Approximating joint marginals in R-INLA

Cristian Chiuchiolo and Haavard Rue

KAUST, Aug 2020

Introduction

The primary use of R-INLA is to approximate univariate marginals of the latent field, so we can compute their marginal summaries and densities. In applications, we sometimes need more than this, as we are interested also in statistics which involve more several of the components of the latent field, and/or, a joint posterior approximation to a subset of the latent field.

The way around this issue, have earlier resolved to stochastic simulation, using the function inla.posterior.sample. This function samples from joint approximation to the full posterior which INLA construct, but do so for the whole latent field. Using these samples, we can compute the density of the relevant statistics and/or use standard methods to represent a joint marginal.

This vignette introduces a new tool which computes a deterministic approximation to the joint marginal for a subset of the latent field using R-INLA. This approximation is explicitly available, and constructed using skew-normal marginals and a Gaussian copula, hence restricted to a joint approximation of a modest dimension.

The key specification is using an argument selection in the inla()-call, which defines the subset, and then the joint marginal approximation is made available in result\$selection.

Theory reference

For any Latent Gaussian Model with data y and set of parameters (x, θ) with x being the latent field and θ the hyperparameters, the resulting joint posterior approximation is stated as

$$\tilde{\pi}(\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y}) \propto \sum_{k} \tilde{\pi}_{G}(\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y}) \tilde{\pi}(\boldsymbol{\theta}_{k}|\boldsymbol{y}) \Delta_{k}$$
 (1)

where $\tilde{\pi}_G(x|\theta,y)$ is the Gaussian approximation. This expression recalls a Gaussian mixture distribution with weights $\tilde{\pi}(\theta_k|y)\Delta_k$ obtained in the grid exploration of the hyperparameter posterior marginals. For more insights, we suggest checking sources like Rue, Martino, and Chopin (2009), Martins et al. (2013), Blangiardo et al. (2013), or the recent review by Martino and Riebler (2019). The Gaussian approximation used in (1) is both mean and skewness corrected since it exploits Skew-Normal marginals of the latent field into a Gaussian copula structure (see Ferkingstad and Rue (2015) for details). These corrections are available in inla.posterior.sample as

inla.posterior.sample(..., skew.corr = TRUE)

First example

We will illustrate this new feature using a simple example.

```
n = 100
x = rnorm(n, mean = 1, sd = 0.3)
xx = rnorm(n, mean = -2, sd = 0.3)
```

```
y = rpois(n, lambda = exp(x + xx))
r = inla(y \sim 1 + x + xx,
         data = data.frame(y, x, xx),
         family = "poisson")
```

Let us compute the joint marginal for the effect of x, xx and the intercept. This is specified using the argument selection, which is a named list of indices to select. Names are those given the formula, plus standard names like (Intercept), Predictor and APredictor. So that

```
selection = list(Predictor = 3:4, x = 1, xx = 1)
```

say that we want the joint marginal for the 3^{rd} and 4^{th} element of Predictor and the first element of x and xx. (Well, x and xx only have one element, so then there is not else we can do in this case.)

If we pass selection, then we have to rerun inla() as

```
rs = inla(y \sim 1 + x + xx,
          data = data.frame(y, x, xx),
          family = "poisson",
          control.predictor = list(compute = TRUE),
          selection = selection)
```

and we obtain

\$marginal.sn.par\$xi

```
#summary(rs$selection)
print(rs$selection)
```

```
$names
[1] "Predictor:3" "Predictor:4" "x:1"
                                              "xx:1"
$mean
[1] -0.7122284 -1.5151395 1.6615015 1.9552196
$cov.matrix
                        [,2]
                                    [,3]
            [,1]
[1,] 0.08282597 0.02406932 0.13371964 -0.03139111
[2,] 0.02406932 0.05741459 -0.06885383 -0.08589099
[3,] 0.13371964 -0.06885383 0.44694642 0.10015724
[4,] -0.03139111 -0.08589099 0.10015724 0.32337257
$skewness
[1] -0.24583152 -0.26877320 -0.00106979 -0.02536701
$marginal.sn.par
```

```
$marginal.sn.par$omega
[1] 0.3740984 0.3153399 0.6746575 0.6102768
```

[1] -0.4732228 -1.3101404 1.7521456 2.1767263

```
$marginal.sn.par$alpha
[1] -1.3366865 -1.4052794 -0.1708292 -0.5108181
```

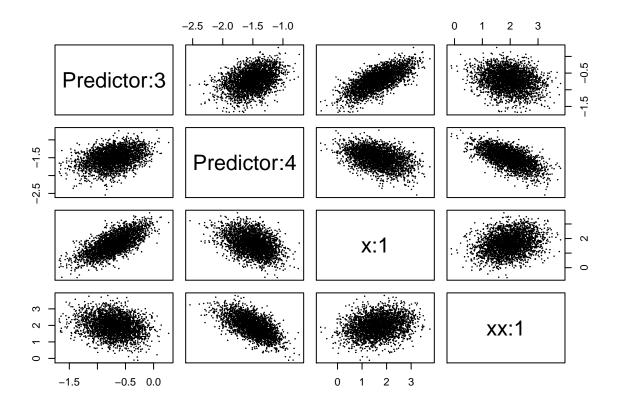
The Gaussian copula is given by the mean and the cov.matrix objects, while the Skew-Normal marginals are given implicitly using the marginal mean and variance in the Gaussian copula and the listed skewness. Moreover the respective Skew-Normal mapping parameters for the marginals (ξ, ω, α) are provided in the object 'marginal.sn.par'. The names are given as a separate entry instead of naming each individual result, to

save some storage.

There are utility functions to sample and evaluate samples from this joint marginal, similar to inla.posterior.sample and inla.posterior.sample.eval.

```
ns = 10000
xs = inla.rjmarginal(ns, rs) ## or rs$selection
str(xs)

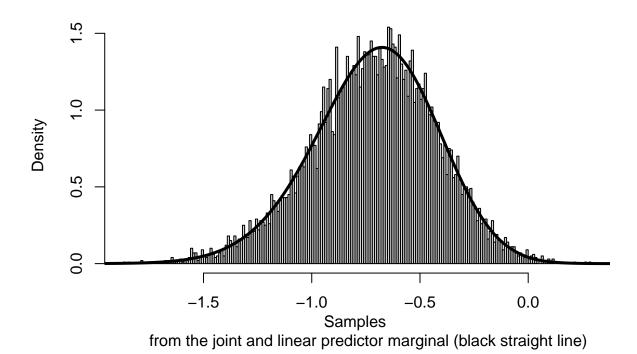
List of 2
$ samples : num [1:4, 1:10000] -0.559 -1.27 1.469 1.367 -0.317 ...
..- attr(*, "dimnames")=List of 2
...$ : chr [1:4] "Predictor:3" "Predictor:4" "x:1" "xx:1"
...$ : chr [1:10000] "sample:1" "sample:2" "sample:3" "sample:4" ...
$ log.density: Named num [1:10000] 4.028 0.836 4.19 3.023 2.064 ...
..- attr(*, "names")= chr [1:10000] "sample:1" "sample:2" "sample:3" "sample:4" ...
whose output is a matrix where each row contains the samples for the variable in each column
pairs(t(xs$samples[, 1:3000]), cex = 0.1)
```



We can compare the approximation of Predictor: 3 to the one computed by the inla() call,

```
hist(xs$samples["Predictor:3",], n = 300, prob = TRUE,
    main = 'Histogram of Predictor:3', xlab = 'Samples
    from the joint and linear predictor marginal (black straight line)')
lines(inla.smarginal(rs$marginals.linear.predictor[[3]]), lwd = 3)
```

Histogram of Predictor:3



These marginals are not *exactly* the same (as they are different approximations), but should be very similar.

Deterministic Joint approximation

As a conclusion to this vignette we show an additional joint posterior related tool. The following INLA function computes a deterministic approximation for the joint posterior sampler and must be considered experimental. The function is strictly related to the selection type INLA setting. Deterministic posterior marginals for the previous example can be obtained as follows

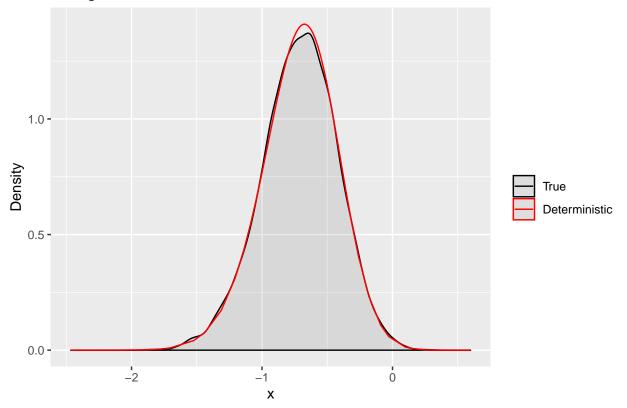
```
dxs = inla.1djmarginal(jmarginal = rs$selection)
str(dxs)
```

```
List of 4
 $ Predictor:3: num [1:63, 1:2] -2.46 -2.3 -2.13 -1.93 -1.78 ...
  ..- attr(*, "dimnames")=List of 2
  .. ..$ : NULL
  ....$ : chr [1:2] "x" "y"
 $ Predictor:4: num [1:63, 1:2] -2.99 -2.85 -2.7 -2.54 -2.41 ...
  ..- attr(*, "dimnames")=List of 2
  .. ..$ : NULL
  ....$ : chr [1:2] "x" "y"
              : num [1:63, 1:2] -1.818 -1.519 -1.192 -0.826 -0.54 ...
  ..- attr(*, "dimnames")=List of 2
  .. ..$ : NULL
  ....$ : chr [1:2] "x" "y"
              : num [1:63, 1:2] -1.0736 -0.8079 -0.5177 -0.1949 0.0574 ...
 $ xx:1
  ..- attr(*, "dimnames")=List of 2
```

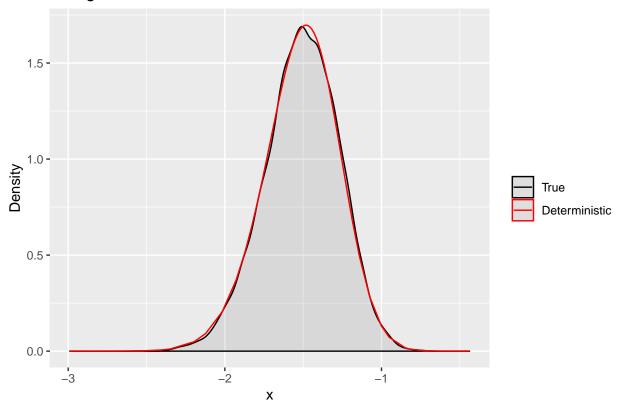
```
....$ : NULL
....$ : chr [1:2] "x" "y"
```

The output is a list with all computed marginals and a matrix summary output in INLA style. Marginal can be accessed and plotted by using the respective names in the selection

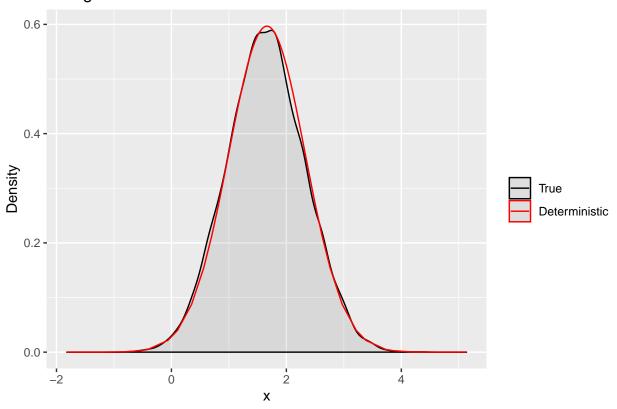
"Marginal Results for Predictor:3"



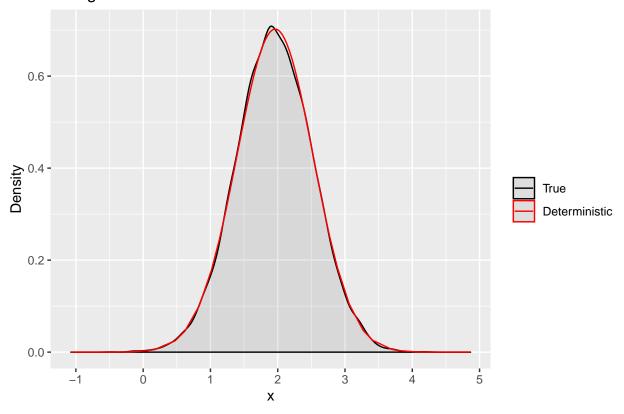
"Marginal Results for Predictor:4"



"Marginal Results for x:1"



"Marginal Results for xx:1"



Here we compare the deterministic marginals with their sampling version. They are quite accurate and can provide more informations. Indeed, a complete summary based on these deterministic results is achievable with a personalized summary function

```
summary(rs$selection)
```

```
[1] "Joint marginal is computed for: "
                              sd 0.025quant
                                              0.5quant 0.975quant
                  mean
                                                                         mode
Predictor:3 -0.7122284 0.2877950 -1.3116547 -0.6999178 -0.1811191 -0.6746855
Predictor:4 -1.5151395 0.2396134 -2.0169092 -1.5038748 -1.0756438 -1.4807078
             1.6615015 0.6685405 0.3508431
x:1
                                             1.6616207
                                                        2.9714825
                                                                    1.6618524
             1.9552196 0.5686586
                                 0.8335949
                                             1.9576304
xx:1
                                                        3.0631556
                                                                   1.9624591
```

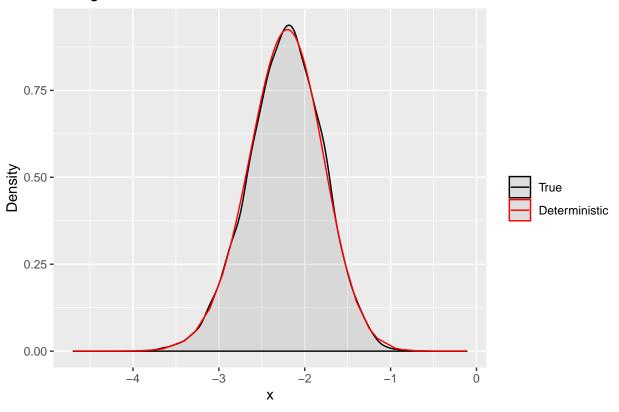
where posterior estimates and quantiles are computed for all the selected marginals. Along the same line, we can easily compute multiple deterministic linear combinations through the function inla.tjmarginaland a matrix object A with the respective indexes

```
A = matrix(c(1,1,0,0,0,0,1,1), nrow = 2, ncol = 4, byrow = T)
rownames(A) <- c("Test1", "Test2")
A</pre>
```

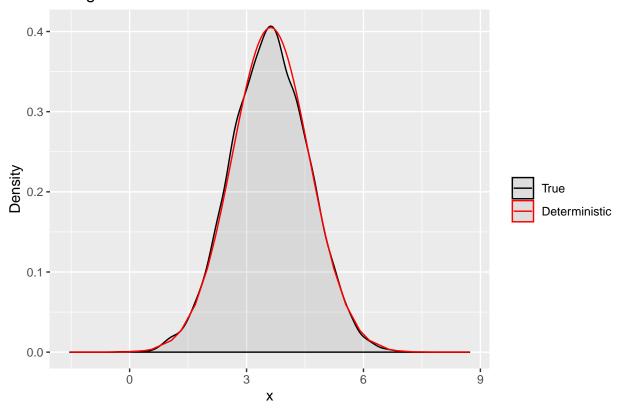
We define two linear combinations: Predictor: 3+Predictor: 4 and x:1+xx:1 respectively. Then we can use the cited function which has the same class of selection object

```
m = inla.tjmarginal(jmarginal = rs, A = A)
class(m)
$names
[1] "Test1" "Test2"
$mean
           [,1]
Test1 -2.227368
Test2 3.616721
$cov.matrix
            Test1
                        Test2
Test1 0.18837920 -0.05241628
Test2 -0.05241628 0.97063347
$skewness
[1] -0.116894186 -0.005212263
$marginal.sn.par
$marginal.sn.par$xi
    Test1
              Test2
-1.946031 3.843176
$marginal.sn.par$omega
    Test1
              Test2
0.5172325 1.0108983
$marginal.sn.par$alpha
[1] -0.9317813 -0.2925254
[1] "inla.jmarginal"
dxs.lin = inla.1djmarginal(jmarginal = m)
str(dxs.lin)
fun1 <- function(...) {Predictor[1]+Predictor[2]}</pre>
fun2 <- function(...) {x+xx}</pre>
xs.lin1 = inla.rjmarginal.eval(fun1, xs)
xs.lin2 = inla.rjmarginal.eval(fun2, xs)
ggplot(data = data.frame(y = xs.lin1[1, ]), aes(y, after_stat(density), colour = "True")) +
  stat_density(alpha = .1) +
  geom_line(data = as.data.frame(dxs.lin$Test1), aes(x = x, y = y, colour = "Deterministic"))+
  labs(title= '"Marginal Results for Lin1"', x='x', y='Density') +
  scale_colour_manual("",
                      breaks = c("True", "Deterministic"),
                      values = c("black", "red"))
```

"Marginal Results for Lin1"



"Marginal Results for Lin2"



```
List of 2
$ Test1: num [1:63, 1:2] -4.69 -4.48 -4.23 -3.96 -3.75 ...
..- attr(*, "dimnames")=List of 2
....$: NULL
....$: chr [1:2] "x" "y"
$ Test2: num [1:63, 1:2] -1.5313 -1.0874 -0.6016 -0.0594 0.3657 ...
..- attr(*, "dimnames")=List of 2
....$: NULL
....$: chr [1:2] "x" "y"
```

and accomplish the job with summaries

summary(m)

```
[1] "Joint marginal is computed for: "

mean sd 0.025quant 0.5quant 0.975quant mode

Test1 -2.227368 0.4340267 -3.103378 -2.218768 -1.399892 -2.201422

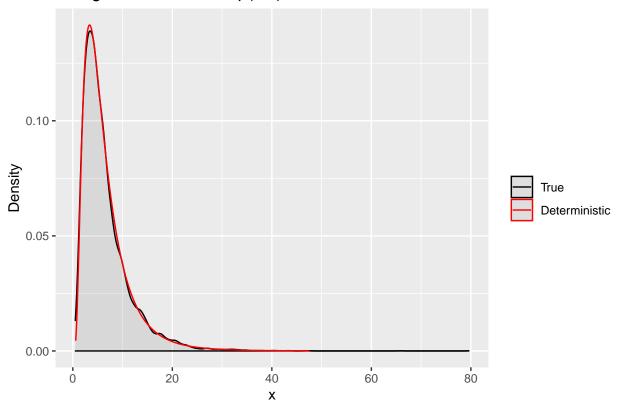
Test2 3.616721 0.9852073 1.683266 3.617577 5.545311 3.619280
```

Transformations of the marginal terms or linear combinations are possible as well. We just need to use inla.tmarginal as follows

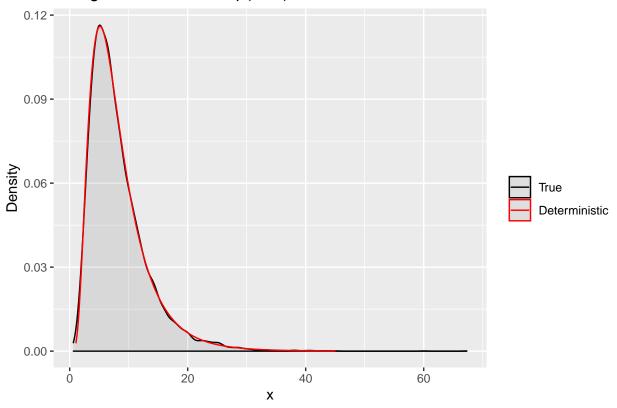
```
fun.exp <- function(x) exp(x)

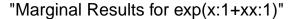
fun5 <- function(...) {exp(x)}
fun6 <- function(...) {exp(xxx)}
fun7 <- function(...) {exp(x+xx)}</pre>
```

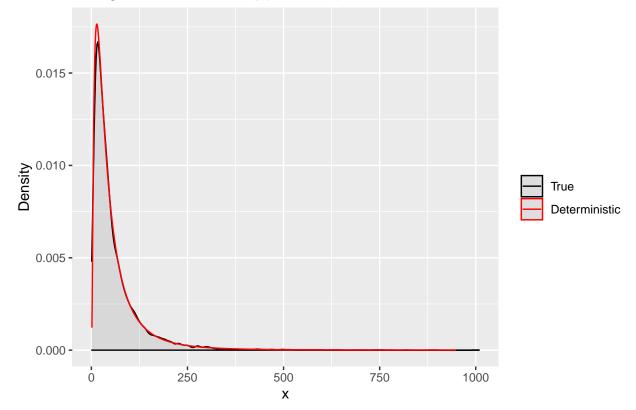
"Marginal Results for exp(x:1)"



"Marginal Results for exp(xx:1)"







Summaries for all marginal transformations can be obtained through inla.zmarginal

```
expx = inla.zmarginal(marginal = tdx, silent = TRUE)
expxx = inla.zmarginal(marginal = tdxx, silent = TRUE)
expx.lin = inla.zmarginal(marginal = tdx.lin, silent = TRUE)

exp.summaries = rbind(expx, expxx, expx.lin)
exp.summaries
```

```
    mean
    sd
    quant0.025
    quant0.25
    quant0.5
    quant0.75
    quant0.75
    quant0.975

    expx
    6.562708
    4.800713
    1.416441
    3.345754
    5.256102
    8.252395
    19.39268

    expxx
    8.28001
    4.985093
    2.301952
    4.813491
    7.071574
    10.36367
    21.2768

    expx.lin
    59.82771
    71.62435
    5.179063
    18.93141
    37.00413
    72.05407
    253.5351
```

References

Blangiardo, M., M. Cameletti, G. Baio, and H. Rue. 2013. "Spatial and Spatio-Temporal Models with R-INLA." Spatial and Spatio-Temporal Epidemiology 3 (December): 39–55.

Ferkingstad, E., and H. Rue. 2015. "Improving the INLA Approach for Approximate Bayesian Inference for Latent Gaussian Models." *Electronic Journal of Statistics* 9: 2706–31. https://doi.org/10.1214/15-EJS1092.

Martino, Sara, and Andrea Riebler. 2019. "Integrated Nested Laplace Approximations (Inla)." http://arxiv.org/abs/1907.01248.

Martins, T. G., D. Simpson, F. Lindgren, and H. Rue. 2013. "Bayesian Computing with INLA: New Features." Csda 67: 68–83.

Rue, Håvard, Sara Martino, and Nicolas Chopin. 2009. "Approximate Bayesian Inference for Latent Gaussian Models by Using Integrated Nested Laplace Approximations." Journal of the Royal Statistical Society: Series B (Statistical Methodology) 71 (2): 319–92.