

# BiiPS

a software for Bayesian inference with interacting  $\ensuremath{\textbf{P}}\xspace{article}$  Systems

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Summary







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#### Summary





3 Example in financial econometry



$$p(X|Y) = \frac{p(X,Y)}{p(Y)}$$

- Many people use MCMC methods with BUGS software
  - Provides a, so-called, BUGS language for describing a graphical model
  - Expert system drives MCMC methods (Gibbs, Slice, Metropolis, ...) in a 'black-box' fashion
  - Very popular among practitioners, applying MCMC methods to a wide range of applications
- Having such a 'black-box' software (generic and easy to use) for SMC methods would be great
- The *BiiPS* project have been trying to bridge this gap for 3 years

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The graph displays a factorization of the joint distribution:

 $p(x_{1:3}, y_{1:2})$ 

Figure : Directed acyclic graph

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- Stochastic relations
- Deterministic relations



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Y  $\sim$  dnorm(mu, tau)



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Y ~ dnorm(mu, tau) tau ~ dgamma(0.01, 0.01)

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Y ~ dnorm(mu, tau) tau ~ dgamma(0.01, 0.01) mu <- alpha + beta \* x



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#### Summary





3 Example in financial econometry



#### BiiPS software

- Core developped in C++ (>30K lines)
- $\bullet$  BUGS language compiler adapted from JAGS  $\bigodot$  M. Plummer
- Multi-platform: Linux, Windows, Mac OSX
- Open-source GPL license
- RBiips interface for R
- MatBiips interface for Matlab (ongoing development)



Figure : *BiiPS*: input/output diagram





Figure : Directed acyclic graph

- Sample  $x_1^{(i)} \sim p(x_1)$ , and set weights  $w_1^{(i)} = 1/N$
- Sample  $x_2^{(i)} \sim p(x_2|x_1^{(i)})$
- Set weights  $w_2^{(i)} = w_1^{(i)} p(y_1 | x_2^{(i)})$  and resample  $\{x_{1:2}^{(i)}, w_2^{(i)}\}$
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More generally:

- Sample nodes in a topological order
- One iteration of SMC corresponds to sampling one unobserved parameter: p(X<sub>k</sub>|parents(X<sub>k</sub>))
- Weight particles with likelihood associated to their observed children p(Y<sub>k</sub>|parents(Y<sub>k</sub>))

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#### Choice of the importance distribution

- Sampling from the prior  $p(X_k|parents(X_k))$  may lead to bad resampling and degeneracy problems
- It is better to sample from  $p(X_k | parents(X_k), children(X_k))$  or any approximation

Prior to running the SMC, *BiiPS* assigns an appropriate importance sampling method (node sampler) to each unobserved parameter

- Finite discrete sampler
- Onjugate sampler (only normal prior implemented yet)
- Prior sampler (default)



#### When does it work?



Figure : State-space model / HMM

- Ok with state-space models (HMM), switching state-space models, etc.
- More generally: ok when the unkown parameters are controlled by a dynamic system
- But this method will not suit all graphical models

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Figure : Switching state-space model

- Ok with state-space models (HMM), switching state-space models, etc.
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- But this method will not suit all graphical models

#### How to deal with fixed parameters?



#### Particle Marginal Metropolis-Hastings [?]

MCMC algorithm using SMC at each iteration. At iteration k:

- Propose a  $\theta^*$
- Run an SMC algorithm conditionally on  $\theta^*$
- Accept or reject θ<sup>\*</sup> with acceptance rate depending on the estimate of the conditional marginal likelihood p̂<sub>θ\*</sub>(Y<sub>1:T</sub>)

#### Summary







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Consider infering the underlying volatility  $X_{1:T}$  from observed incremental price or rate  $Y_{1:T}$ 

$$\begin{array}{l} X_1 \sim \mathcal{N}(0,\sigma^2) \\ X_t | X_{t-1} \sim \mathcal{N}(\alpha x_{t-1}, \frac{\sigma^2}{1-\alpha^2}) \quad t > 1 \\ Y_t | X_t \sim \mathcal{N}(0, \beta^2 \exp(x_t)) \quad t > 1 \end{array}$$









#### BUGS language "volatility.bug"

```
model
{
    x[1] ~ dnorm(0, 1 / sigma^2)
    prec.x <- (1-alpha^2) / sigma^2
    for (t in 2:t.max)
    {
        x[t] ~ dnorm(alpha * x[t-1], prec.x)
        prec.y[t] <- 1 / (beta^2 * exp(x[t]))
        y[t] ~ dnorm(0, prec.y[t])
    }
}</pre>
```

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```
# Compile the model and load the data
model <- biips.model("volatility.bug", data)</pre>
```

```
# Run SMC algorithm
out.smc <- smc.samples(model, "x", n.part=1000)</pre>
```

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Figure : Summary statistics

# # Kernel density estimates plot(density(out.smc\$x, adjust=2))



Figure : Kernel density estimates



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#### Estimation of the fixed parameter $\boldsymbol{\alpha}$

#### BUGS language "volatility\_param.bug"

```
model {
    x[1] ~ dnorm(0, 1 / sigma^2)
    prec.x <- (1-alpha^2) / sigma^2
    for (t in 2:t.max) {
        f[t] <- alpha * x[t-1]
        x[t] ~ dnorm(f[t], prec.x)
        prec.y[t] <- 1 / (beta^2 * exp(x[t]))
        y[t] ~ dnorm(0, prec.y[t]) }
    alpha ~ dunif(0, 0.99) }
</pre>
```



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Estimation of the fixed parameter  $\boldsymbol{\alpha}$ 

```
# Define data and compile the model
data <- list(t.max=100, sigma=1.0,</pre>
              beta=0.5, v=v)
model <- biips.model("volatility param.bug", data)</pre>
# Burn in PMMH algorithm
update.pmmh(model, "alpha", n.iter=1000, n.part=100)
# Generate PMMH samples
out.pmmh <- pmmh.samples(model, "alpha", n.iter=10000,</pre>
                           n.part=100)
# Summary statistics
alpha.mean <- mean(out.pmmh$alpha)</pre>
```

```
alpha.var <- var(out.pmmh$alpha)</pre>
```

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#### Estimation of the fixed parameter $\boldsymbol{\alpha}$

# # PMMH trace plot and histogram plot(out.pmmh\$alpha) hist(out.pmmh\$alpha)

Histogram of 10000 BIPS PMMH samples

Trace of 10000 BijPS PMMH samples

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#### Figure : $\alpha$ PMMH samples: trace plot and histogram

0.85

value

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0.80

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0.90

0.95

## THANK YOU

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