

Hierarchical Monte Carlo Fusion

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**The
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Outline

Monte Carlo Fusion

- Fork-and-join

- Constructing a rejection sampler

- Double Langevin Approach

Hierarchical Fusion

- Time-adapting Monte Carlo Fusion

- Divide-and-Conquer SMC with Fusion

Logistic Regression Example

Ongoing directions

Fusion Problem

- Target:

$$\pi(\mathbf{x}) \propto \prod_{c=1}^C f_c(\mathbf{x})$$

where each *sub-posterior*, $f_c(\mathbf{x})$, is a density representing one of the C distributed inferences we wish to unify

- Assume we can sample $\mathbf{x}^{(c)} \sim f_c(\mathbf{x})$
- Applications:
 - Expert elicitation: combining views of multiple experts
 - Privacy setting
 - Big Data (by construction)
 - Tempering (by construction)

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Fusion for Big Data

- Consider we have data \mathbf{x} with a large number of observations n
- The likelihood $\ell(\mathbf{x} | \theta)$ becomes **expensive** to calculate
 - This makes MCMC prohibitively **slow** for big data
- **Potential solution:**

$$\pi(\theta | \mathbf{x}) \propto \prod_{i=1}^n \ell(\mathbf{x}_i | \theta) \pi(\theta) = \prod_{c=1}^C \ell(\mathbf{x}_c | \theta) \pi(\theta)^{\frac{1}{C}}$$

where \mathbf{x}_c denotes the c -th subset for $c = 1, \dots, C$ and $\pi(\theta) = \prod_{c=1}^C \pi(\theta)^{\frac{1}{C}}$ is the prior

- Advantage: inference on each smaller dataset can be conducted independently and in **parallel**

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Fusion for Tempering

- Consider the *power-tempered target distribution*
 $\pi_\beta(\mathbf{x}) = [\pi(\mathbf{x})]^\beta$ for $\beta \in (0, 1]$
- MCMC can become computationally expensive to sample from *multi-modal densities* and can get stuck in modes
- Potential solution:

$$\pi(\mathbf{x}) = \pi(\mathbf{x})^{\frac{1}{\beta} \cdot \beta} = \prod_{c=1}^{\frac{1}{\beta}} \pi_\beta(\mathbf{x})$$

where $\frac{1}{\beta} \in \mathbb{N}$

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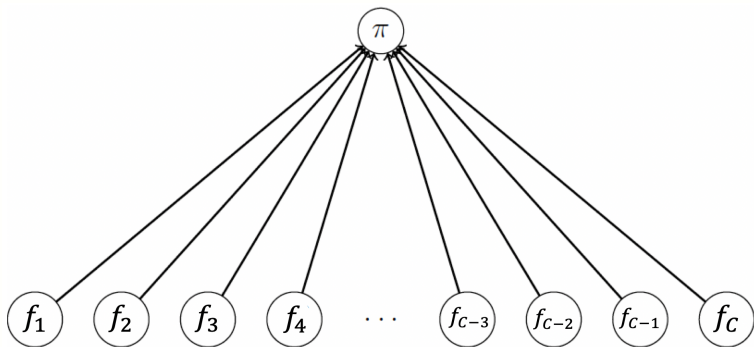
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Fork-and-join

The **fork-and-join** approach:



Current Fork-and-Join Methods

- Several fork-and-join methods have been developed. For instance
 - Kernel density averaging [Neiswanger et al., 2013]
 - Weierstrass sampler [Wang and Dunson, 2013]
 - Consensus Monte Carlo [Scott et al., 2016]
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Constructing a rejection sampler - An Extended Target

Proposition

Suppose that $p_c(\mathbf{y} \mid \mathbf{x}^{(c)})$ is the transition density of a *stochastic process with stationary distribution* $f_c^2(\mathbf{x})$. The $(C + 1)d$ -dimensional (fusion) density proportional to the integrable function

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Rejection Sampling (Double Langevin Approach)

- There are many possible choices for $p_c(\mathbf{y} | \mathbf{x})$
- Let $p_c(\mathbf{y} | \mathbf{x}) := p_{T,c}(\mathbf{y} | \mathbf{x})$, the transition density of the d -dimensional (double) Langevin (DL) diffusion processes $\mathbf{X}_t^{(c)}$ for $c = 1, \dots, C$, from \mathbf{x} to \mathbf{y} for a pre-defined time $T > 0$ given by

$$d\mathbf{X}_t^{(c)} = \nabla \log f_c(\mathbf{X}_t^{(c)}) dt + d\mathbf{W}_t^c,$$

- $\mathbf{W}_t^{(c)}$ is d -dimensional Brownian motion
- ∇ is the gradient operator over \mathbf{x}
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- Consider the proposal density h for the extended target g :

$$h(\mathbf{x}^{(1)}, \dots, \mathbf{x}^{(C)}, \mathbf{y}) \propto \prod_{c=1}^C \left[f_c(\mathbf{x}^{(c)}) \right] \cdot \exp\left(-\frac{C \cdot \|\mathbf{y} - \bar{\mathbf{x}}\|^2}{2T}\right)$$

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- T is an arbitrary positive constant

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Rejection Sampling - acceptance probability

- Acceptance probability:

$$\frac{g(\mathbf{x}^{(1)}, \dots, \mathbf{x}^{(C)}, \mathbf{y})}{h(\mathbf{x}^{(1)}, \dots, \mathbf{x}^{(C)}, \mathbf{y})} \propto \rho \cdot Q$$

where

$$\begin{cases} \rho := e^{-\frac{C\sigma^2}{2T}}, & \sigma^2 = \frac{1}{C} \sum_{c=1}^C \|\mathbf{x}^{(c)} - \bar{\mathbf{x}}\|^2 \\ Q := \mathbb{E}_{\bar{\mathbb{W}}} \left(\prod_{c=1}^C \left[\exp \left\{ - \int_0^T \left(\phi_c(\mathbf{x}_t^{(c)}) - \Phi_c \right) dt \right\} \right] \right) \end{cases}$$

where $\bar{\mathbb{W}}$ denotes the law of C independent Brownian bridges $\mathbf{x}_t^{(1)}, \dots, \mathbf{x}_t^{(C)}$ with $\mathbf{x}_0 = \mathbf{x}^{(c)}$ and $\mathbf{x}_T^{(c)} = \mathbf{y}$

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Q Acceptance Probability

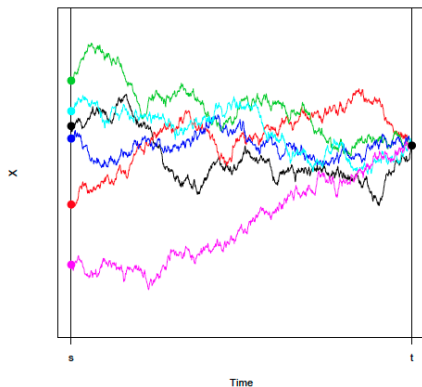
$$Q := \mathbb{E}_{\mathbb{W}} \left(\prod_{c=1}^C \left[\exp \left\{ - \int_0^T \left(\phi_c(\mathbf{x}_t^{(c)}) - \Phi_c \right) dt \right\} \right] \right)$$

where

- $\phi_c(\mathbf{x}) = \frac{1}{2} \left(\|\nabla \log f_c(\mathbf{x})\|^2 + \Delta \log f_c(\mathbf{x}) \right)$
- Φ_c are constants such that for all \mathbf{x} , $\phi_c(\mathbf{x}) \geq \Phi_c$ for $c \in \{1, \dots, C\}$
- Events of probability Q can be simulated using **Poisson thinning** and methodology called **Path-space Rejection Sampling (PSRS)** or the **Exact Algorithm** (Beskos et al. [2005], Beskos et al. [2006], Pollock et al. [2016])

Interpretation

- Correct a simple average \bar{x} of sub-posterior values to a Monte Carlo draw from $\pi(x)$ with acceptance probability $\rho \cdot Q$



Double Langevin Approach - Summary

- Proposal:

$$h(\mathbf{x}^{(1)}, \dots, \mathbf{x}^{(C)}, \mathbf{y}) \propto \prod_{c=1}^C [f_c(\mathbf{x}^{(c)})] \cdot \exp\left(-\frac{C \cdot \|\mathbf{y} - \bar{\mathbf{x}}\|^2}{2T}\right)$$

- Accept \mathbf{y} as a draw from fusion density π with probability:

$$\frac{g(\mathbf{x}^{(1)}, \dots, \mathbf{x}^{(C)}, \mathbf{y})}{h(\mathbf{x}^{(1)}, \dots, \mathbf{x}^{(C)}, \mathbf{y})} \propto \rho \cdot Q$$

- Monte Carlo Fusion Algorithm:
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Limitations of above rejection sampler

1. **Scalability**: the acceptance probability of Monte Carlo fusion **can** be small, especially when C is large or d is large
2. **Use of simple average** \bar{x} of sub-posterior samples as the proposal
 - but should we use a weighted average?
3. **Use of same time** T for each sub-posterior
 - but different cores / sub-posteriors could contain different amounts of information
4. **PSRS**: methodology for PSRS can be computationally expensive

Aim: To construct a fusion algorithm / framework to alleviate some of these limitations

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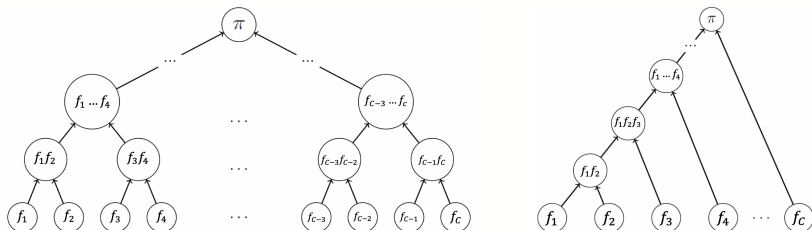
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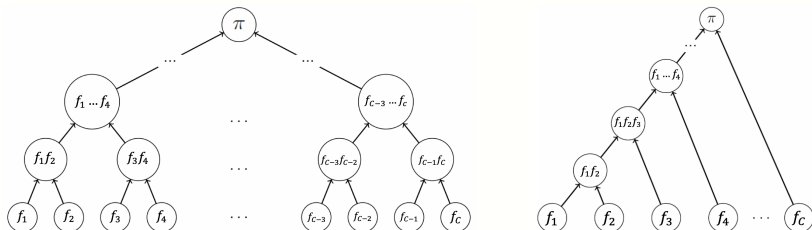
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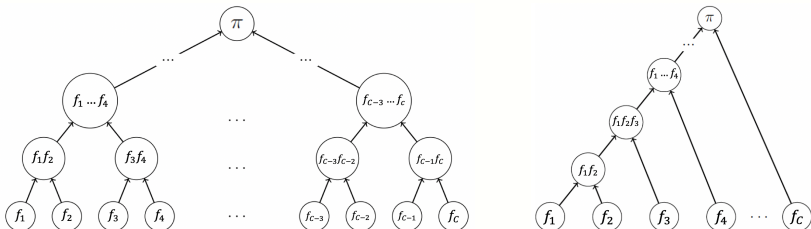
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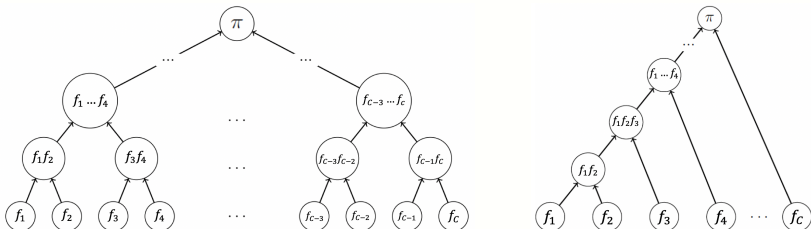
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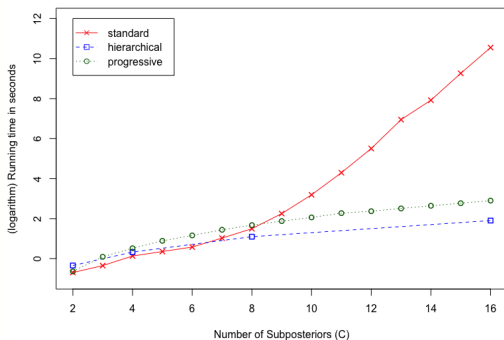
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Example

- Target: $\pi(x) \propto e^{-\frac{x^4}{2}}$
- Sub-posteriors: $f_c(x) = e^{-\frac{x^4}{2c}}$ for $c = 1, \dots, C$

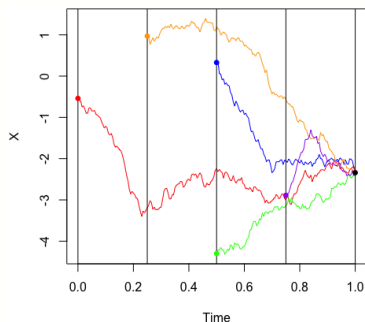
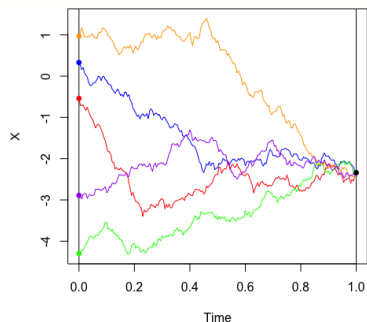


Time-adapting Monte Carlo Fusion

2. Use of simple average \bar{x} of sub-posterior samples as the proposal
3. Use of same time T for each sub-posterior

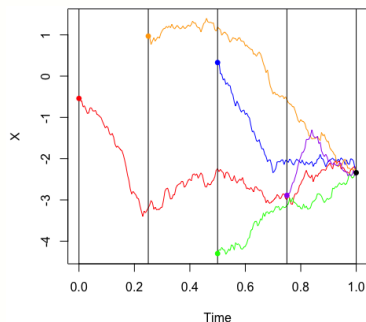
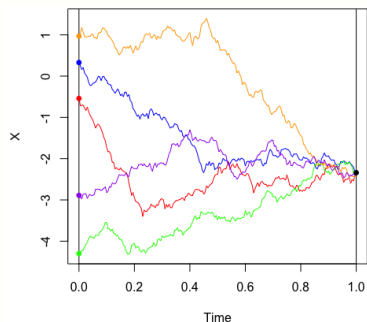
Time-adapting Monte Carlo Fusion

- **Solution:** Time-adapting Monte Carlo Fusion
 - We assign **weights** to each sub-posterior and use a weighted average: $\tilde{x} = \sum_c w_c x^{(c)} / \sum_c w_c$
 - Time T is **adapted** for each posterior: $T_c = \frac{T}{w_c}$ for $c = 1, \dots, C$



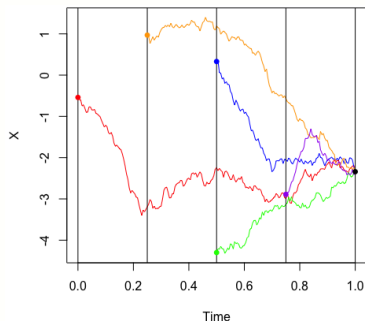
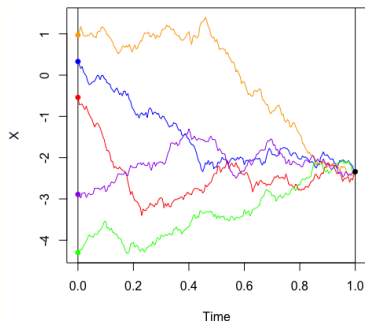
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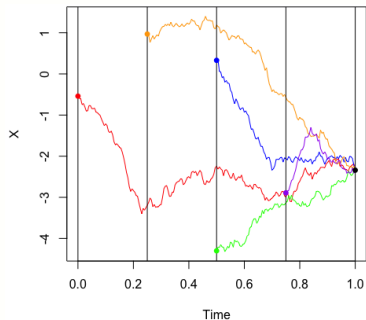
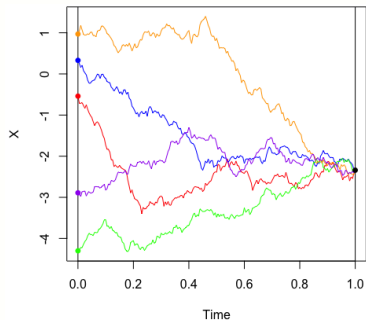
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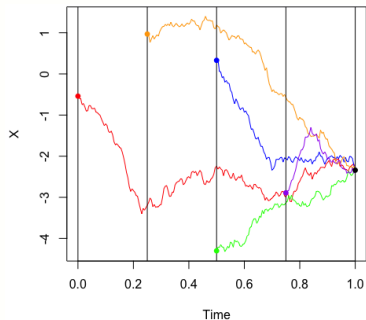
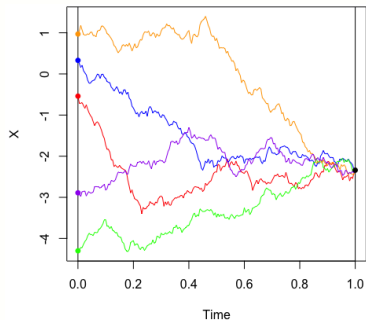
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- Time-adapting Monte Carlo Fusion Algorithm:
 1. Choose time T and weights for sub-posteriors $w_c, c = 1, \dots, C$
 2. Simulate $\mathbf{x}^{(c)} \sim f_c(\mathbf{x})$ and $\mathbf{y} \sim \mathcal{N}(\bar{\mathbf{x}}, \frac{T \mathbb{I}_d}{\sum_c w_c})$
 3. Accept \mathbf{y} with probability $\rho^{ta} \cdot Q^{ta}$



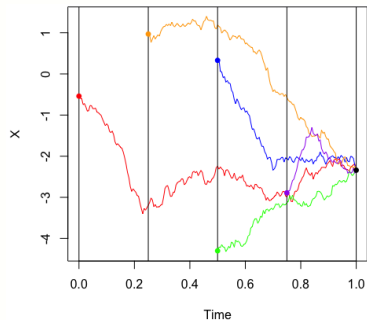
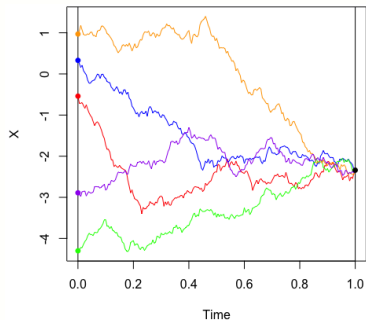
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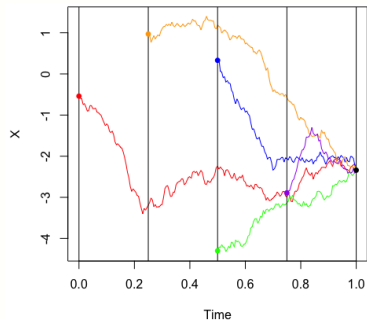
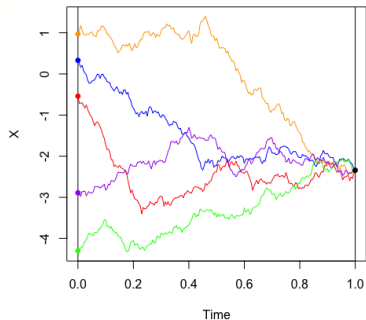
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Divide-and-Conquer SMC

4. PSRS: methodology for PSRS can be computationally expensive

- **Solution:** Sequential Monte Carlo in the hierarchical fusion framework
- Rejection sampling can be **wasteful**: large number of proposed samples are rejected
- Motives the use of **Sequential Importance Sampling / Resampling** ideas
 - **Replace** the rejection sampling steps with importance sampling steps
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- Predicting if customers defaulted on their payments using gender and education levels ($n = 30,000$ and $d = 5$)
- Split into $C = 32$ subsets and apply the following methods:
 1. Consensus Monte Carlo [Scott et al., 2016]
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Logistic Regression Example - Taiwan default payments

How to apply Hierarchical Time-adapting SMC Fusion

- Choose a time $T = 0.5$
- Use **balanced hierarchical tree** where we combine $m = 2$ sub-posteriors at each level, i.e. have $L = 6$ levels in the tree
- Weights are chosen according to **how much data they have relative to the bottom level**:
 - At $L = 6$: sub-posteriors are given weight $w_c = 1$ for $c = 1, \dots, 32$, i.e. $T_c = 0.5$
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- To compare methods we calculate the **integrated absolute distance**

$$IAD = \frac{1}{2^d} \sum_{j=1}^d \int |\hat{f}(\mathbf{x}_j) - f(\mathbf{x}_j)| d\mathbf{x}_j$$

where $\hat{f}(\mathbf{x}_j)$ is the marginal density for \mathbf{x}_j based on the method applied and $f(\mathbf{x}_j)$ is the benchmark estimate (obtained using NUTS with Stan)

- **Fix computational cost** by restricting run-time allowed for algorithm

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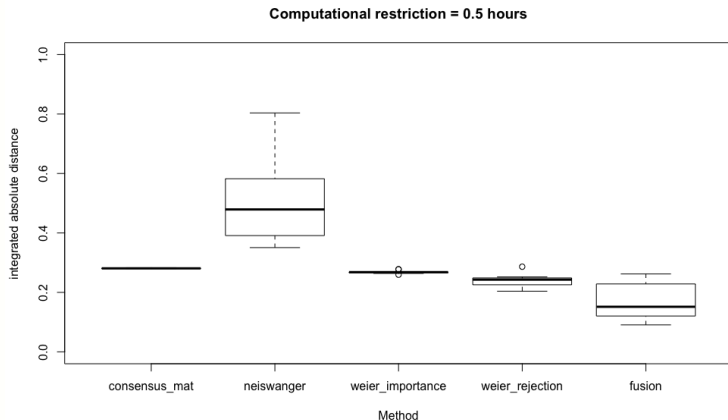
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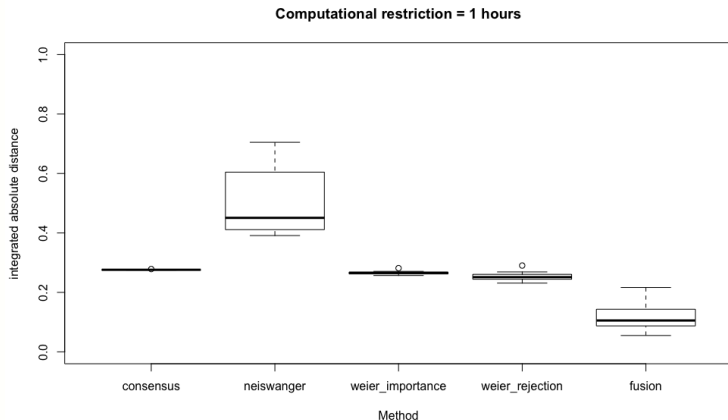
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Ongoing directions

- Combining **conflicting sub-posteriors**
- **Confidential fusion (Con-fusion)**: where sharing information/data between cores is **not** permitted
- **Bayesian Fusion**: tailored to big data Bayesian problems
- **Different proposals**: e.g. **Ornstein-Uhlenbeck** bridges, more general Langevin diffusion with **pre-conditioning matrix** for each sub-posterior
- **Theory** for D&C-SMC with Monte Carlo Fusion

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References

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