Asymptotic analysis of genealogies induced by sequential Monte Carlo algorithms

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1 Introduction

- what is SMC and what is it used for?
- basic SMC algorithm
- the three main tasks: prediction, filtering, smoothing
- the problem with smoothing: ancestral vs. weight degeneracy
- motivating plots
- organisation of the report

. . . .

For our purposes it is sufficient to describe sequential Monte Carlo algorithms in the context of inference on a time-homogeneous hidden Markov model. Consider the following model:

$$X_0 \sim \mu(\cdot)$$

$$X_{t+1} \mid (X_t = x_t) \sim f(\cdot | x_t) \qquad t = 0, \dots, T-1$$

$$Y_t \mid (X_t = x_t) \sim g(\cdot | x_t) \qquad t = 0, \dots, T$$

where $\{X_t\}_{t=0}^T$ is an unobservable Markov chain and the observables $\{Y_t\}_{t=0}^T$ satisfy $Y_t \perp \{Y_{-t}, X_{-t}\} \mid X_t$. The conditional independence structure is described by the following graphical model.

$$Y_{t-1} \qquad Y_t$$

$$\uparrow g \qquad \uparrow g$$

$$\cdots \xrightarrow{f} X_{t-1} \xrightarrow{f} X_t \xrightarrow{f} \cdots$$

We assume that the transition and emission kernels have densities which are denoted by f and g respectively, but this is not necessary in general. We only require that we can sample from $\mu(\cdot)$ and $f(\cdot|x)$, and calculate unnormalised potentials g(y|x), for all x, y.

2 SMC as a coalescent

- k-coalescent & Kingman coalescent
- pop gen literature about large population cts time limits of various models
- resampling viewed backwards in time: branching process \rightarrow coalescent process
- asymptotic properties of SMC lit review: CLT, path storage, coalescence etc.
- the gap in knowledge that we aim to fill
- (remark: although SMC has other problems in high dimension, the coalescence rate doesn't depend on the dimension...)

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3 Conditional SMC

- motivation: particle MCMC, need for multiple lineages
- conditional (multinomial) SMC algorithm and its context within particle Gibbs

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- result: coalescence rate etc in terms of standard multinomial one; verification of assumptions of KJJS theorem (but exile horrible calculations to appendix)
- simulations & conclusions thence

4 Alternative resampling schemes

- multinomial not really used in practice, but other schemes hard to analyse
- overview of the main variance-reducing schemes
- results: theorem for residual resampling (hopefully)
- maybe results for other schemes
- simulation comparing all of them, and conclusions thence

5 Discussion

- results so far
- impact of this work: to practitioners, to enriching the SMC literature, interpretation within pop gen.
- future directions

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