>
## Zero-variance for error term
zero.variance = list(prec=list(initial = 25, fixed=TRUE))
## Compute eigenvalues for SLM model, used to obtain rho.min and
## rho.max
e = eigenw(lw)
re.idx = which(abs(Im(e)) < 1e-6)
rho.max = 1/max(Re(e[re.idx]))
rho.min = 1/min(Re(e[re.idx]))
rho = mean(c(rho.min, rho.max))
## Precision matrix for beta coefficents' prior
betaprec <- .0001
Q.beta = Diagonal(n=ncol(mmatrix), betaprec)
## Priors on the hyperparameters
hyper = list(
        prec = list(
                prior = "loggamma",
                param = c(0.01, 0.01)),
        rho = list(
                initial=0,
                prior = "logitbeta",
                param = c(1,1))
## Fit model
slmm1 <- inla( log(CMEDV) ~ -1 +</pre>
              f(idx, model="slm",
                args.slm=list(
                         rho.min = rho.min,
                         rho.max = rho.max,
                         W=W,
                         X=mmatrix,
                         Q.beta=Q.beta),
                hyper=hyper),
              data=boston.c, family="gaussian",
              control.family = list(hyper=zero.variance),
              control.compute=list(dic=TRUE, cpo=TRUE)
summary(slmm1)
## Summary of the coefficients (at the end of the vector of random effects)
slmm1$summary.random$idx[n+1:ncol(mmatrix),]
## Re-scale rho to real scale
rhomarg <- inla.tmarginal(function(x){rho.min+x*(rho.max-rho.min)},</pre>
```

### slmm1\$marginals.hyperpar[[2]])

inla.zmarginal(rhomarg)

```
## Maximum likelihood estimate of model (used for comparison)
summary(m2 <- lagsarlm(f1, boston.c, lw))</pre>
```

#### Notes

The estimates of  $\beta$  are included at the end of the vector of random effects. See the example for details on how to extract them.