

# INTRODUCING THE BAYESIAN ANALYSIS TOOLKIT AND PYPMC

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Excellence cluster universe

Belle II workshop, Karlsruhe, Feb 2015

BAT men: A. Caldwell, D. Greenwald (**Belle I/II**), D. Kollár, K. Kröninger  
pypmc: S. Jahn



F. Beaujean

Bayes: basics,  
numerics

C. Bobeth

global fit of rare B decays:  
motivation, results

D. van Dyk

computing observables  
with EOS

- ① Given the data from Belle, BaBar, LHCb, (... Belle II), what are the likely values of Wilson coefficients, or CKM phases, or ... ?
- ② If there is a deviation from the SM, which NP model is preferred?

## PARAMETER INFERENCE

$$P(\theta|D, M) = \frac{P(D|\theta, M)P_0(\theta|M)}{P(D|M)}$$

posterior  $\propto$  likelihood  $\times$  prior

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## MODEL COMPARISON

$$\text{evidence } P(D|M) = \int d\theta P(D|\theta, M)P_0(\theta|M)$$

$$\frac{P(M_1|D)}{P(M_2|D)} = \frac{P(D|M_1)}{P(D|M_2)} \times \frac{P(M_1)}{P(M_2)}$$

posterior odds = Bayes factor  $\times$  prior odds

## PARAMETERS

- $\theta$  = Wilson coefficients, CKM parameters, theory uncertainties, detector effects . . .
- $\theta = (\mu, \nu)$  parameters of interest + nuisance

# APPLYING BAYES' THEOREM

## PARAMETERS

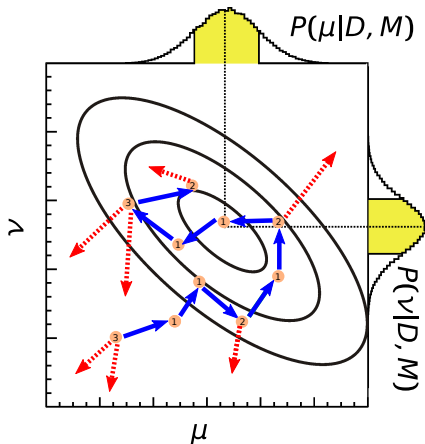
- $\theta$  = Wilson coefficients, CKM parameters, theory uncertainties, detector effects ...
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## INTEGRATION

- marginalization  $P(\mu|D, M) = \int d\nu P(\mu, \nu|D, M)$
- evidence  $P(D|M) = \int d\theta P(D|\theta, M)P_0(\theta|M)$
- curse of dimensionality

⇒ need samples from posterior

# MARKOV CHAIN MONTE CARLO



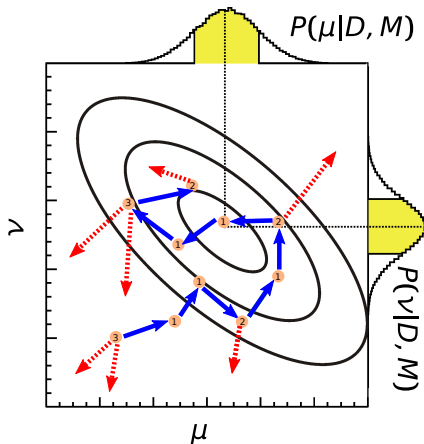
## METROPOLIS HASTINGS ALGORITHM

one sample for each step

- 1 propose move
- 2 **accept** or **stay**



# MARKOV CHAIN MONTE CARLO

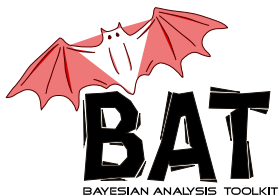


## METROPOLIS HASTINGS ALGORITHM

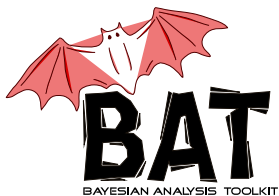
one sample for each step

- 1 propose move
- 2 **accept** or **stay**

- sample near mode  $\Rightarrow$  seed for optimization
- uncertainty propagation  
 $f(\mu, \nu) \rightarrow P(f|D, M)$



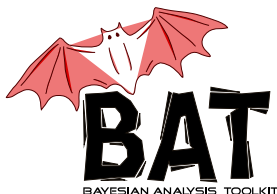
- home page <http://mpp.mpg.de/bat>
- fork me on <https://github.com/bat/bat>
- first release in 2008, latest **0.9.4.1** in Jan 2015



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## MOTIVATION

- reinventing the wheel time waster, error prone
- create C++ toolkit to supply algorithms/models, so user can focus on the problem



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## FEATURES

- implemented: MCMC (multithreaded), simulated annealing ...
- depends on ROOT: I/O, plots, optimization (Minuit) ...
- optional: roostats, CUBA (integration)
- docs, tutorials, examples ... on web page

$$P(\theta|D, M) \propto P(D|\theta, M)P_0(\theta|M)$$

## USER DEFINED

- create model
- read data

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## DEFINE MYMODEL : BCMODEL

- AddParameter("mu", 0, 1)
- LogLikelihood()
- LogAPrioriProbability()

## READ DATA

- text file
- ROOT tree
- anything in C++

$$P(\theta|D, M) \propto P(D|\theta, M)P_0(\theta|M)$$

## USER DEFINED

- create model
- read data

## COMMON TOOLS

- `Normalize()`
- `FindMode()`
- `MarginalizeAll()`
- `PrintAllMarginalized()`
- `PrintKnowledgeUpdatePlots()`

## DEFINE MYMODEL : BCMODEL

- `AddParameter("mu", 0, 1)`
- `LogLikelihood()`
- `LogAPrioriProbability()`

## READ DATA

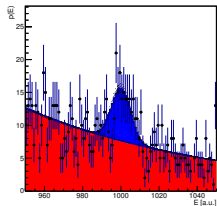
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## PREDEFINED MODEL

### template fit: signal + bkg

```
// define the model
BCMTF m("SingleChannelMTF");
m.AddChannel("channel1");
m.SetData("channel1", hist_data);
m.AddProcess("background", 200., 400.);
m.SetTemplate("channel1", "background",
             hist_background, 1.0);
m.SetPriorGauss("background", 300., 10.);
m.AddProcess("signal", 0., 200.);
m.SetTemplate("channel1", "signal", hist_signal, 1.0);
m.SetPriorConstant("signal");
```

```
// analyze model
m.MarginalizeAll();
m.PrintAllMarginalized("marg.pdf");
m.FindMode(m.GetBestFitParameters());
m.PrintStack(0, m.GetBestFitParameters(), "stack.pdf")
```



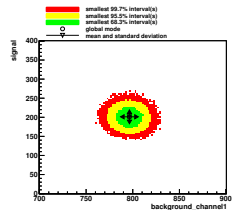
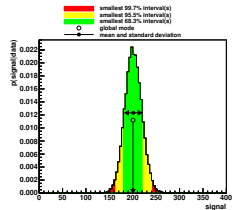
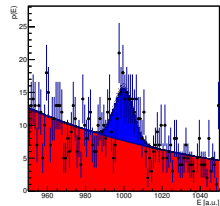


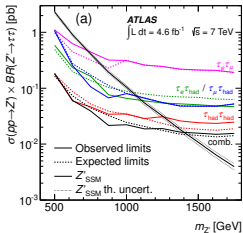
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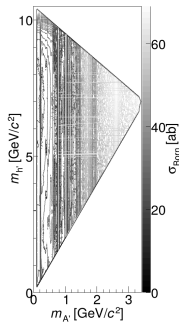
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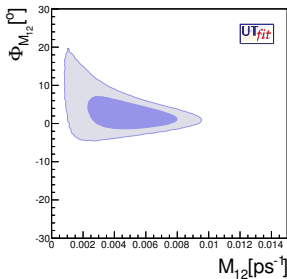




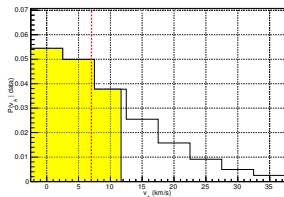
ATLAS:  $Z'$  search  
Phys. Lett. B 719 (2013)



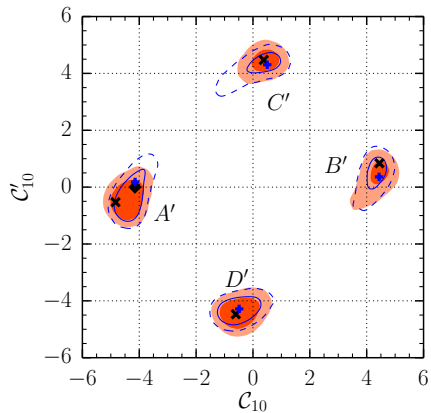
Belle: dark-photon search  
[arXiv:1502.00084](https://arxiv.org/abs/1502.00084)



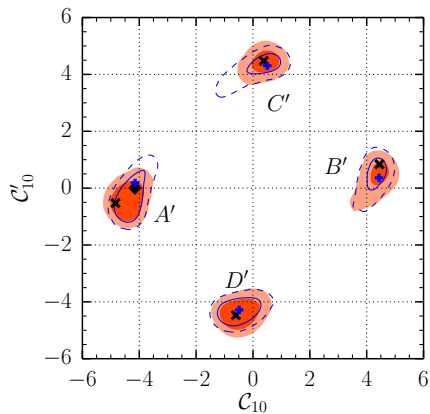
UTFIT: D meson mixing  
[arXiv:1402.1664](https://arxiv.org/abs/1402.1664)



PAMELA: cosmic-ray proton spectrum  
[arXiv:1306.1354](https://arxiv.org/abs/1306.1354)



see C. Bobeth's talk



see C. Bobeth's talk

### CHALLENGES

- high dimensional: (30-40)D
- isolated modes
- slow likelihood  $\mathcal{O}(1s)$

⇒ Metropolis Hastings not enough

## IMPORTANCE SAMPLING

$$\int P = \int \frac{P}{q} q \approx \frac{1}{N} \sum_i \frac{P(\theta_i)}{q(\theta_i)}, \theta \sim q$$

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## MIXTURE PROPOSAL

$$q(\theta) = \sum_{k=1}^K \alpha_k \mathcal{N}(\theta | \mu_k, \Sigma_k)$$

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## ADVANTAGES

- evidence + marginals
- massively parallel
- multiple modes

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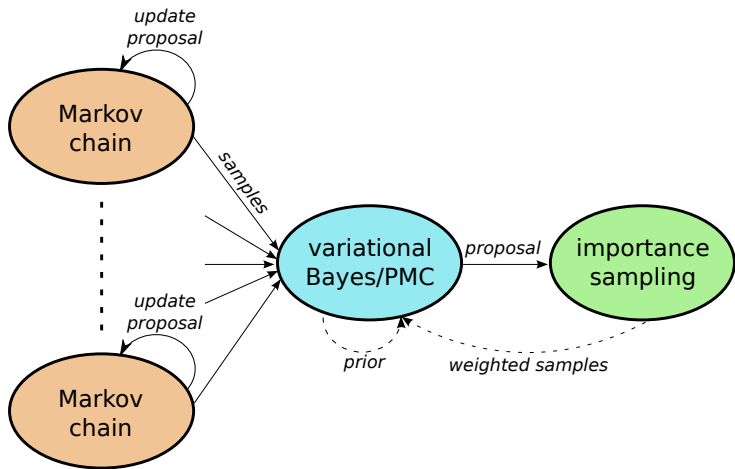
## ADVANTAGES

- evidence + marginals
- massively parallel
- multiple modes

⇒ how to adjust proposal  $q$ ?



# ALGORITHM OVERVIEW



hierarchical clustering + PMC: [arXiv:1304.7808](https://arxiv.org/abs/1304.7808)

variational Bayes: S. Jahn's master's thesis (TU Munich, Feb 2015)

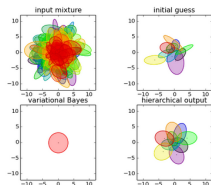
[pypmc 1.0 documentation](#) »

## pypmc

pypmc is a python package focusing on adaptive importance sampling. It can be used for integration and sampling from a user-defined target density. A typical application is Bayesian inference, where one wants to sample from the posterior to marginalize over parameters and to compute the evidence. The key idea is to create a good proposal density by adapting a mixture of Gaussian or student's t components to the target density. The package is able to efficiently integrate multimodal functions in up to about 30-40 dimensions at the level of 1% accuracy or less. For many problems, this is achieved without requiring any manual input from the user about details of the function. Importance sampling supports parallelization on multiple machines via mpi4py.

Useful tools that can be used stand-alone include:

- importance sampling (sampling & integration)
- adaptive Markov chain Monte Carlo (sampling)
- variational Bayes (clustering)



### 4.6. MCMC + variational Bayes

```
'''This example illustrates how pypmc can be used to int
non-negative function. The presented algorithm needs ver
analytical knowledge about the function.
'''
```

```
...
from __future__ import print_function
import numpy as np
import pypmc
```

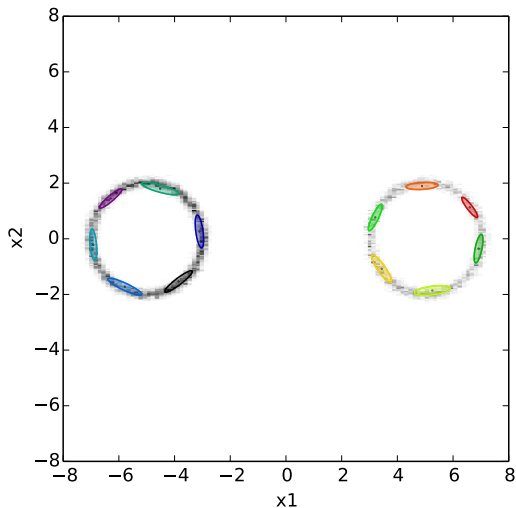
## MAIN FEATURES

- Markov chains
- importance sampling (with MP I)
- variational Bayes (GMM)
- population Monte Carlo (Gauss + Student's t)

## ORDER NOW

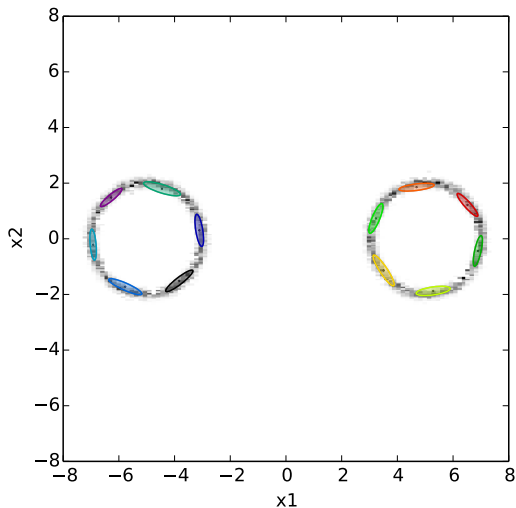
```
pip install pypmc
```

# VARIATIONAL BAYES AND IMPORTANCE SAMPLING



MCMC + variational Bayes (VB)

# VARIATIONAL BAYES AND IMPORTANCE SAMPLING



(MCMC + VB) + (IS + VB)

## IMPROVEMENTS UNDER DEVELOPMENT

- factorized priors  $P(\theta|M) = \prod_i P(\theta_i|M)$ 
  - ⇒ community extensible
- sharing samples as ROOT files (even w/o the model)
  - ⇒ uncertainty propagation, replotting
- multivariate proposal ⇒ big speed-up in high dimensions
- evidence from MCMC [arXiv:1410.7149](https://arxiv.org/abs/1410.7149)

⇒ release in summer 2015

## WISHLIST FOR THE FUTURE

- performance: threads + MPI for tough problems
- interface from C++ to python, mathematica ...
- sampling algorithms: MCMC, Hamiltonian MC, nested sampling, variational Bayes + importance sampling ...

⇒ BAT 2.0 ready for Belle II analyses

- ① Bayes: random numbers
- ② BAT well usable for small-medium scale problems
- ③ many more powerful sampling algorithms in BAT 2.0