

# User defined integration points

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## Introduction

This short note describe a new option that allow the user to use user-defined integration points (or “design” points), instead of the default ones. The relevant integration in INLA does

$$\int f(x|\theta, y) \pi(\theta|y) d\theta = f(x|y)$$

where  $\pi(\theta|y)$  is the approximated posterior marginal for the hyperparameters, and where  $f(x|\theta, y)$  is the approximated marginal for  $x$  for that configuration. The output of this integral is the posterior marginal  $f(x|y)$ . In practice, we use a discrete set of integration points for  $\theta$ , and corresponding weights  $w$ , to get

$$f(x|y) \approx \sum_i f(x|\theta_i, y) w_i \pi(\theta_i|y)$$

for which we require  $w_i \geq 0$  and  $\sum_i w_i = 1$ . Usually, the integration is done in a *standardised scale*,

$$z = A(\theta - \gamma)$$

i.e. with respect to  $\pi(z|y)$ . Here,  $\gamma$  is the mode of  $\pi(\theta|y)$  and the matrix  $A$  is the negative square root of the approximated covariance matrix for  $\theta|y$  at the mode.

The relevant options are

```
opts = control.inla(int.strategy = "user", int.design = Design)
```

where **Design** is a matrix with the integration points and the integration weights. The  $j$ th row of **Design** consists of the values  $\theta_j = (\theta_{1j}, \dots, \theta_{mj})$ , and the integration weight for this configuration,  $w_j$ . The values are in the  $\theta$ -scale, meaning that you have to know exactly what you are doing, including knowing the ordering of the hyperparameters.

Another version, is to define the points in the standardised scale  $z$ . To do this, use

```
opts = control.inla(int.strategy = "user.std", int.design = Design)
```

instead. The meaning of **Design** is unchanged, except that these can be given in standardised coordinates. This version is more relevant if you want to implement a generic new integration design instead of the ones already provided.

## First example

In this artificial example, we want to compute the change of the marginal variance of one component,  $x_1$ , of a hidden AR(1) process, with respect to lag one correlation  $\rho$ . So we want to compute

$$\frac{\partial \text{SD}(x_1|y)}{\partial \rho}$$

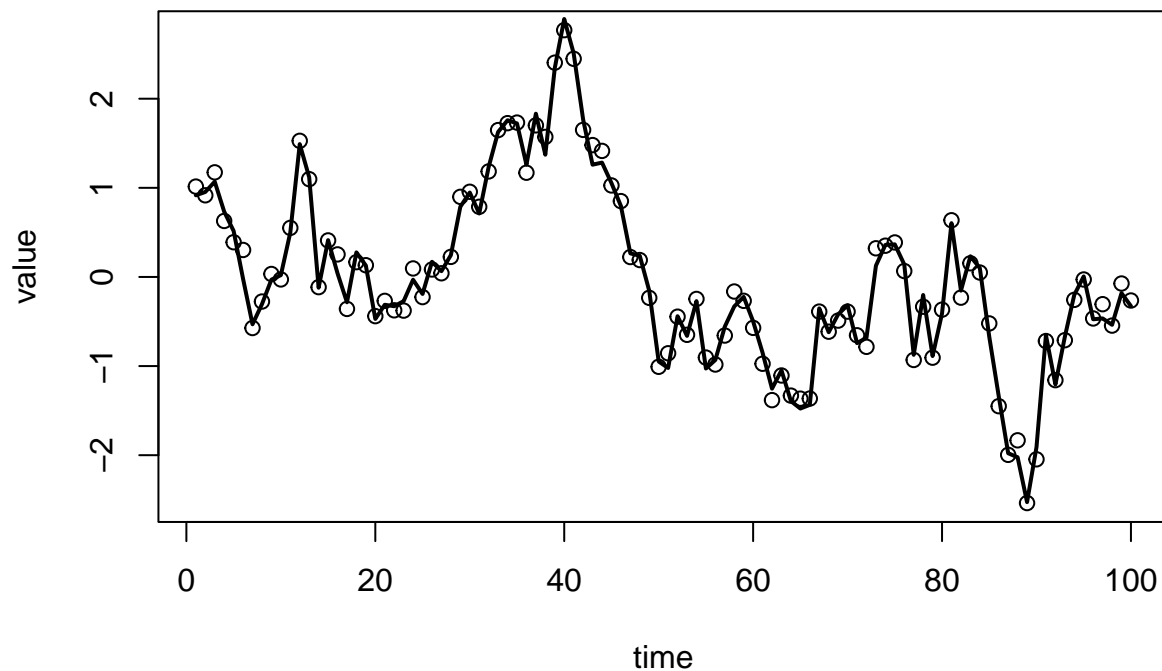
for a fixed value of  $\rho = \rho_0$ . We have to compute a numerical approximation, using finite difference. While doing this, it is a good idea to keep the design fixed, to avoid introducing an error for changing that part as well.

Let us first setup the experiment

```
n = 100
rho = 0.9
x = scale(arima.sim(n, model = list(ar = rho)))
y = x + rnorm(n, sd = 0.1)
```

this gives the following

```
plot(y, xlab = "time", ylab = "value")
lines(x, lwd=2)
```



To compute the derivative, we do

```
rho.0 = rho
to.theta = inla.models()$latent$ar1$hyper$theta2$to.theta
rho.0.internal = to.theta(rho.0)

r = inla(y ~ -1 + f(time, model="ar1",
  hyper = list(
    theta1 = list(prior = "loggamma",
      param = c(1,1)),
    theta2 = list(initial = rho.0.internal,
      fixed=TRUE))),
  control.inla = list(int.strategy = "grid"),
```

```
data = data.frame(y, time = 1:n))
```

```
sd.0 = r$summary.random$time[1,"sd"]
print(sd.0)
```

```
## [1] 0.01381139
```

The ordering of the hyperparameters are as follows,

```
nm = names(r$joint.hyper)
nm = nm[-length(nm)]
print(nm)
```

```
## [1] "Log precision for the Gaussian observations"
## [2] "Log precision for time"
## [3] "Log posterior density"
```

which may sometimes be useful to know about.

Anyway, we will now change  $\rho$  a little, while we keep the same integration points,

```
Design = as.matrix(cbind(r$joint.hyper[, seq_along(nm)], 1))
head(Design)
```

```
##      Log precision for the Gaussian observations Log precision for time
## [1,]                                9.860095                -0.16733931
## [2,]                                8.979033                -0.17108829
## [3,]                                10.741158                -0.16359032
## [4,]                                9.860541                -0.27216755
## [5,]                                9.859649                -0.06251107
## [6,]                                8.979479                -0.27591653
##      Log posterior density 1
## [1,]                -72.03461 1
## [2,]                -72.35434 1
## [3,]                -72.50780 1
## [4,]                -72.30674 1
## [5,]                -72.32276 1
## [6,]                -72.64800 1
```

where the last column is the (un-normalised) integration weights. Design has dimension 70, 4. We call `inla()` again reusing the previous found mode

```
h.rho = 0.01
rho.1.internal = to.theta(rho.0 + h.rho)
rr = inla(y ~ -1 + f(time, model="ar1",
  hyper = list(
    theta1 = list(prior = "loggamma",
      param = c(1,1)),
    theta2 = list(initial = rho.1.internal,
      fixed=TRUE))),
  control.mode = list(result = r, restart=FALSE),
  data = data.frame(y, time = 1:n),
  control.inla = list(
    int.strategy = "user",
    int.design = Design))
sd.1 = rr$summary.random$time[1,"sd"]
print(sd.1)
```

```
## [1] 0.01270946
```

and then our estimate of the derivative is

```
deriv.1 = (sd.1 - sd.0) / h.rho
print(deriv.1)
```

```
## [1] -0.1101938
```

PS: In the logfile of the `inla()`-call, the configurations are shown in the  $z$  scale even for `int.strategy="user"`.

## Second example

There is also another (experimental) option, that is

```
control.inla = list(int.strategy = "user.expert")
```

for which the weights **includes**  $\pi(\theta_i|y)$ . The following example show how to use it, fitting the same model in three different ways.

```
n = 50
x = rnorm(n)
y = 1 + x + rnorm(n, sd = 0.2)
param = c(1, 0.04)
dz = 0.1
r.std = inla(y ~ 1 + x, data = data.frame(y, x),
             control.inla = list(int.strategy = "grid",
                                 dz = dz,
                                 diff.logdens = 8),
             control.family = list(
               hyper = list(
                 prec = list(
                   prior = "loggamma",
                   param = param))))

s = r.std$internal.summary.hyperpar[1,"sd"]
m = r.std$internal.summary.hyperpar[1,"mean"]
theta = m + s*seq(-4, 4, by = dz)
weight = dnorm(theta, mean = m, sd = s)

r = rep(list(list()), length(theta))
for(k in seq_along(r)) {
  r[[k]] = inla(y ~ 1 + x,
               control.family = list(
                 hyper = list(
                   prec = list(
                     initial = theta[k],
                     fixed=TRUE))),
               data = data.frame(y, x))
}
r.merge = inla.merge(r, prob = weight)
```

```
## Warning in inla.merge(r, prob = weight): This function is experimental.
```

```
## Warning in inla.merge(r, prob = weight): Merging 'cpo' and 'pit'-results are/can
## be, approximate only
```

```

r.design = inla(y ~ 1 + x,
               data = data.frame(y, x),
               control.family = list(
                 hyper = list(
                   prec = list(
                     ## the prior here does not really matter, as we will override
                     ## it with the user.expert in any case.
                     prior = "pc.prec",
                     param = c(1, 0.01))),
                 control.inla = list(int.strategy = "user.expert",
                                     int.design = cbind(theta, weight)))
for(k in 1:2) {
  plot(inla.smarginal(r.std$marginals.fixed[[k]]),
       lwd = 2, lty = 1, type = "l",
       xlim = inla.qmarginal(c(0.0001, 0.9999), r.std$marginals.fixed[[k]]))
  lines(inla.smarginal(r.design$marginals.fixed[[k]]),
        lwd = 2, col = "blue", lty = 1)
  lines(inla.smarginal(r.merge$marginals.fixed[[k]]),
        lwd = 2, col = "yellow", lty = 1)
}

```

