The Expectation-Propagation Algorithm: a tutorial Day II: Recent Advances

Simon Barthelmé, Gipsa-lab, CNRS

2nd March 2016

▲□▶ ▲□▶ ▲□▶ ▲□▶ ▲□ ● ● ●

Outline

- ABC-EP: EP for likelihood-free problems
- MCMC-EP: speeding up MCMC for large datasets

▲□▶ ▲□▶ ▲□▶ ▲□▶ ▲□ ● ● ●

Average-EP and stochastic EP: simpler EP

Likelihood-free Bayesian inference

- ► A class of techniques that can be applied when
 - Likelihood evaluation is impossible or very, very slow

・ロト ・ 日 ・ ・ 日 ・ ・ 日 ・ ・ つ へ ()

Sampling from the model is comparatively easy

Likelihood-free Bayesian inference

- A class of techniques that can be applied when
 - Likelihood evaluation is impossible or very, very slow

ション ふゆ く 山 マ チャット しょうくしゃ

- Sampling from the model is comparatively easy
- Most famous incarnation: the Approximate Bayesian Computation alg. of Pritchard et al. (1999)

Approximate Bayesian Computation

There are many intractable-likelihood models in Population Genetics, where researchers are interested for example in reconstructing evolutionary trees from molecular data.

▲□▶ ▲□▶ ▲□▶ ▲□▶ ▲□ ● ● ●

Approximate Bayesian Computation

- There are many intractable-likelihood models in Population Genetics, where researchers are interested for example in reconstructing evolutionary trees from molecular data.
- Enter [?], with an algorithm now know as ABC, for Approximate Bayesian Computation, now perhaps the hottest topic in computational and applied statistics.

Approximate Bayesian Computation

- There are many intractable-likelihood models in Population Genetics, where researchers are interested for example in reconstructing evolutionary trees from molecular data.
- Enter [?], with an algorithm now know as ABC, for Approximate Bayesian Computation, now perhaps the hottest topic in computational and applied statistics.
- ► Idea brilliantly simple if one thinks of Bayesian modelling as defining a *joint* distribution $p(\mathbf{y}, \boldsymbol{\theta}) = p(\boldsymbol{\theta})p(\mathbf{y}|\boldsymbol{\theta})$ over data and parameters.

(日) (伊) (日) (日) (日) (0) (0)

1. Sample $heta \sim p(heta)$



- 1. Sample $\theta \sim p(\theta)$
- 2. Sample $\mathbf{y} \sim p(\mathbf{y}|\boldsymbol{\theta})$

▲□▶ ▲圖▶ ▲臣▶ ★臣▶ ―臣 …の�?

- 1. Sample $\theta \sim p(\theta)$
- 2. Sample $\mathbf{y} \sim p(\mathbf{y}|\boldsymbol{\theta})$
- 3. Accept $\boldsymbol{\theta}$ iff $||\mathbf{y} \mathbf{y}^{\star}|| < \epsilon$

◆□▶ ◆□▶ ◆臣▶ ◆臣▶ 三臣 - のへで

- 1. Sample $\theta \sim p(\theta)$
- 2. Sample $\mathbf{y} \sim p(\mathbf{y}|\boldsymbol{\theta})$
- 3. Accept $\boldsymbol{\theta}$ iff $||\mathbf{y} \mathbf{y}^{\star}|| < \epsilon$

This algorithm produces samples from

$$p_{\epsilon}\left(oldsymbol{ heta}|\mathbf{y}^{\star}
ight) \propto p\left(oldsymbol{ heta}
ight) \int p(\mathbf{y}|oldsymbol{ heta}) \mathbb{I}\left(||\mathbf{y}-\mathbf{y}^{\star}|| < \epsilon
ight) \mathrm{d}\mathbf{y}$$

▲ロト ▲圖ト ▲ヨト ▲ヨト ヨー のへで

which tends to $p(\theta|\mathbf{y})$ with $\epsilon \to 0$.

ABC in pictures



<□▶ <□▶ < □▶ < □▶ < □▶ < □ > ○ < ○

ABC in pictures



θ

イロト イロト イヨト イヨト

æ

$\mathsf{ABC} \text{ in pictures}$



・ロト ・ 日 ・ ・ 日 ・ ・

물 🛌 🗄

ABC in pictures



A problem with basic ABC

- 1. Sample $oldsymbol{ heta} \sim p(oldsymbol{ heta})$
- 2. Sample $\mathbf{y} \sim p(\mathbf{y}|\boldsymbol{\theta})$
- 3. Accept $\boldsymbol{\theta}$ iff $||\mathbf{y} \mathbf{y}^{\star}|| < \epsilon$.

If there are many datapoints, then either ϵ is enormous or the probability of acceptance is going to be impractically small (the model will never reproduce *exactly* a large dataset).

ション ふゆ く 山 マ チャット しょうくしゃ

Introducing summary statistics

Solution found by [?]: reduce the dimension of y by computing some summary statistics s (y), and modify the algorithm:

◆□▶ ◆□▶ ◆□▶ ◆□▶ ● ● ●

- 1. Sample $oldsymbol{ heta} \sim p(oldsymbol{ heta})$
- 2. Sample $\mathbf{y} \sim p(\mathbf{y}|\boldsymbol{\theta})$
- 3. Accept $\boldsymbol{\theta}$ iff $||\mathbf{s}(\mathbf{y}) \mathbf{s}(\mathbf{y}^{\star})|| < \epsilon$.

Introducing summary statistics

- Solution found by [?]: reduce the dimension of y by computing some summary statistics s (y), and modify the algorithm:
- 1. Sample $oldsymbol{ heta} \sim p(oldsymbol{ heta})$
- 2. Sample $\mathbf{y} \sim p(\mathbf{y}|\boldsymbol{\theta})$
- 3. Accept $\boldsymbol{\theta}$ iff $||\mathbf{s}(\mathbf{y}) \mathbf{s}(\mathbf{y}^{\star})|| < \epsilon$.
- Provided that the choice of summary statistics is appropriate, this behaves reasonably and is actually computationally feasible.

ション ふゆ く 山 マ チャット しょうくしゃ

More advanced variants

- There are by now many, more efficient, variants on the original algorithm, based on MCMC, Sequential Monte Carlo, etc. [ref.]
- All of them require the definition of summary statistics, and are rather slow and difficult to tune.
- Using EP you can get rid of summary statistics, and obtain substantial speed-ups (10-100x, Barthelmé & Chopin, 2011, Barthelmé & Chopin, 2014, Barthelmé, Chopin, Cottet, 2015).
 - Caveat I: you can't get rid of summary statistics in all models

- Caveat II: implementation is a bit of work
- Caveat III: you get a Gaussian approximation (it's still EP)

How to get rid of summary statistics

- In the ABC-reject algorithm, we needed summary statistics because we had more than one datapoint.
- In EP we only integrate one datapoint at a time, so
 - No need for summary statistics!
 - We can just compute all hybrid moments using ABC-reject

(日) (伊) (日) (日) (日) (0) (0)

Our objective is

 $p_{\epsilon}(\boldsymbol{\theta}|\mathbf{y}^{\star}) \propto p(\boldsymbol{\theta}) \prod_{i=1}^{n} \left\{ \int f_{i}(y_{i}|\boldsymbol{\theta}) \mathbb{I}_{\left\{ \left\| y_{i}-y_{i}^{\star} \right\| \leq \epsilon
ight\}} dy_{i}
ight\}$

ABC-EP in one slide

- 1. Initialise site parameters $\lambda_1 \dots \lambda_n$. Global parameter: $\lambda = \sum \lambda_i$.
- 2. While not converged, loop over *i*:
 - 2.1 Form cavity: $\lambda_{-i} = \lambda \lambda_i$, hybrid $h_i(\theta) \propto l_i(\theta) \exp(s(\theta)^t \lambda_{-i})$
 - 2.2 Compute moments: $\eta_i = E_{h_i}(s(\theta))$ USING REJECTION ABC, transform back to natural parameters $\lambda_i = \nu(\eta_i) - \lambda_{-i}$

ション ふゆ く 山 マ チャット しょうくしゃ

2.3 Update global approximation: $\lambda = \lambda_{-i} + \lambda_i$

First example: alpha-stable distributions

 Alpha-stable densities are a class of univariate densities with potentially very heavy tails, that are popular in economics, because...

▲□▶ ▲圖▶ ▲臣▶ ★臣▶ ―臣 …の�?

First example: alpha-stable distributions

 Alpha-stable densities are a class of univariate densities with potentially very heavy tails, that are popular in economics, because...



◆□▶ ◆□▶ ◆□▶ ◆□▶ ● ● ●

(fig. from Brad DeLong's blog).

Alpha-stable densities

- ► No closed-form for the density function.
- ► We take data: n = 1200 AUD/GBP log-returns computed from daily exchange rates.
- Data assumed IID from alpha-stable distribution with parameters θ
- ▶ θ is α (tail heaviness), β (skewness), δ (location), γ (scale)

・ロト ・ 日 ・ ・ 日 ・ ・ 日 ・ ・ つ へ ()

Results from alpha-stable example



MCMC-ABC: 50 times more alpha-stable simulations than EP. "Exact" MCMC takes 60 hours.

э

Example II: race model for reaction times

- Hierarchical model with independent data (but not IID)
- Subject must choose between k alternatives. Evidence e_j(t) in favour of choice j follows a Brownian motion with drift:

$$\tau de_j(t) = m_j dt + dW_t^j.$$

Decision is taken when one evidence "wins the race"; see plot.



Data

1860 Observations (courtesy of M Maertens, TU Berlin), from a single human being, who must choose between "signal absent", and "signal present".



The hierarchical model

- ► The relative speed of the two racing diffusions changes according to experimental condition (≈random effect).
- Global parameters: boundaries, noise variance.
- ► 33 parameters in total (3 shared, 30 condition-specific).

・ロト ・ 日 ・ ・ 日 ・ ・ 日 ・ ・ つ へ ()

- ▶ 1860 observations, 30 subgroups.
- CPU time ~ 40 min

Reaction times: results



◆□▶ ◆□▶ ◆豆▶ ◆豆▶ ̄豆 _ のへぐ

Reaction times: results



◆□▶ ◆□▶ ◆臣▶ ◆臣▶ ─臣 ─ のへで

Reaction times: results

- Hierarchical model with ~ 30 parameters would be very challenging for standard ABC
- ABC-EP makes it do-able.
- Has actually been used again in an actual neuroscience paper (Park et al. 2016)

・ロト ・ 日 ・ ・ 日 ・ ・ 日 ・ ・ つ へ ()

Example III: ABC-EP with summary statistics

- Sometimes we can't get rid of summary statistics entirely:
 - Spatial extremes: each observation is a set of extreme rainfall values over different weather stations

・ロト ・ 日 ・ エ ヨ ・ ト ・ 日 ・ うらつ

- IID over years but not over stations (because of spatial dependencies)
- We want to infer spatial dependencies

Swiss rainfall



▲□▶ ▲□▶ ▲□▶ ▲□▶ ▲□ ● ● ●

ABC-EP in spatial extremes

- In recent work (Barthelmé, Chopin, Cottet, 2015) we suggest using ABC-EP with "local" summary statistics: summarise the observations over stations, but keep the successive years as IID sites
- Summary statistics is a robust estimate of spatial dependence: estimated value of a, b in following regression

 $\log |F(y_i(x_j)) - F(y_i(x_k))| = a + b \log ||x_j - x_k|| + \epsilon_{jk}, \quad 1 \le j < k \le d$ (F is the Fréchet CDF).

ABC-EP in spatial extremes

- ▶ We used the Swiss rainfall dataset (79 sites, 1962-2008).
- MCMC-ABC approach by Ehrardt & Smith (2012) essentially returns the prior after running for a week
- ABC-EP gives you something in about 3 hours
- Posterior is over the parameters of the covariance function (length-scale, height)

うして ふゆう ふほう ふほう うらつ

ABC-EP in spatial extremes



▲□▶ ▲圖▶ ▲臣▶ ▲臣▶ 三臣 - のへ(や)

ABC-EP with summary statistics

- Even if you can't get rid of summary statistics entirely, you can get still choose summary statistics that are "local" and low-dimensional.
- Because you're still integrating the data bit-by-bit, the acceptance rate is high and you get considerable speed-ups over MCMC
- Caveat: in this example, it took us a while to find the right set of local summary statistics

・ロト ・ 日 ・ エ ヨ ・ ト ・ 日 ・ うらつ

We are taking a deterministic algorithm and making it stochastic: have to be careful about Monte Carlo variance.

▲□▶ ▲圖▶ ▲臣▶ ★臣▶ ―臣 …の�?

- We are taking a deterministic algorithm and making it stochastic: have to be careful about Monte Carlo variance.
- In the paper we highlight a set of tricks, among which:
 - Ways to "recycle" previous model simulations or exploit Markov structure

・ロト ・ 日 ・ ・ 日 ・ ・ 日 ・ ・ つ へ ()

- We are taking a deterministic algorithm and making it stochastic: have to be careful about Monte Carlo variance.
- In the paper we highlight a set of tricks, among which:
 - Ways to "recycle" previous model simulations or exploit Markov structure

・ロト ・ 日 ・ ・ 日 ・ ・ 日 ・ ・ つ へ ()

Quasi-Monte Carlo

- We are taking a deterministic algorithm and making it stochastic: have to be careful about Monte Carlo variance.
- In the paper we highlight a set of tricks, among which:
 - Ways to "recycle" previous model simulations or exploit Markov structure
 - Quasi-Monte Carlo
 - Rao-Blackwellisation A.K.A. Conditional Monte Carlo

・ロト ・ 日 ・ エ ヨ ・ ト ・ 日 ・ うらつ

- We are taking a deterministic algorithm and making it stochastic: have to be careful about Monte Carlo variance.
- In the paper we highlight a set of tricks, among which:
 - Ways to "recycle" previous model simulations or exploit Markov structure
 - Quasi-Monte Carlo
 - Rao-Blackwellisation A.K.A. Conditional Monte Carlo
- A lot of these tricks are model-specific and require a bit of work.

うして ふゆう ふほう ふほう うらつ

Conclusion on ABC-EP

► ABC-EP can bring tremendous speed improvements, but:

- It is not trivial to implement
- If your likelihood leads to a multi-modal posterior, the best you can do is recover one mode (hard to know in advance in ABC settings)
- When you switch from deterministic moment computations to Monte Carlo ones, stability becomes a problem
 - We'll see that matters a lot for the algorithms in the next part of this tutorial

・ロト ・ 日 ・ エ ヨ ・ ト ・ 日 ・ うらつ

MCMC in large datasets

- A lot of attention currently on how to scale MCMC to large datasets (Angelino, Johnson, Adams, 2016)
- As everybody knows, we need to parallelise
- In 2014 two groups came up with the same idea
 - Split datasets, run independent MCMC chains
 - Use EP to synchronise
- Xu, M., Lakshminarayanan, B., Teh, Y. W., Zhu, J., and Zhang, B. (2014). Distributed Bayesian posterior sampling via moment sharing. In Advances in Neural Information Processing Systems
- Gelman, A., Vehtari, A., Jylanki, P., Robert, C., Chopin, N., and Cunningham, J. P. (2014). Expectation propagation as a way of life. ArXiv:1412.4869

Parallel EP

- EP parallelises trivially
- All we need to do is compute all site updates in parallel rather than sequentially

▲□▶ ▲圖▶ ▲臣▶ ★臣▶ ―臣 …の�?

Parallel EP in one slide

- 1. Initialise site parameters $\lambda_1 \dots \lambda_n$. Global parameter: $\lambda = \sum \lambda_i$.
- 2. While not converged:
 - 2.1 **Split:** For all *i*'s , do:
 - 2.1.1 Form cavity: $\lambda_{-i} = \lambda \lambda_i$, hybrid $h_i(\theta) \propto l_i(\theta) \exp(s(\theta)^t \lambda_{-i})$
 - 2.1.2 Compute moments: $\eta_i = E_{h_i}(s(\theta))$, transform back to natural parameters $\lambda_i \leftarrow \nu(\eta_i) \lambda_{-i}$

2.2 **Combine:** Update global approximation: $\boldsymbol{\lambda} \leftarrow \sum \boldsymbol{\lambda}_i$

MCMC-EP

- In our logistic regression application, a site was just a single datapoint and we could compute moments almost exactly using 1d quadrature
- What if we had sites with k datapoints?
 - ▶ We could maybe still do k = 3 using quadrature but it gets expensive

・ロト ・ 日 ・ ・ 日 ・ ・ 日 ・ ・ つ へ ()

- There's no hope of doing k = 500
- ► Use MCMC!

MCMC-EP

There's an additional insight: in hierarchical models, you only need to synchronise *global*, *shared* parameters using EP.

・ロト ・ 日 ・ ・ 日 ・ ・ 日 ・ ・ つ へ ()

 The parameters that are private to each batch can just be integrated over

MCMC-EP

- Strategy is very simple: you have *m* workers and *n* datapoints.
 - Split dataset into *m* batches of $k \approx n/m$ datapoints.
 - Each hybrid is now a Gaussian pseudo-prior times k likelihood terms
 - Compute moments using *m* parallel MCMC chains over your workers

・ロト ・ 日 ・ ・ 日 ・ ・ 日 ・ ・ つ へ ()

Update global approximation once everybody's done

Does it work?



From Gelman et al. Logistic regression example, k = 50, n = 2500

◆□▶ ◆□▶ ◆臣▶ ◆臣▶ 三臣 - のへで

Does it work?



Teh et al. (2016). Fully connected neural net on MNIST dataset.

◆□▶ ◆圖▶ ◆厘▶ ◆厘▶

3

Conclusion on combining EP and MCMC

- The results so far are proof-of-concept
 - Either good results on toy models
 - Or so-so results on non-toy models
- Stability is a problem, just like in ABC-EP, which required model-specific work
 - ▶ Need better theory, right now we have a collection of hacks
- It's potentially very promising, but we are still quite far from effective black-box EP (The Stan team is working on it, though, pers. communication)
- Extension to sequential settings: De Freitas et al. (2015) (haven't had time to look at it yet)

aEP, sEP: Even more approximate EP

- aEP is a simpler version of EP we originally introduced to study asymptotic properties of EP (Dehaene & Barthelmé, 2015)
 - Forget about individual site parameters, use an average cavity parameter

- Average cavity $oldsymbol{\lambda}_c = rac{n-1}{n}oldsymbol{\lambda}$
- It has a nice interpretation as an approximate projection algorithm:
 - ► Form hybrids, approximate all hybrids as Gaussians
 - Average hybrids
- Same asymptotic properties as EP

Is aEP practical?

- We originally didn't make much of it, but noted
 - ► aEP is easier to implement
 - It cuts on the linear algebra by half
- Hernandez-Lobato et al. (2015) introduced stochastic EP (sEP)
 - essentially a EP with a random update schedule
 - obtained good results on neural networks
 - claim reduction in memory footprint (true but of limited consequence)

▲□▶ ▲□▶ ▲□▶ ▲□▶ = ● ● ●

- M. Beaumont (talk at NIPS) found aEP more stable in ABC setting
- We also found that generic fixed-point acceleration schemes (SQUAREM, Varadhan, 2014) worked well on aEP
- Easier in aEP than EP because there are far fewer parameters

・ロト ・ 日 ・ ・ 日 ・ ・ 日 ・ ・ つ へ ()

Potentially promising

General conclusions

- The original EP algorithm is extremely successful in GLMs, GAMs, etc., everybody should be using it
 - (still need quality implementation comparable to INLA or mgcv)
- ► EP is very promising as a generic black-box inference scheme

うして ふゆう ふほう ふほう うらつ

- in ABC settings
- for large hierarchical models
- Early days
 - Either a lot of model-specific work (ABC-EP)
 - Or proof-of-concept
 - aEP, sEP interesting direction

Things I wish I had time to mention

Corrections to EP

find EP approximation and improve it using expansions

・ロト ・ 日 ・ ・ 日 ・ ・ 日 ・ ・ つ へ ()

- Reviewed in Opper's lecture notes
- Marginal likelihood approximation
- Double-loop algorithms
- EP for bilinear models
- EP for Gibbs distributions
 - e.g. Ising model, see Opper's lecture notes

Collaborators

Thanks to Nicolas Chopin, Vincent Cottet, Guillaume Dehaene, Gina Gruenhage, Manfred Opper

▲□▶ ▲圖▶ ▲臣▶ ★臣▶ ―臣 …の�?