Smooth-Copy of another model component: "scopy"

This model is a generalization of `copy`, please refer to `inla.doc("copy")` first.

This describes the way to copy another model component with an optional smooth/spline scaling, like with

$$\eta = u + v$$

where v is a smooth copy of u (component-wise)

$$v = \beta(z) \times \text{copy}(u)$$

where $\beta(z)$, a smooth/spline function of the covariate z. The smooth scaling is done **component-wise** for u, so if u are defined with domain $(1, 2, \ldots, m)$, i.e. $u = (u_1, u_2, \ldots, u_m)$, then z must be $z = (z_1, z_2, \ldots, z_m)$, so that

$$v_i = \beta(z_i)u_i, \qquad i = 1, 2, \ldots, m.$$

Hyperparameters

The hyperparameters are the value of the spline at n fixed locations, (l_i, β_i), for $i = 1, \ldots, n$. The function $\beta(z)$, is defined as (using z as the covariate)

```
zr <- range(z)
l <- seq(zr[1], zr[2], len=n)
beta.z <- splinefun(l, beta, method = "natural")
```

We can control β and its prior distribution using argument `control.scopy` within `f()`,

```
control.scopy = list(
covariate = ...,
n = 5,
model = "rw2",
mean = 1.0,
prec.mean = 1.0,
prec.betas = 10.0)
```

where

covariate gives the covariate that is used

n is the number of hyperparameters used in the spline ($3 \le n \le 15$).

model the prior model for $\{\beta_i\}$, either `rw1` or `rw2`. This model is scaled (like with `scale.model=TRUE`).

mean The prior mean for the (weighted-)mean[1] of $\{\beta_i\}$

prec.mean The prior precision for the (weighted-)mean of $\{\beta_i\}$

prec.betas The prior precision for the `rw1/rw2` model for $\{\beta_i\}$

Note that the prior mean and both prior precisions, are *fixed* and not *random*.

The `f()`-argument `precision`, defines how close the copy is, is similar as for model `copy`.

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[1]The mean of $\{\beta_i\}$ is defined to approximate the integral of the RW, hence its $\left(\frac{1}{2}(\beta_1 + \beta_n) + \sum_{j=2}^{n-1} \beta_j\right)/(n-1)$.

Spesification

doc `Create a scopy of a model component`

hyper

    theta1

        hyperid `36101`
        name `beta1`
        short.name `b1`
        initial `0.1`
        fixed `FALSE`
        prior `none`
        param
        to.theta `function(x) x`
        from.theta `function(x) x`

    theta2

        hyperid `36102`
        name `beta2`
        short.name `b2`
        initial `0.1`
        fixed `FALSE`
        prior `none`
        param
        to.theta `function(x) x`
        from.theta `function(x) x`

    theta3

        hyperid `36103`
        name `beta3`
        short.name `b3`
        initial `0.1`
        fixed `FALSE`
        prior `none`
        param
        to.theta `function(x) x`
        from.theta `function(x) x`

    theta4

        hyperid `36104`
        name `beta4`
        short.name `b4`
        initial `0.1`
        fixed `FALSE`
        prior `none`
        param
        to.theta `function(x) x`

from.theta `function(x) x`

theta5

    hyperid `36105`

    name `beta5`

    short.name `b5`

    initial `0.1`

    fixed `FALSE`

    prior `none`

    param

    to.theta `function(x) x`

    from.theta `function(x) x`

theta6

    hyperid `36106`

    name `beta6`

    short.name `b6`

    initial `0.1`

    fixed `FALSE`

    prior `none`

    param

    to.theta `function(x) x`

    from.theta `function(x) x`

theta7

    hyperid `36107`

    name `beta7`

    short.name `b7`

    initial `0.1`

    fixed `FALSE`

    prior `none`

    param

    to.theta `function(x) x`

    from.theta `function(x) x`

theta8

    hyperid `36108`

    name `beta8`

    short.name `b8`

    initial `0.1`

    fixed `FALSE`

    prior `none`

    param

    to.theta `function(x) x`

    from.theta `function(x) x`

theta9

    hyperid `36109`

    name `beta9`

short.name `b9`

initial `0.1`

fixed `FALSE`

prior `none`

param

to.theta `function(x) x`

from.theta `function(x) x`

theta10

hyperid `36110`

name `beta10`

short.name `b10`

initial `0.1`

fixed `FALSE`

prior `none`

param

to.theta `function(x) x`

from.theta `function(x) x`

theta11

hyperid `36111`

name `beta11`

short.name `b11`

initial `0.1`

fixed `FALSE`

prior `none`

param

to.theta `function(x) x`

from.theta `function(x) x`

theta12

hyperid `36112`

name `beta12`

short.name `b12`

initial `0.1`

fixed `FALSE`

prior `none`

param

to.theta `function(x) x`

from.theta `function(x) x`

theta13

hyperid `36113`

name `beta13`

short.name `b13`

initial `0.1`

fixed `FALSE`

prior `none`

param
to.theta `function(x) x`
from.theta `function(x) x`

theta14

hyperid `36114`
name `beta14`
short.name `b14`
initial `0.1`
fixed `FALSE`
prior `none`
param
to.theta `function(x) x`
from.theta `function(x) x`

theta15

hyperid `36115`
name `beta15`
short.name `b15`
initial `0.1`
fixed `FALSE`
prior `none`
param
to.theta `function(x) x`
from.theta `function(x) x`

constr `FALSE`

nrow.ncol `FALSE`

augmented `FALSE`

aug.factor `1`

aug.constr

n.div.by

n.required `FALSE`

set.default.values `FALSE`

status `experimental`

pdf `scopy`

Example

Just simulate some data and estimate the parameters back.

```
## Use the spline part of scopy to estimate a spline.
## This example is rather artifical, but illustrate
## well the idea.
N <- 100
s <- 0.1
x <- 1:N
y <- 1 + sin(x * 0.25) * exp(-2*x/N) + rnorm(N, sd = s)
m <- 15
r <- inla(y ~ -1 +
              ##
              ## this model will just define the overall mean level, but
              ## with one value for each i. We need this as as can then
              ## scale this one with the spline
              ##
              f(idx,
                model = "rw1",
                scale.model = TRUE,
                constr = FALSE,
                values = 1:N,
                hyper = list(prec = list(initial = 20, fixed = TRUE))) +
              ##
              ## the 'overall level' is scaled by a spline
              ##
              f(idx.scopy,
                scopy = "idx",
                control.scopy = list(covariate = x,
                                     n = m,
                                     mean = 1,
                                     prec.mean = 1,
                                     prec.betas = 10,
                                     model = "rw2")),
          ##
          data = list(idx = rep(NA, N),
                      idx.scopy = 1:N,
                      x = x,
                      m = m),
          ##
          control.family = list(hyper = list(
                                 prec =  list(
                                     initial = log(1/s^2),
                                     fixed = TRUE))),
          ##
          control.compute = list(config = TRUE),
          verbose = TRUE)
plot(x, y, pch = 19)
## note that the locations are not stored in the results, hence we can set them here. This is
## just for the ease of the output, the results are unchanged. 1. we can just rescale
beta <- inla.summary.scopy(r, "idx.scopy", range = c(1, N))
s <- mean(r$summary.random$idx$mean)
lines(beta$x, s * beta$mean, lwd = 3, col = "blue")
lines(beta$x, s * beta$mean + 2 * s * beta$sd, lwd = 1, col = "black")
lines(beta$x, s * beta$mean - 2 * s * beta$sd, lwd = 1, col = "black")
```

Notes

This model is experimental.