

Using Bagged Posteriors for Robust Inference

Jonathan Huggins
Harvard University

Joint work with Jeff Miller

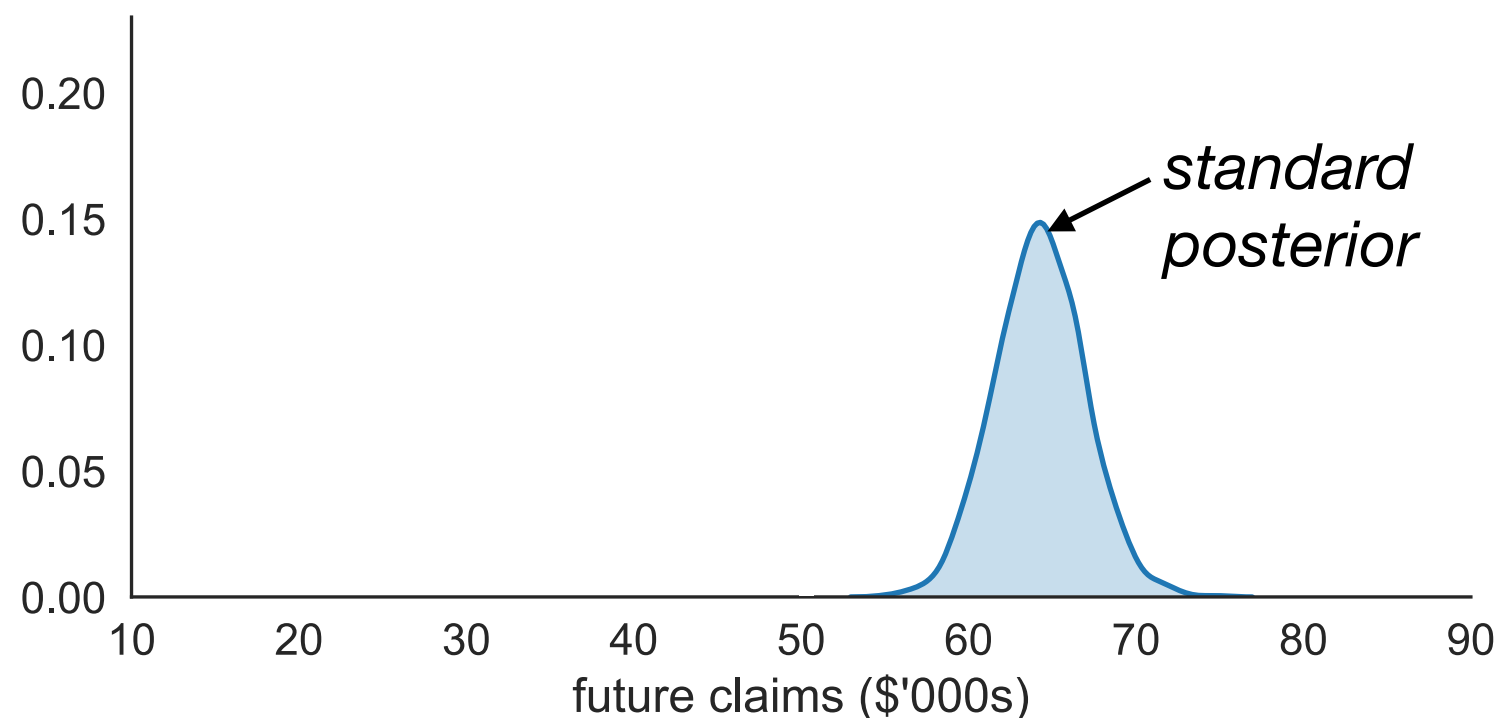
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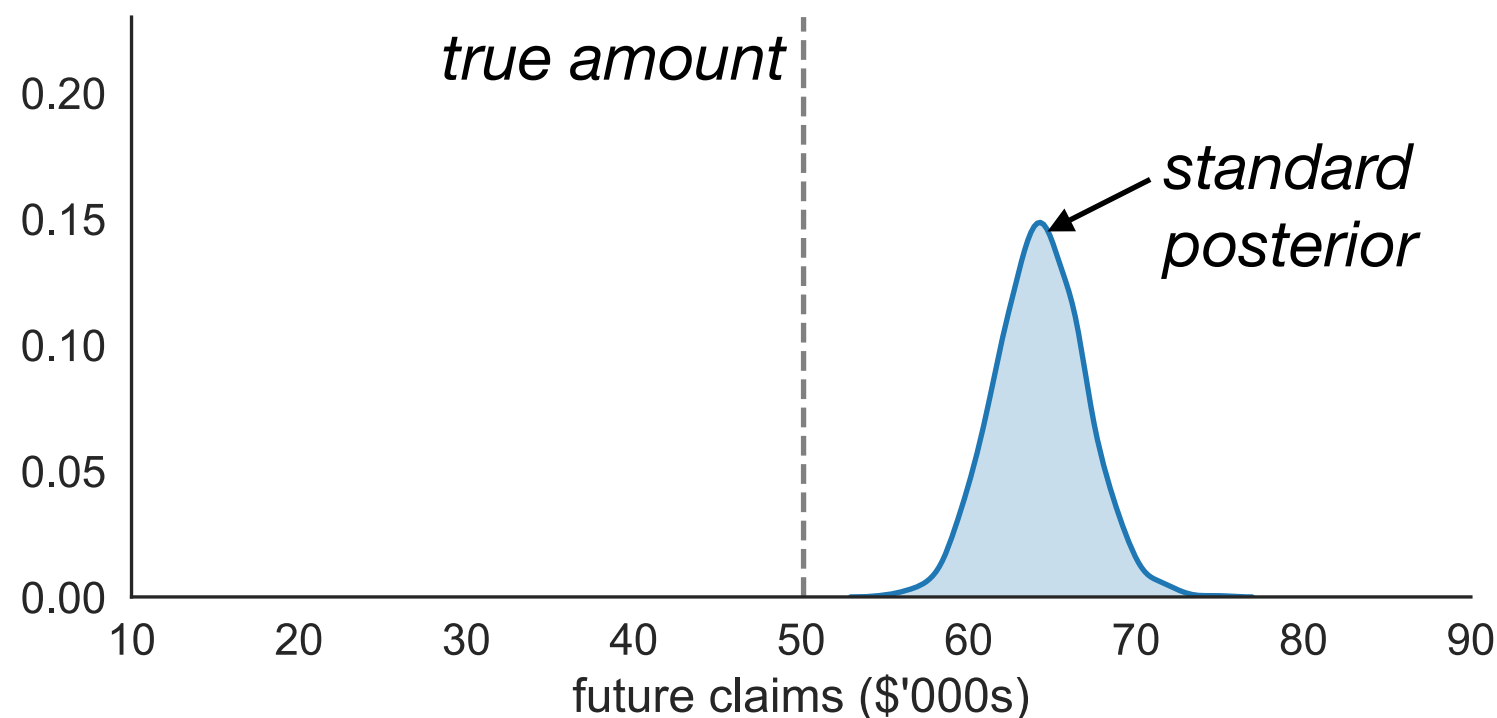
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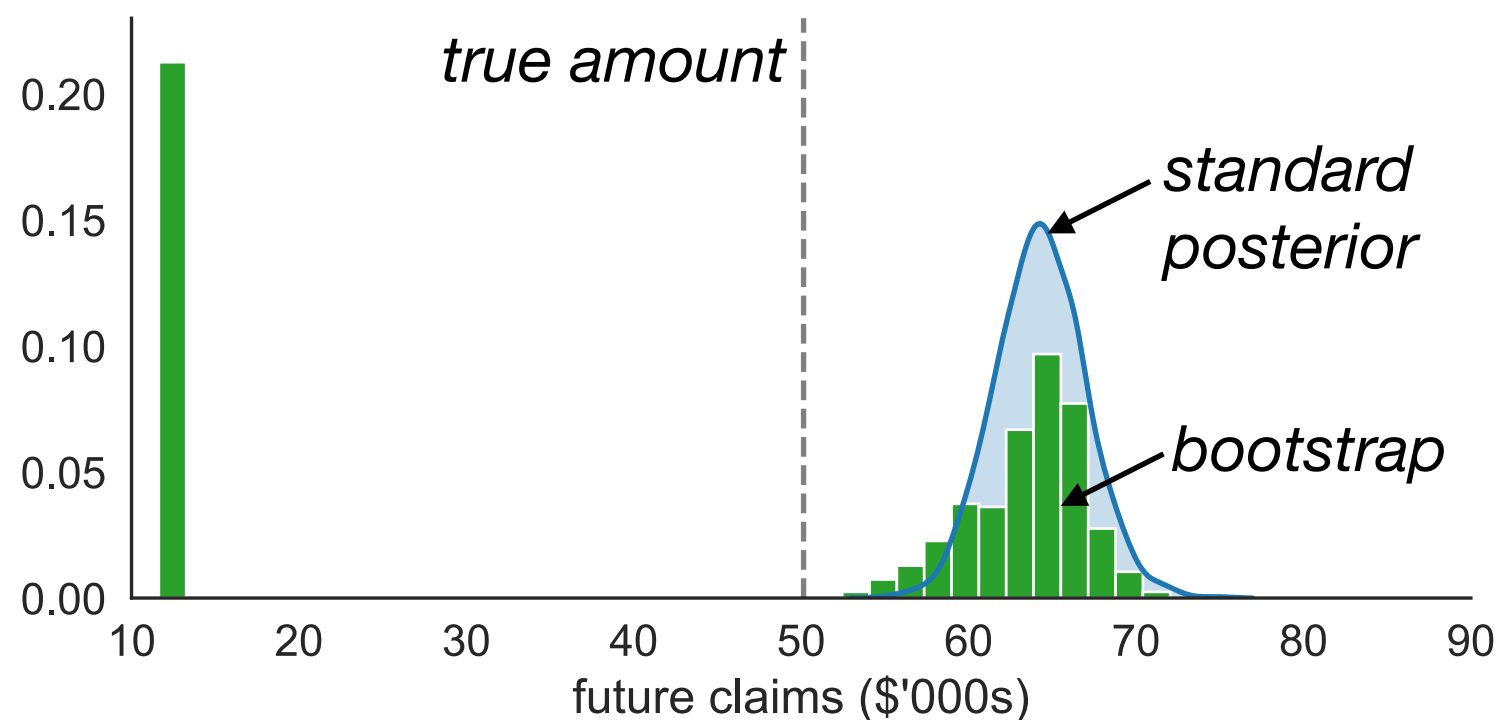
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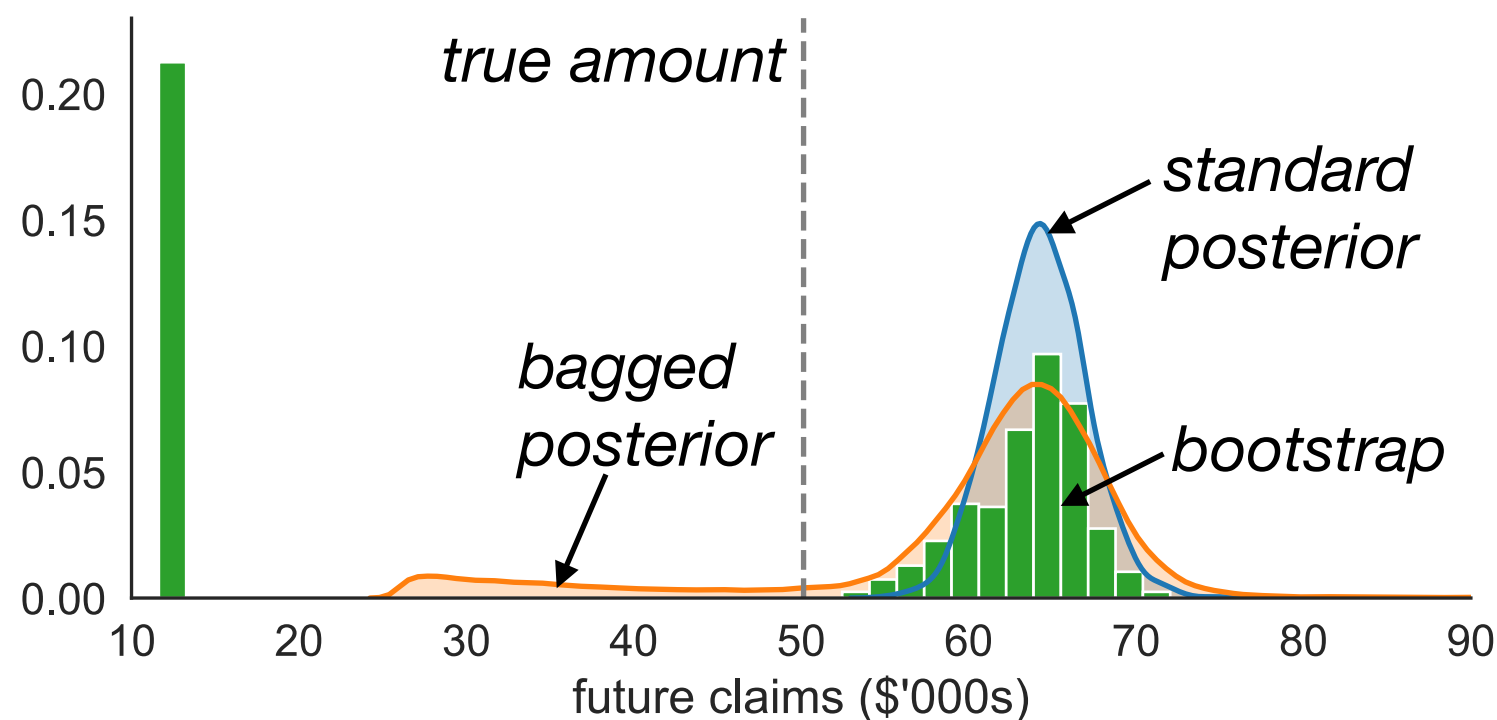
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- **Solution:** use the *bagged posterior* (BayesBag)



Agenda

- ➡ **BayesBag for parameter inference (and prediction)**
- BayesBag theory and methodology
- BayesBag for model selection

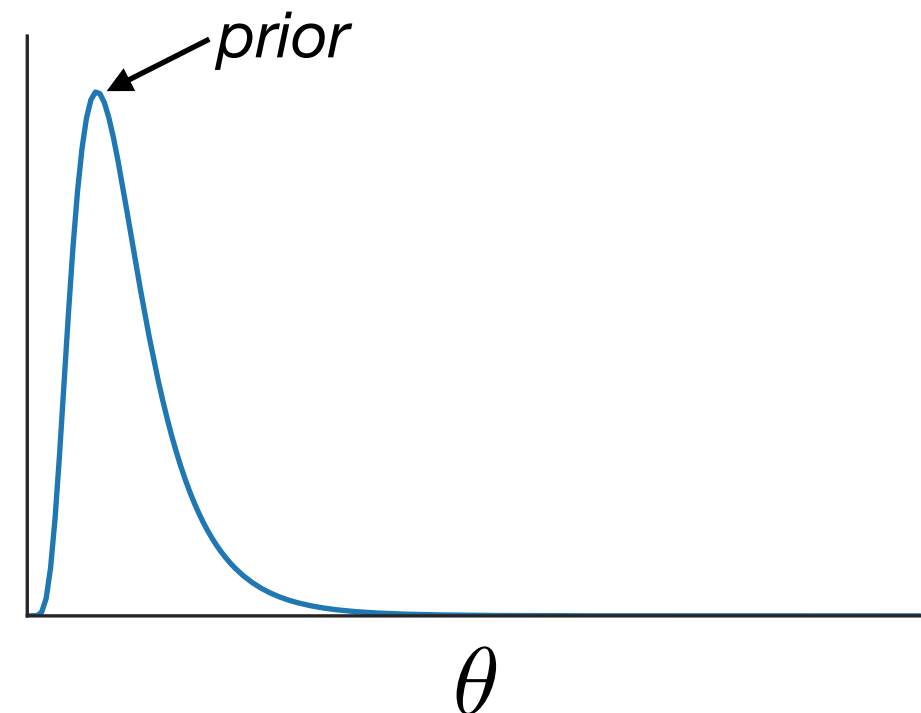
Bayesian inference

Bayesian inference

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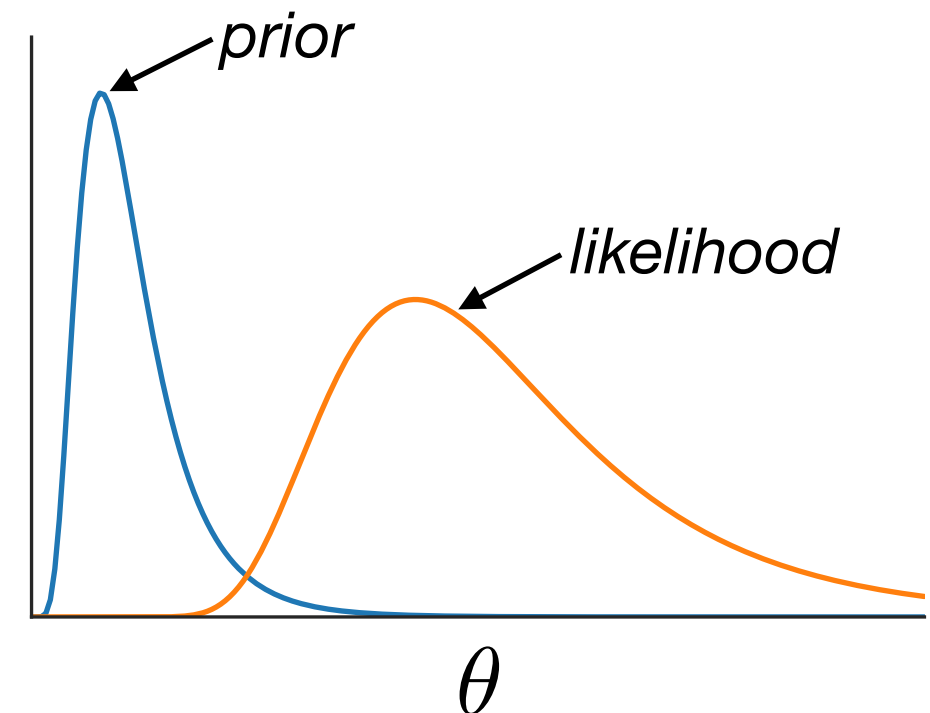
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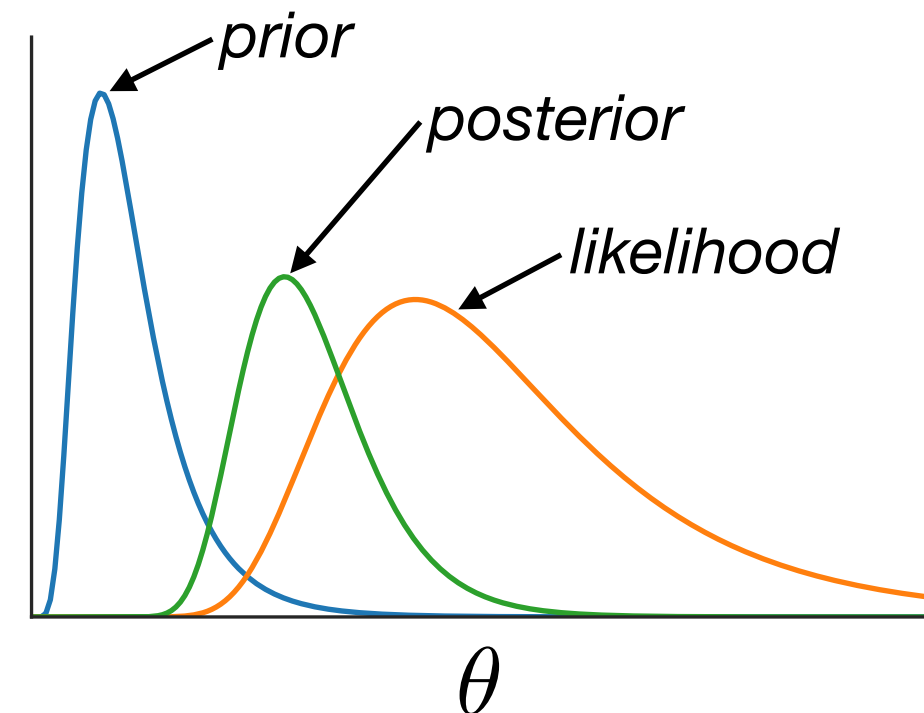
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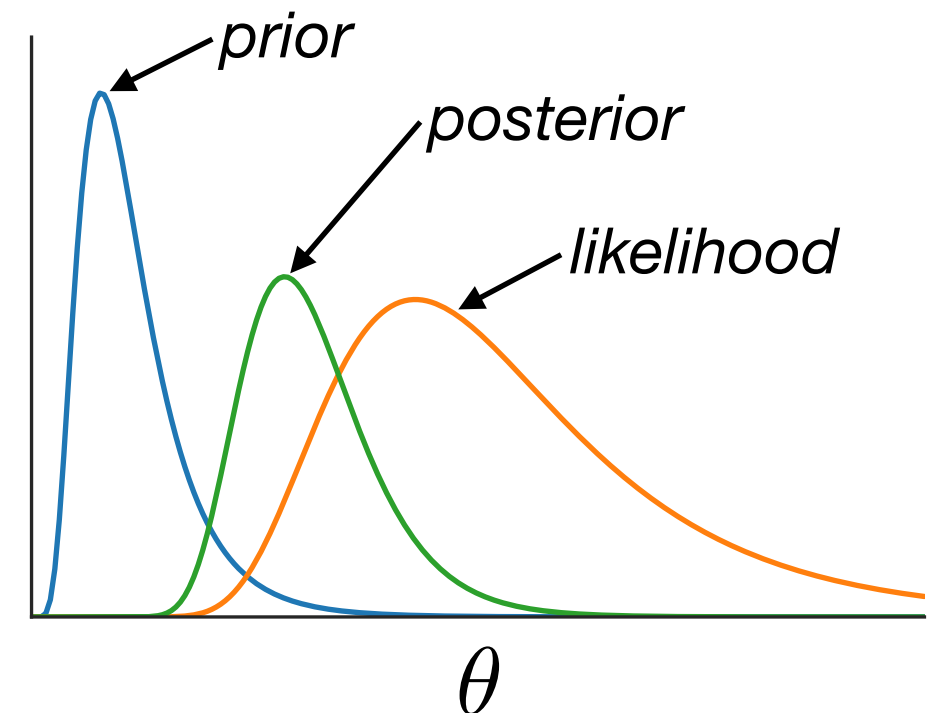
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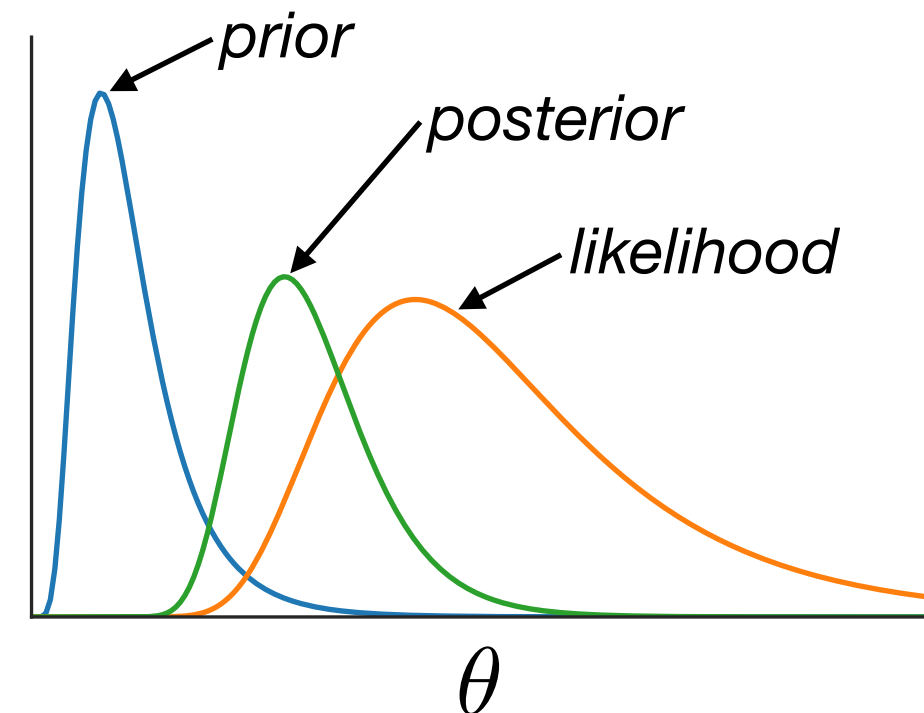


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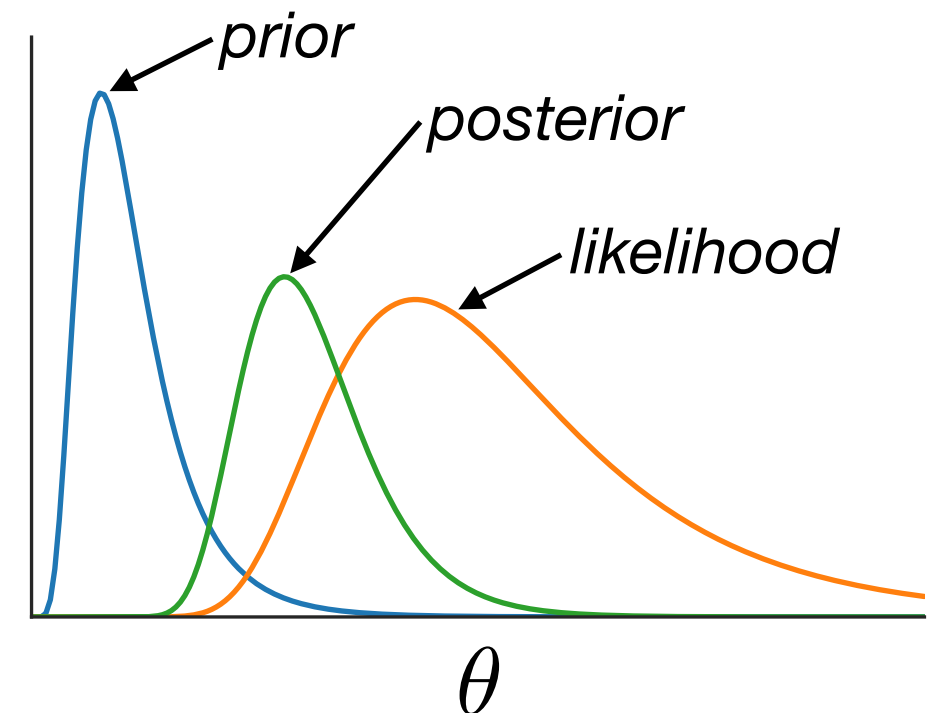


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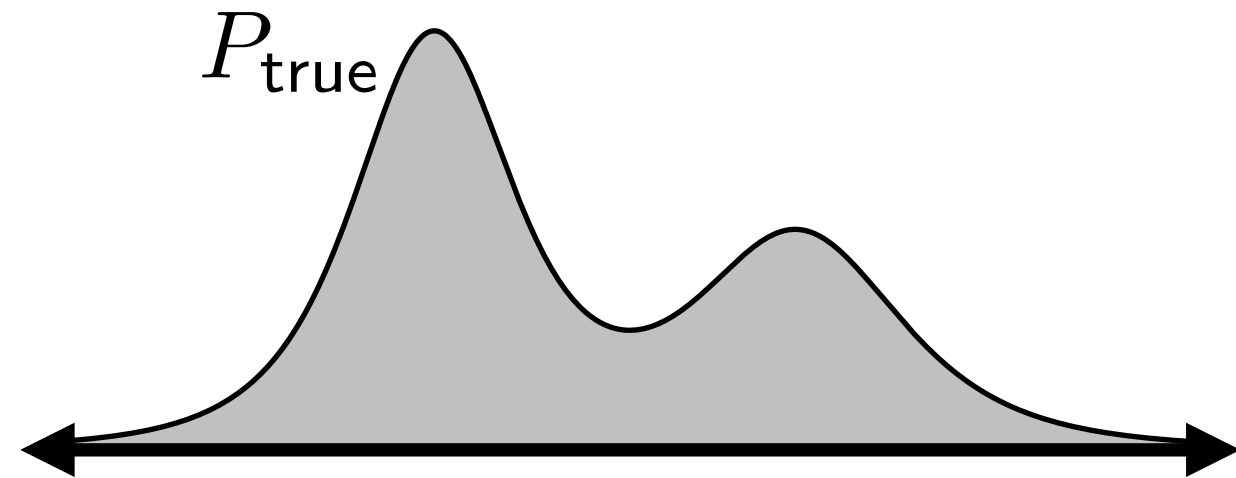
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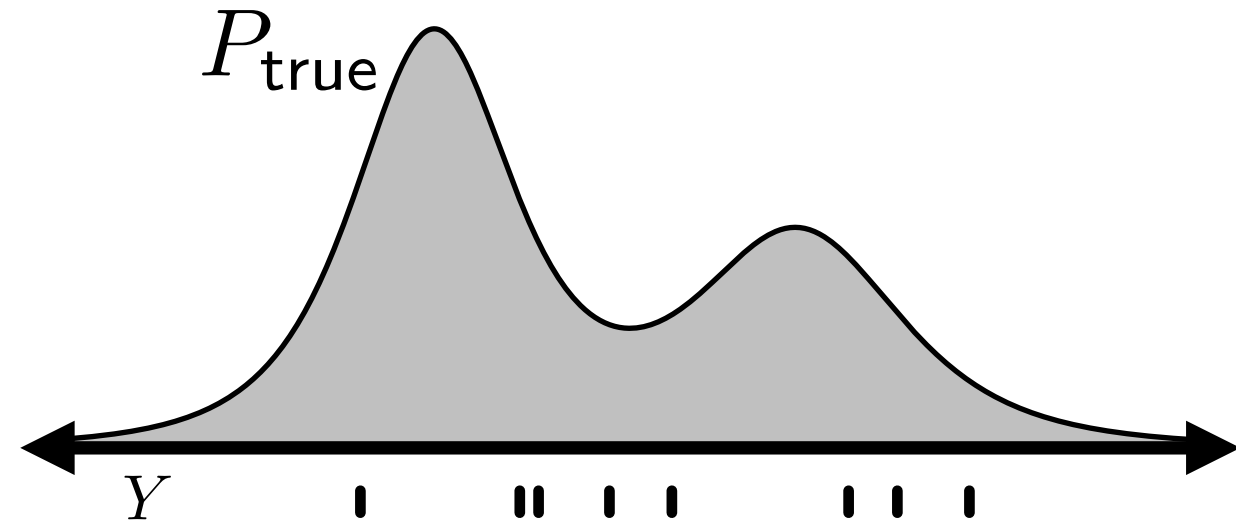


Bootstrapping



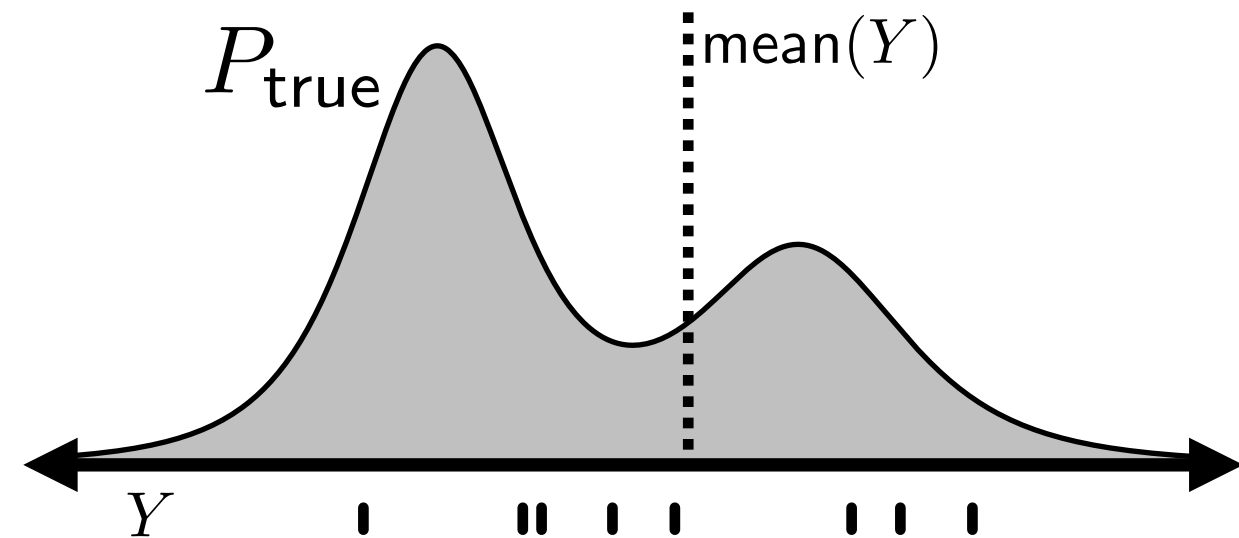
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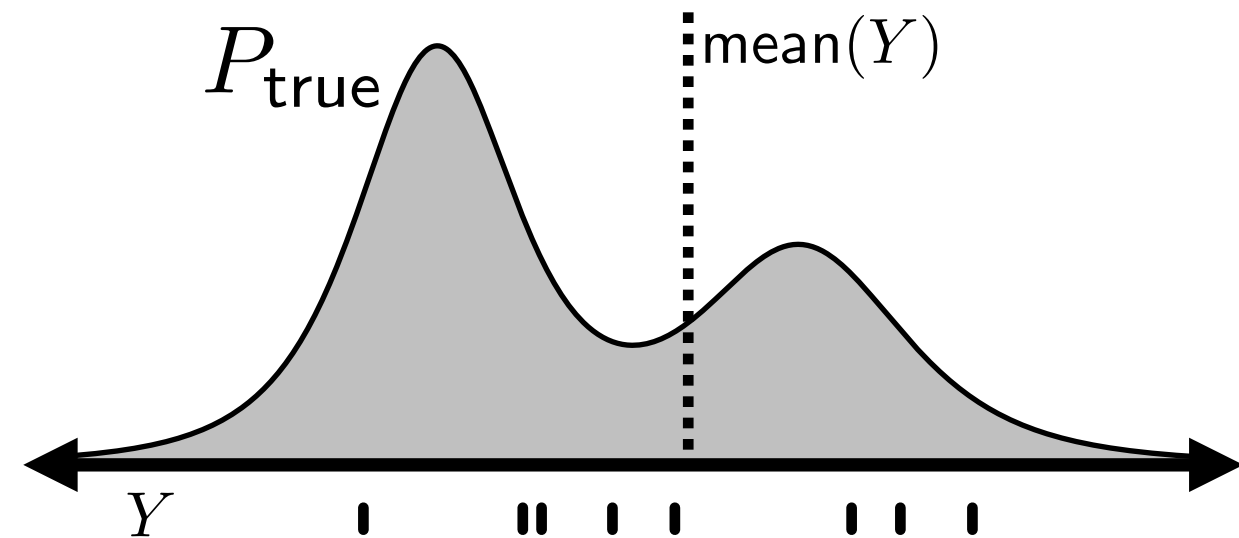
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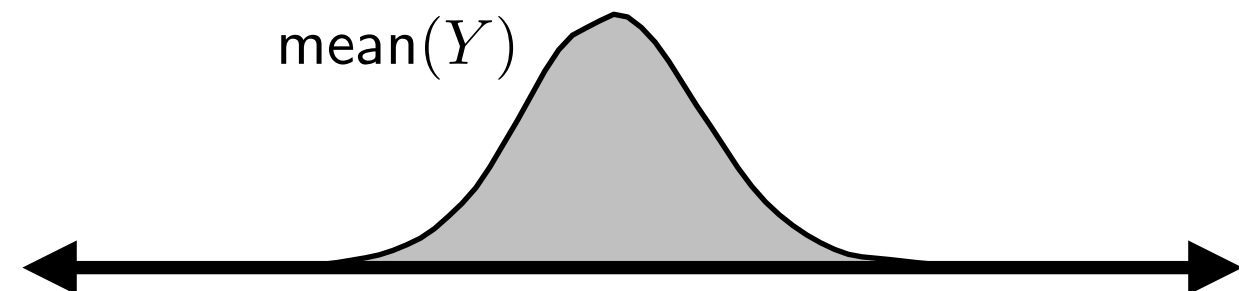


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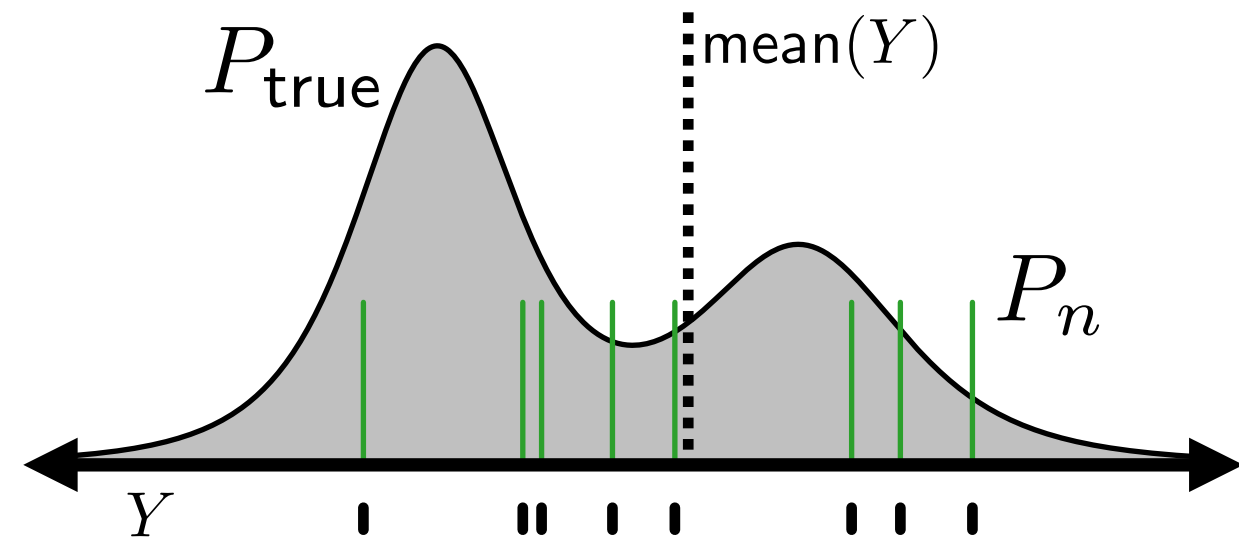


distribution of...

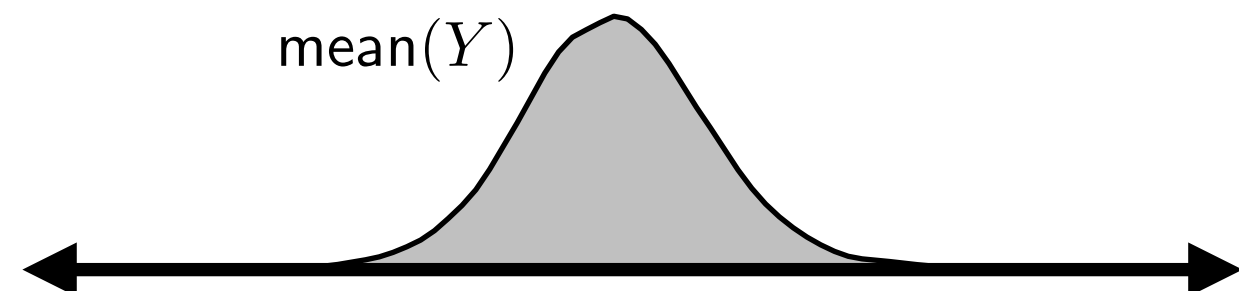


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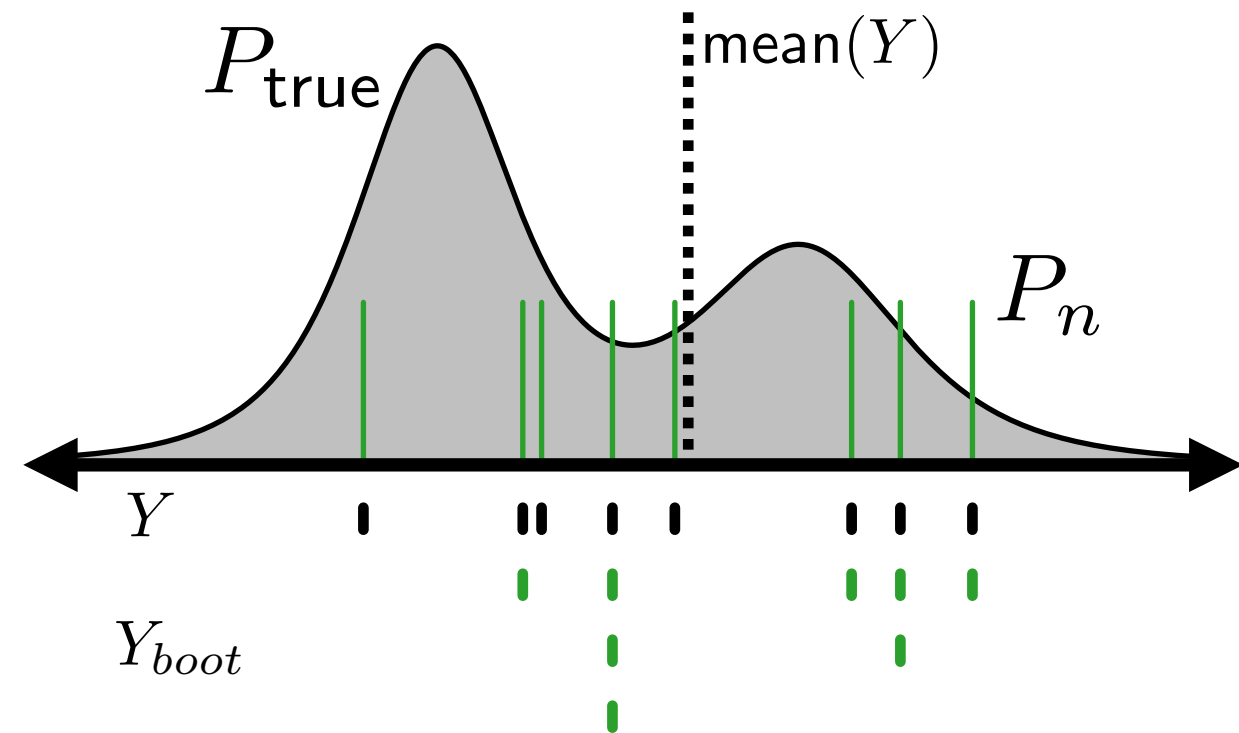


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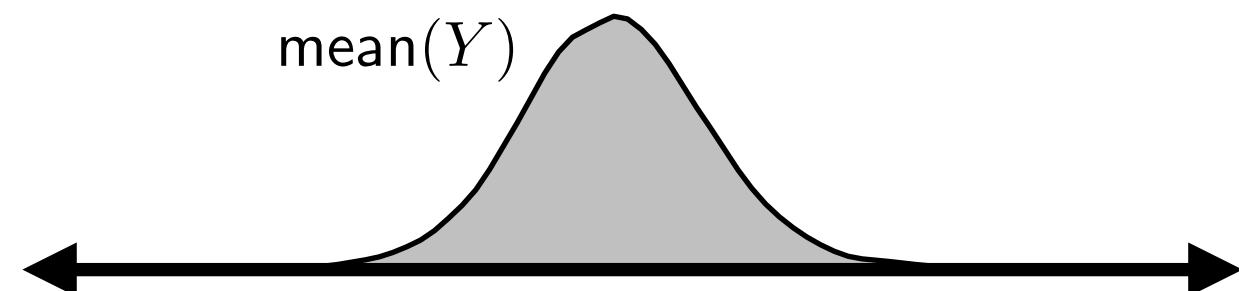


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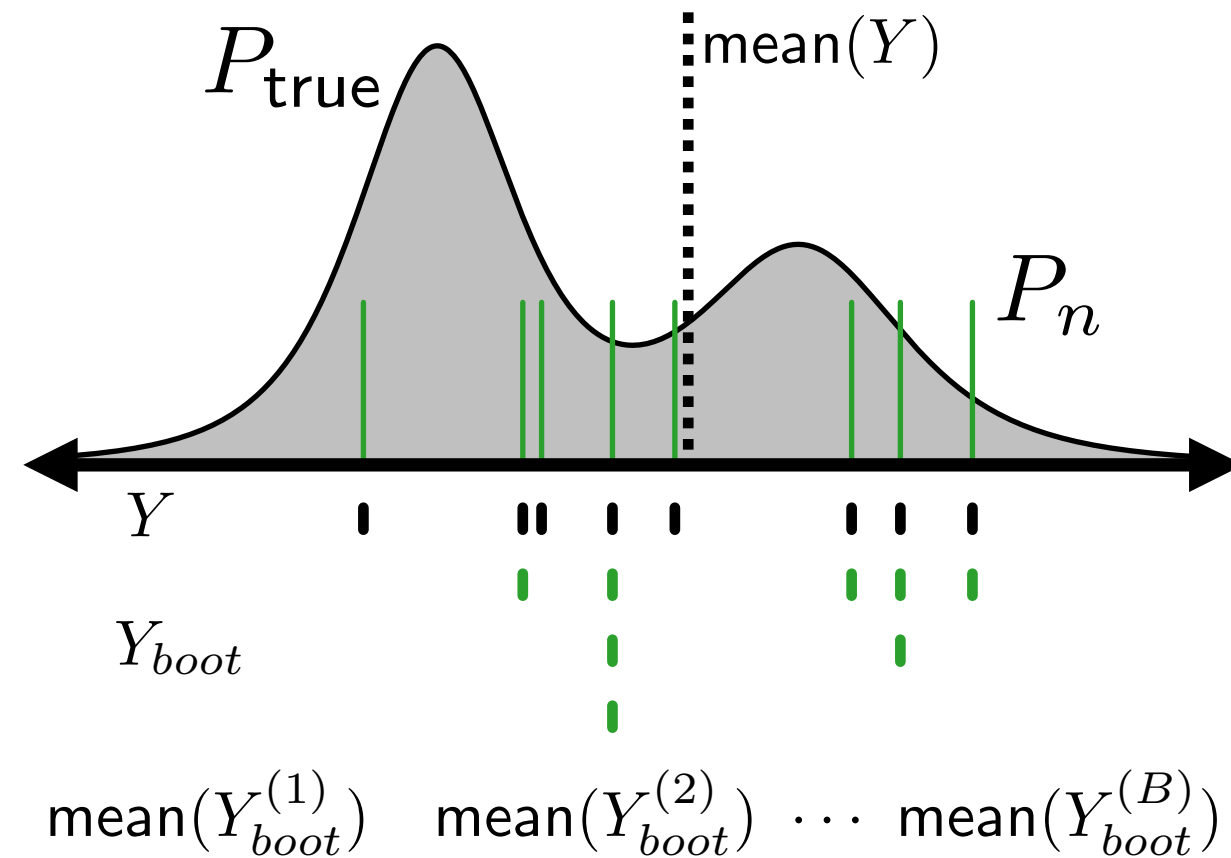


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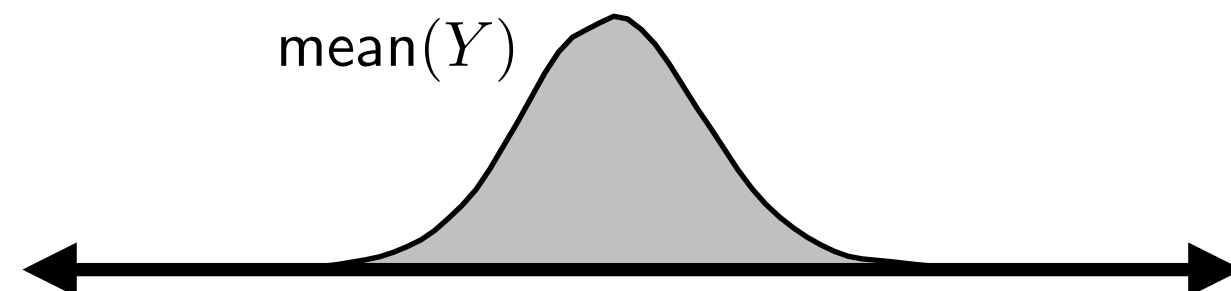


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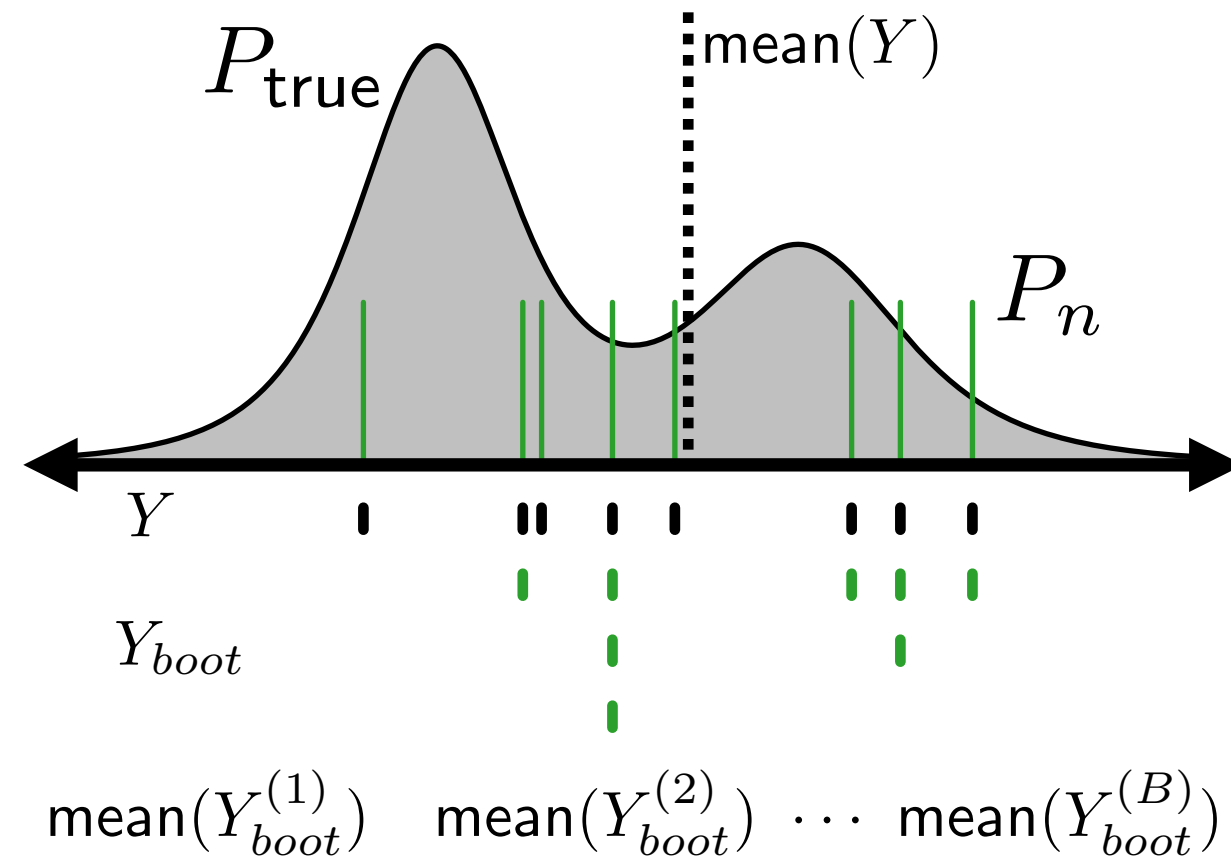


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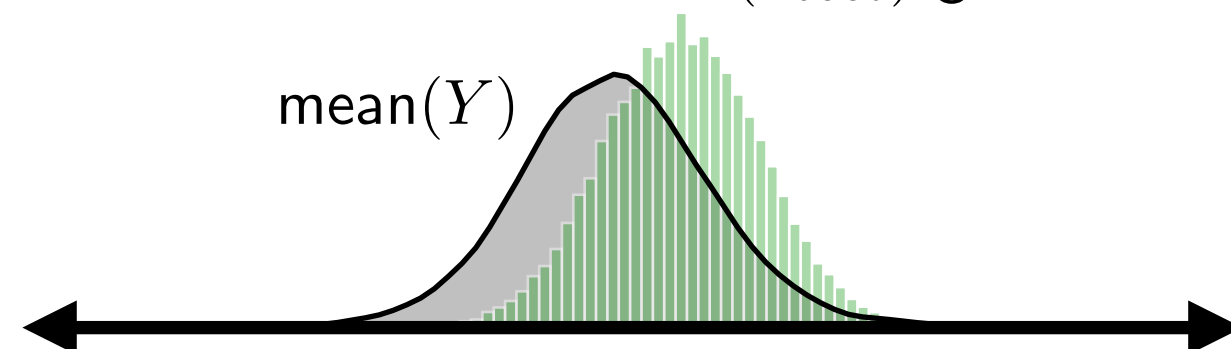
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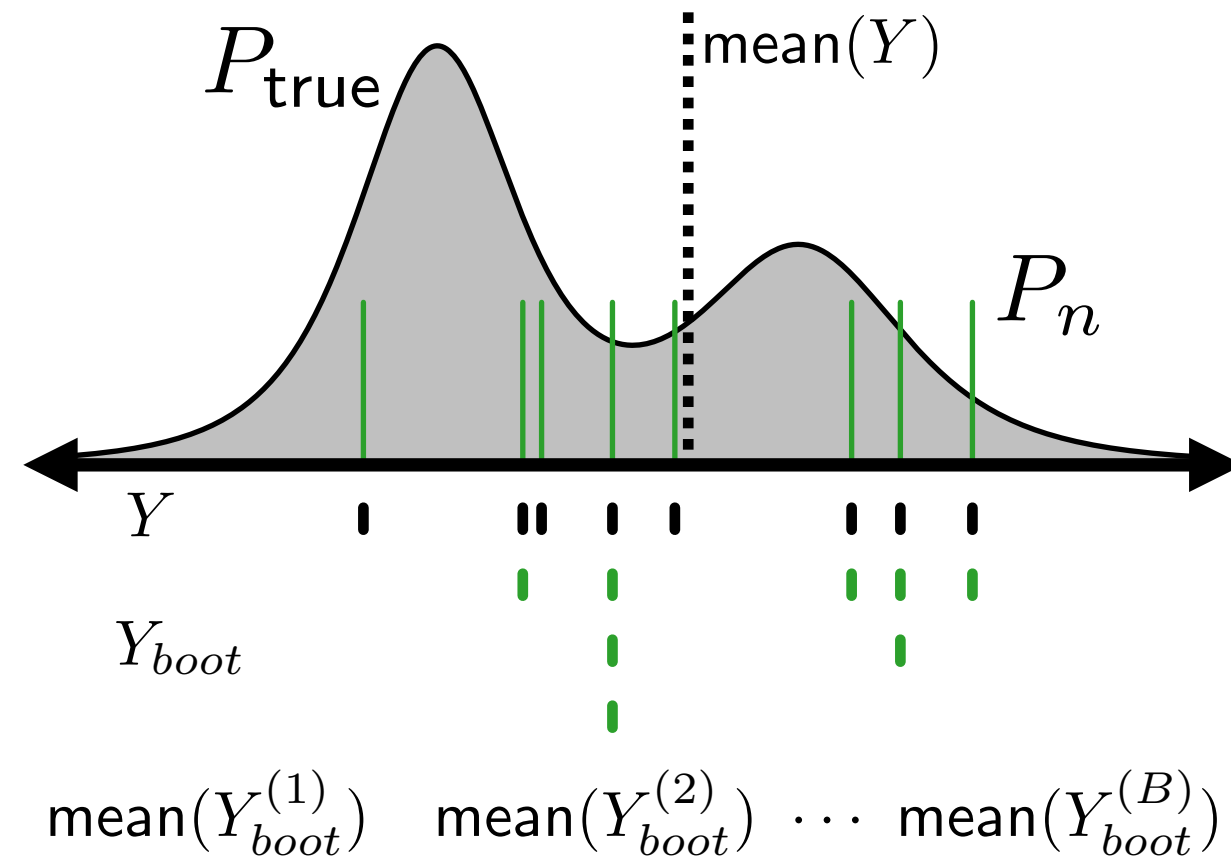
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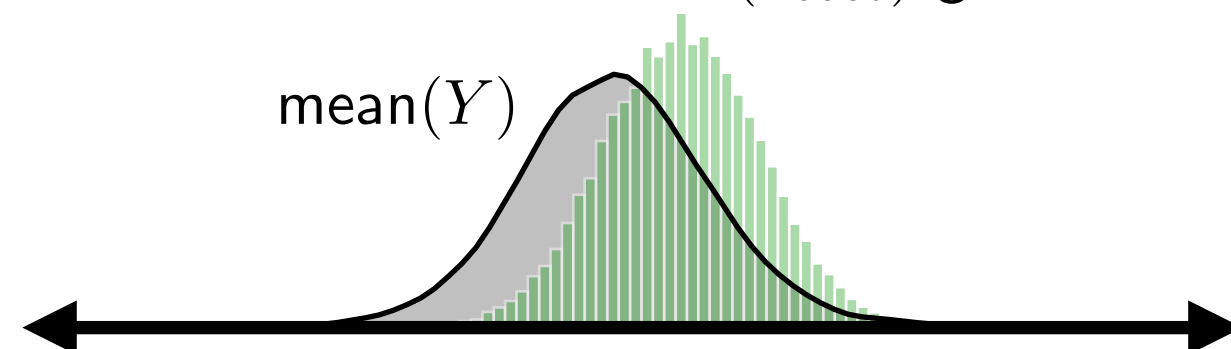
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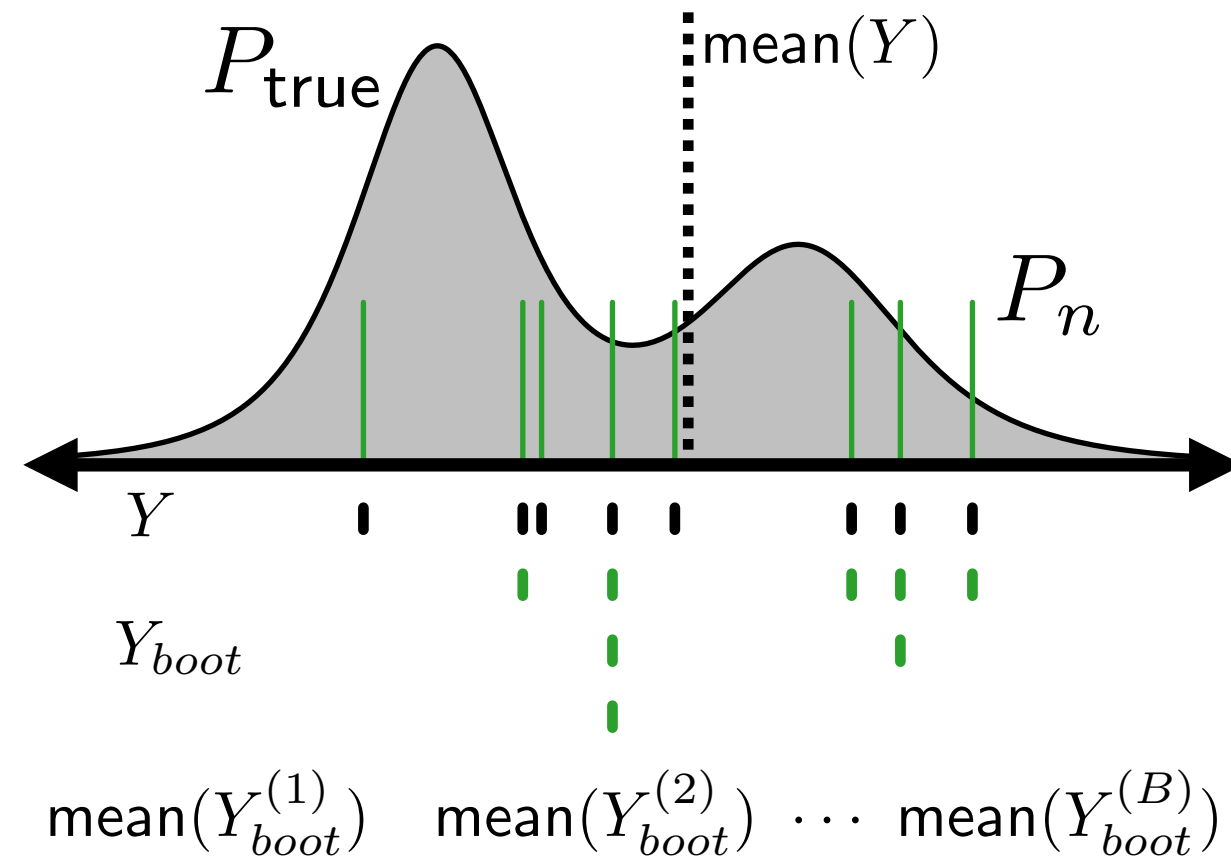
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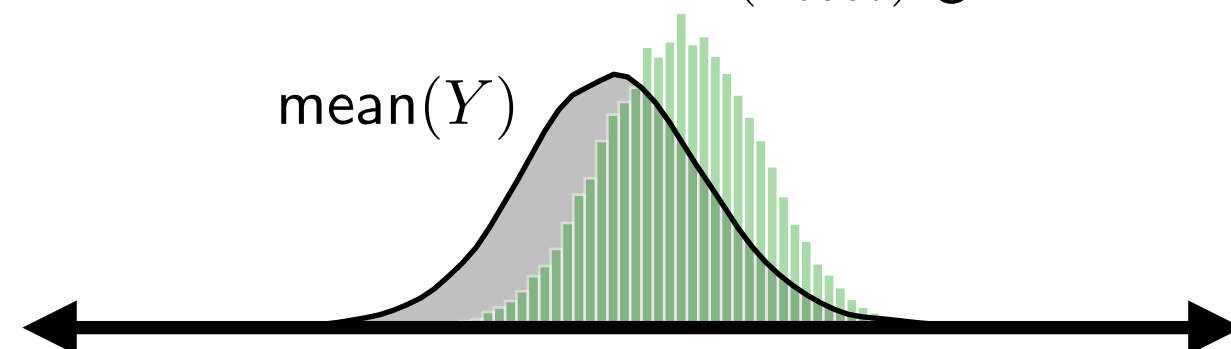
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- **Challenges:** B large (1,000-100,000), finite-sample properties

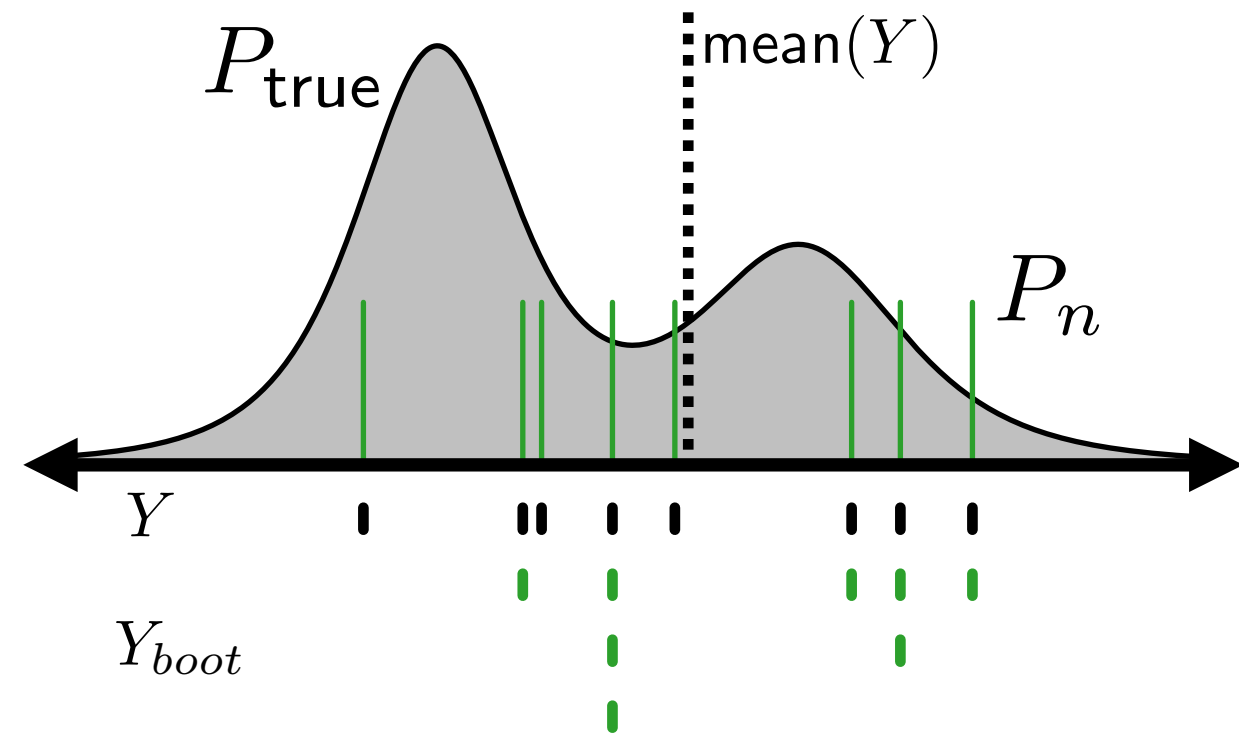


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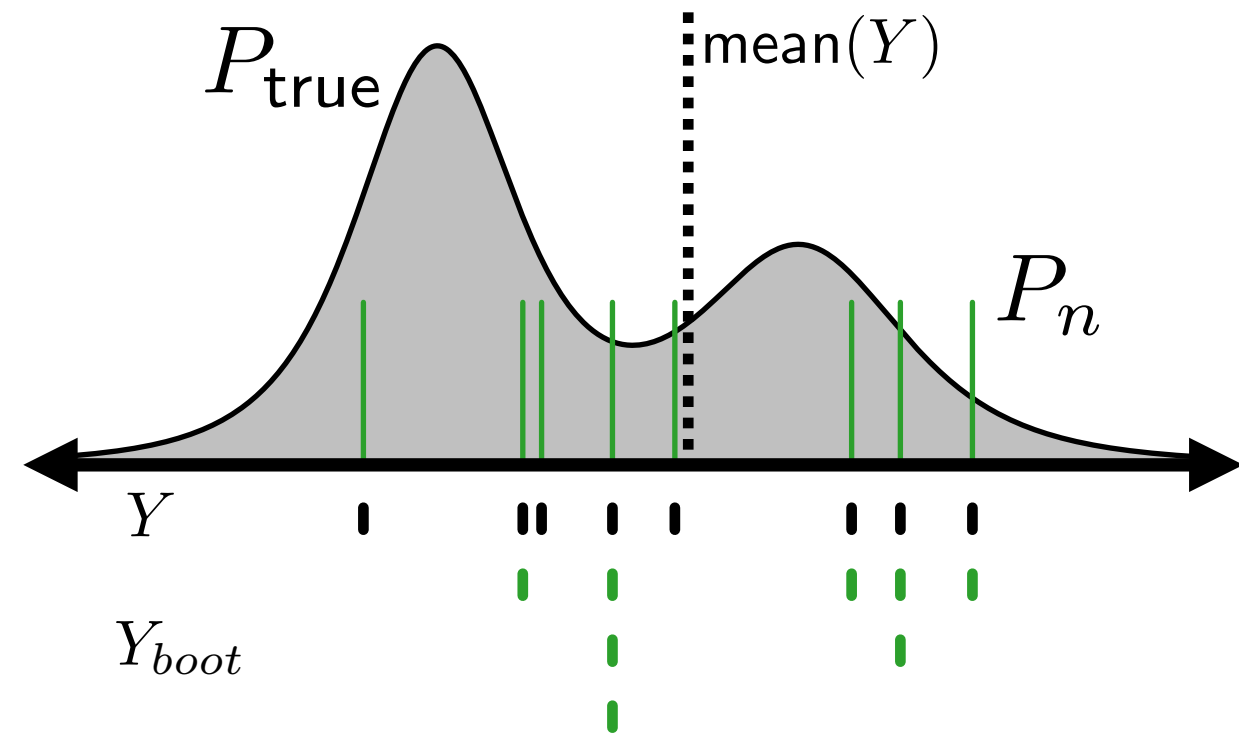


Bootstrapping Bayes

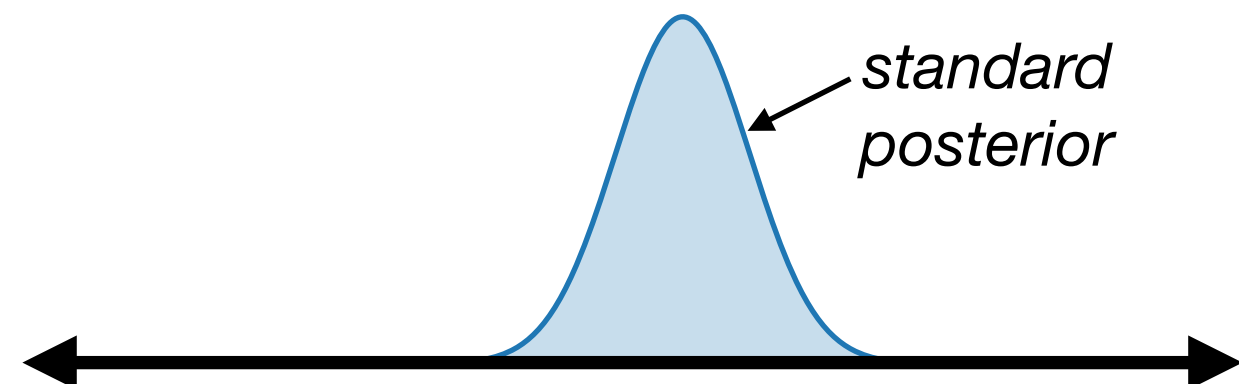


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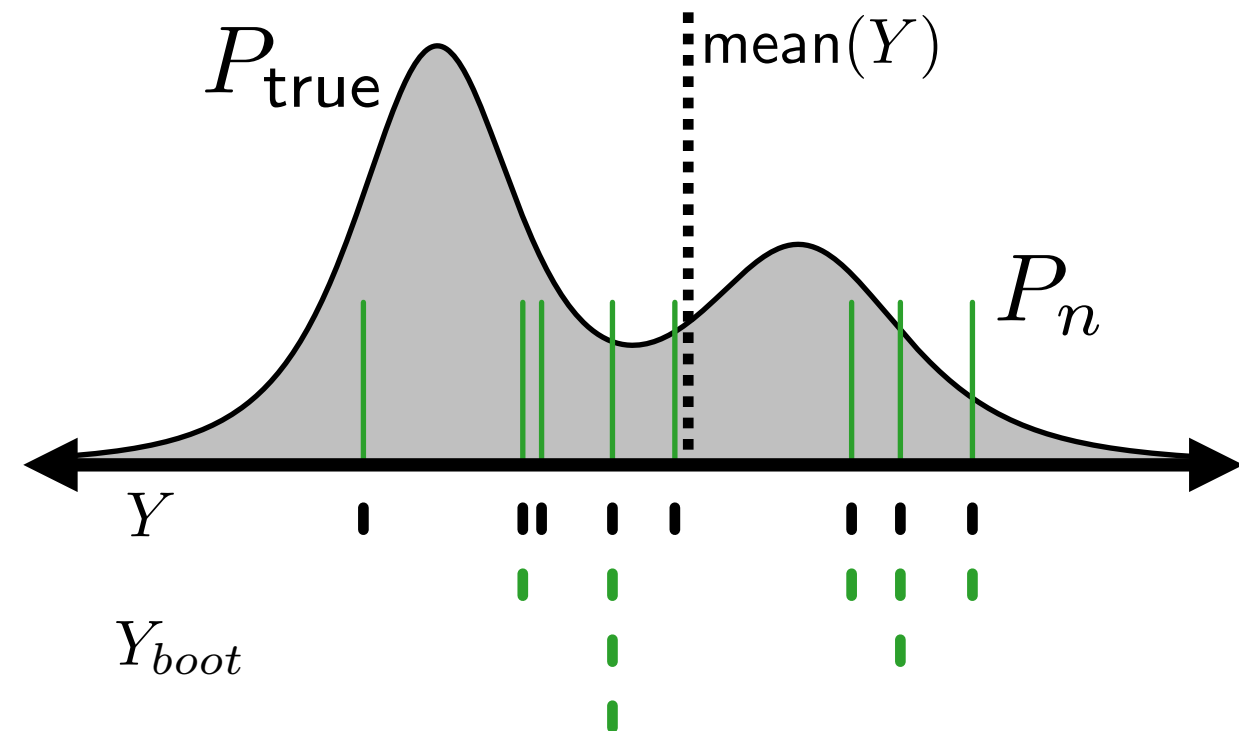
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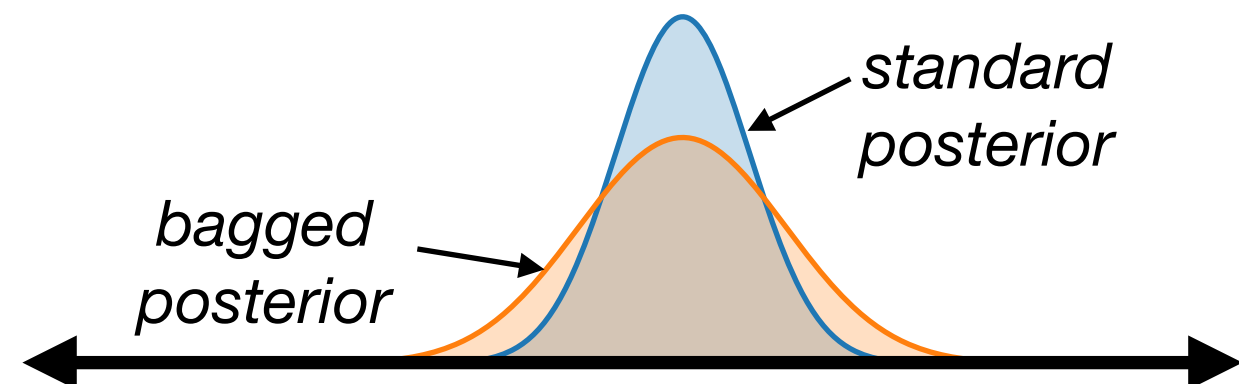
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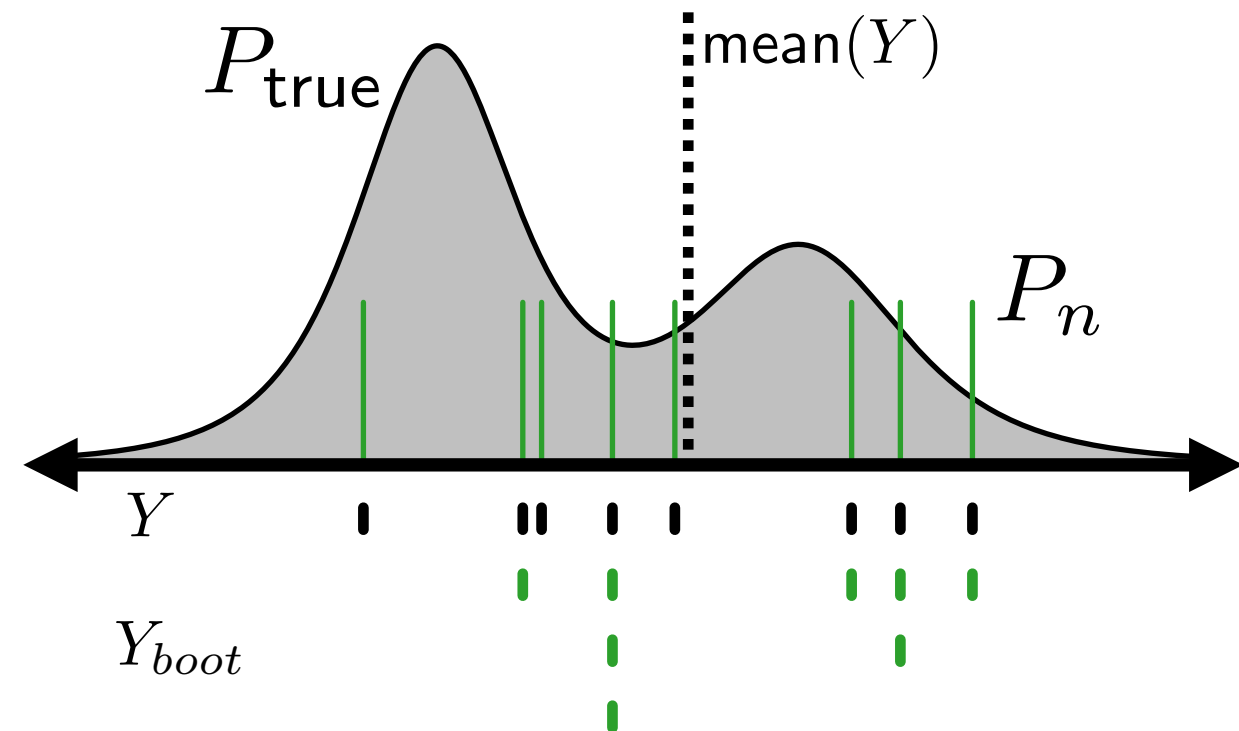


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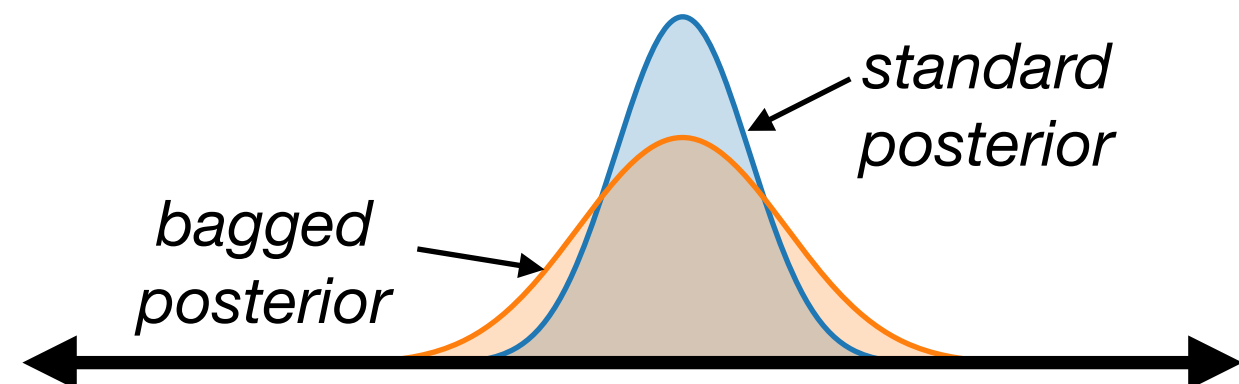
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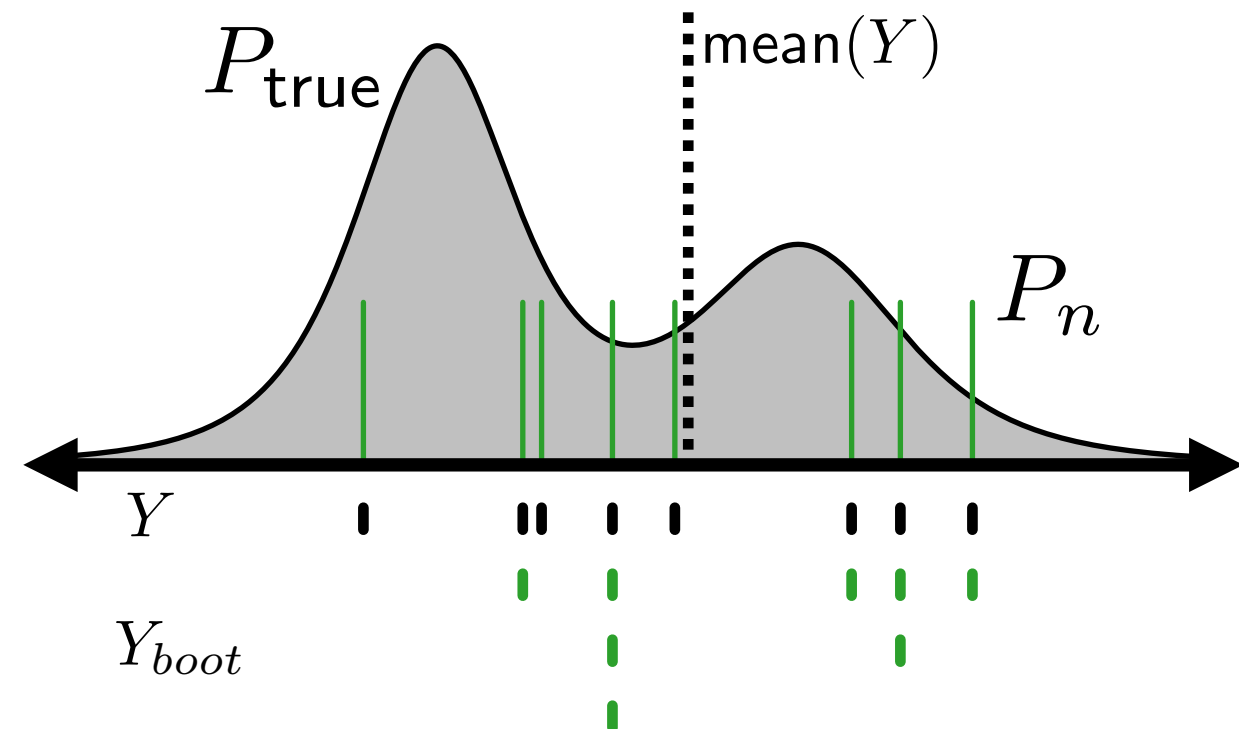


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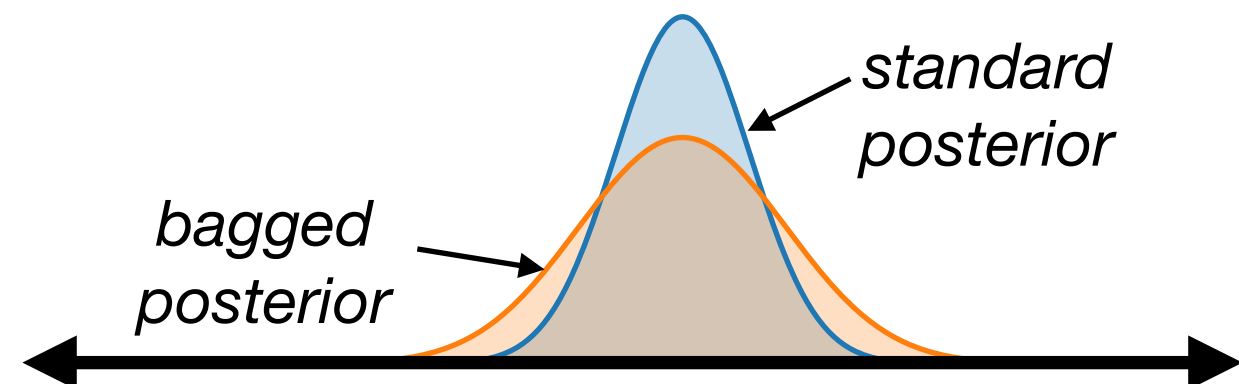
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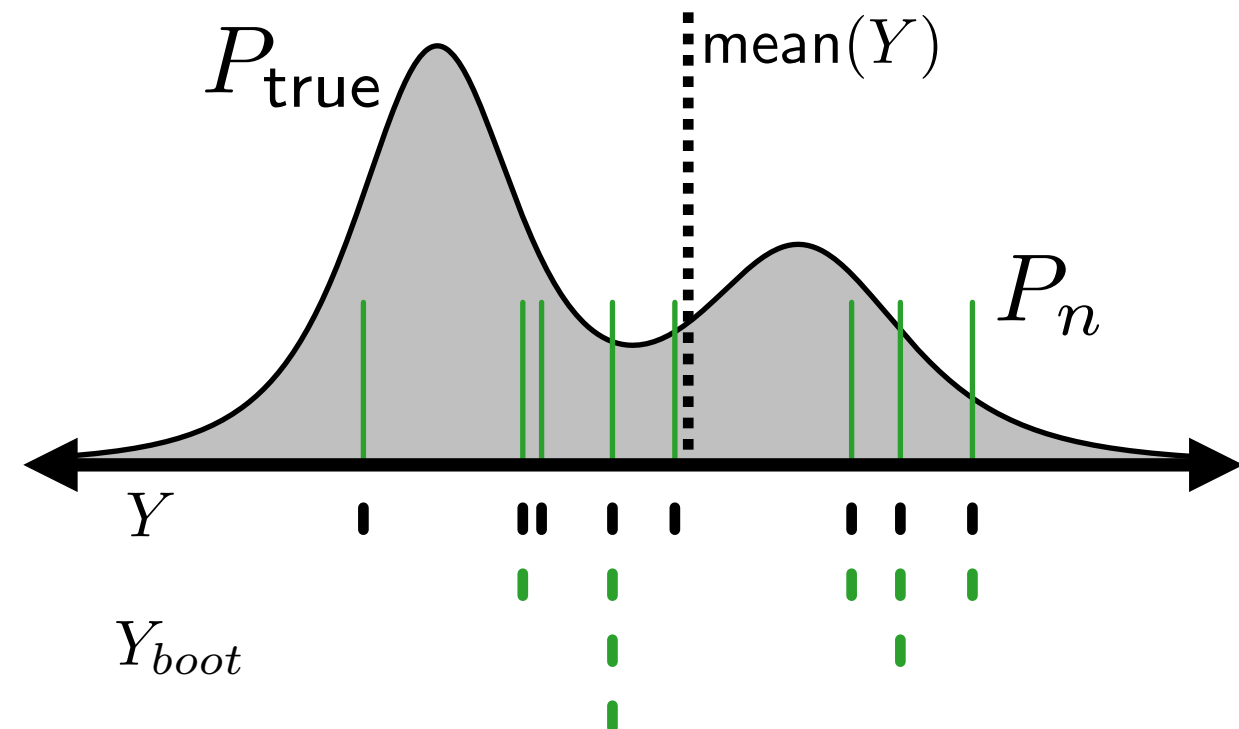


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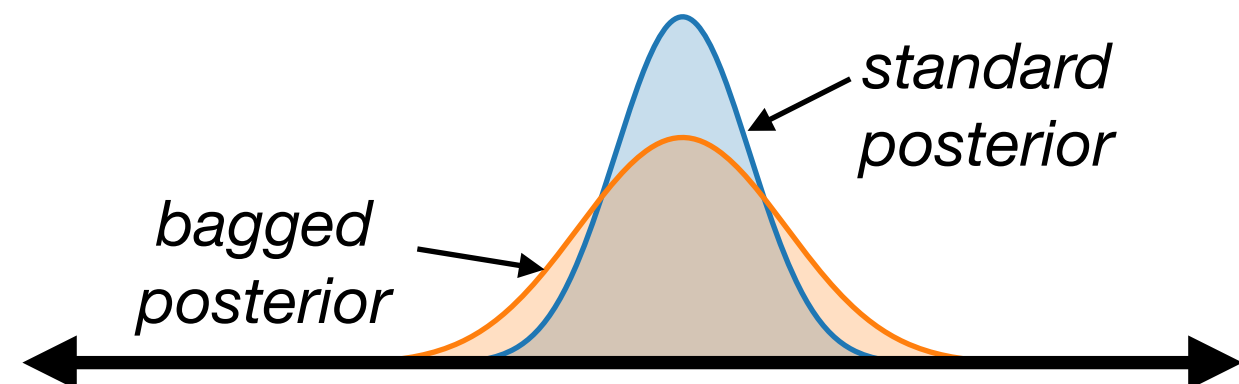
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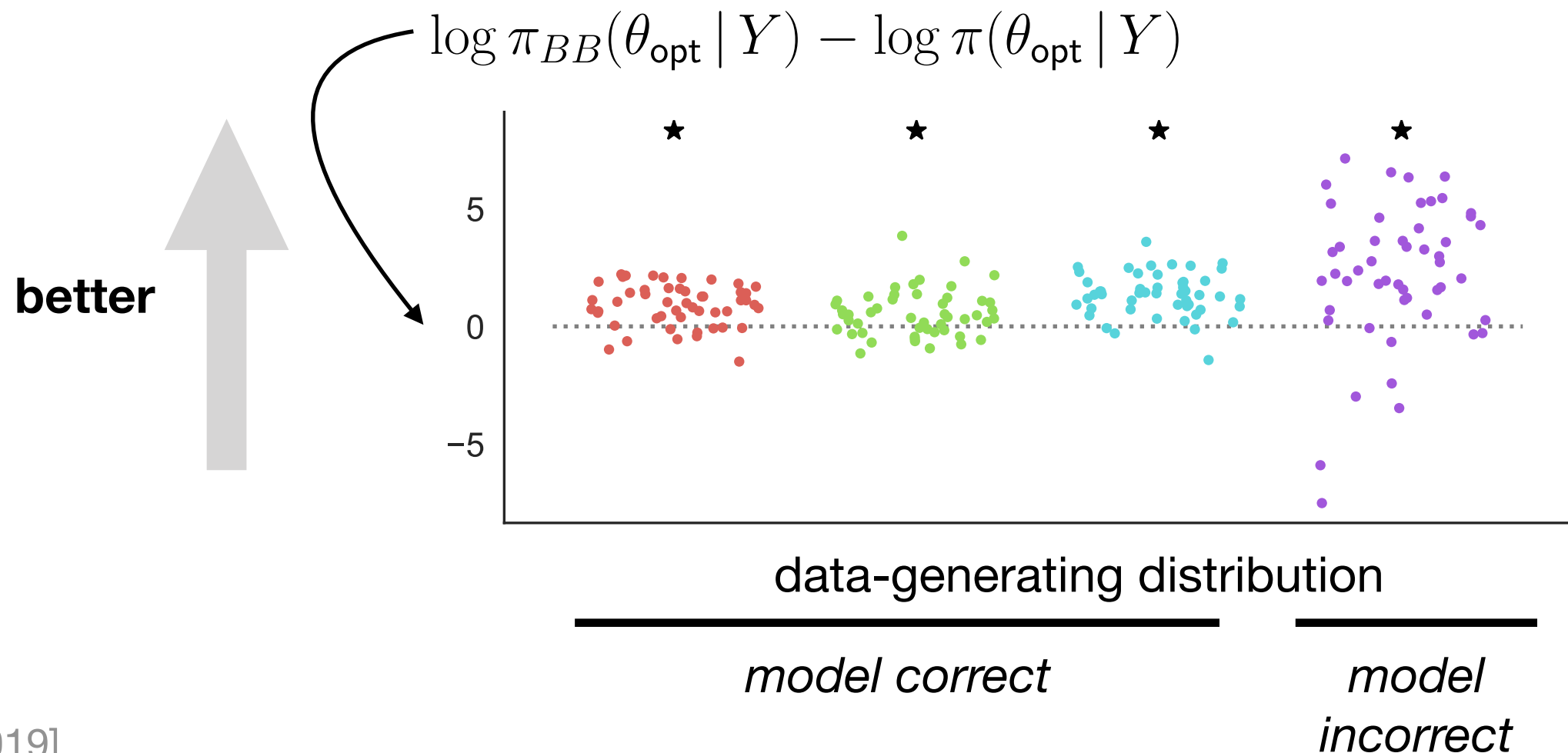
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$$\log \pi_{BB}(\theta_{\text{opt}} | Y) - \log \pi(\theta_{\text{opt}} | Y)$$

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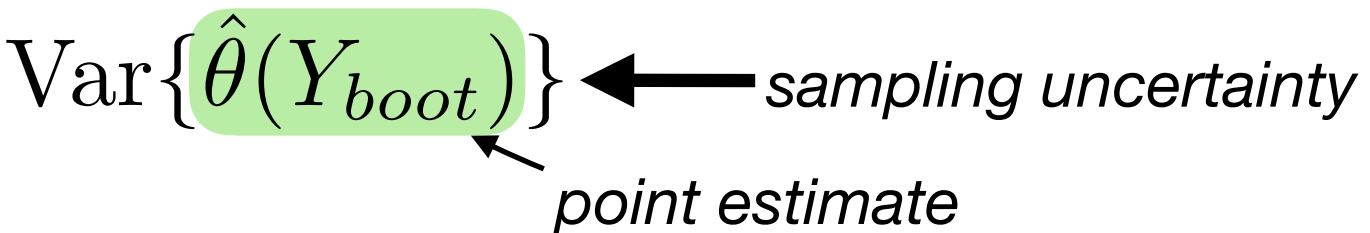
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 *point estimate*

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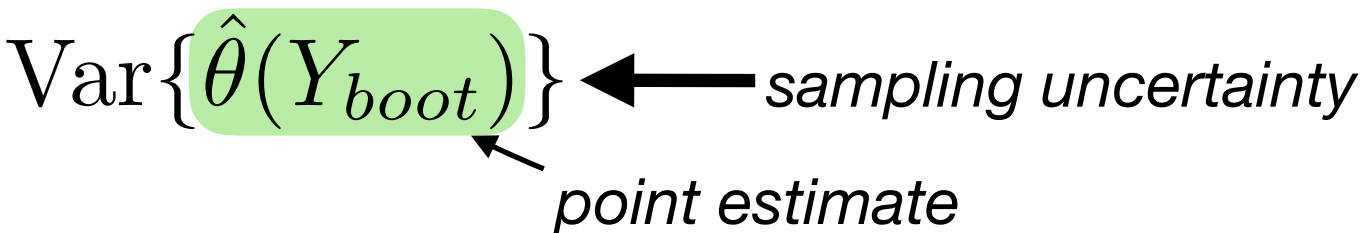
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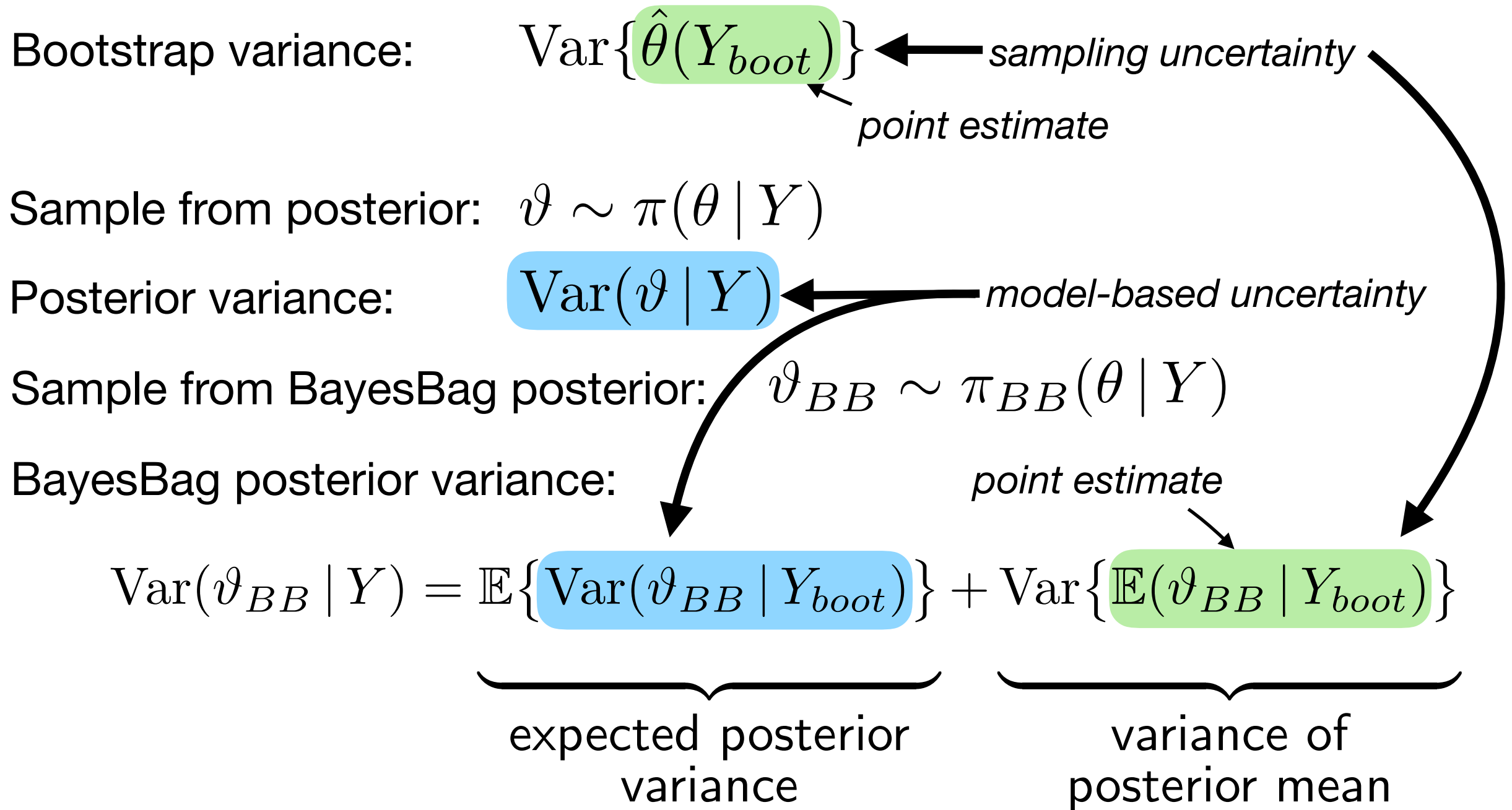
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- Summarizing the previous slide...

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Posterior variance = *model-based uncertainty*

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Posterior variance = *model-based uncertainty*

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Posterior variance = *model-based uncertainty*

Bootstrap variance = *sampling-based uncertainty*

BayesBag variance = *model-based + sampling-based uncertainty*

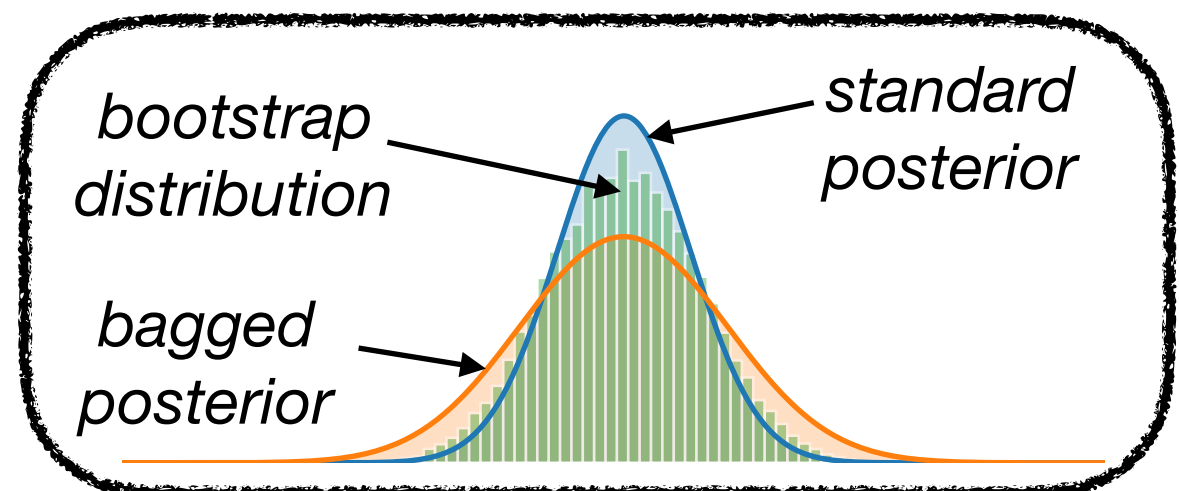
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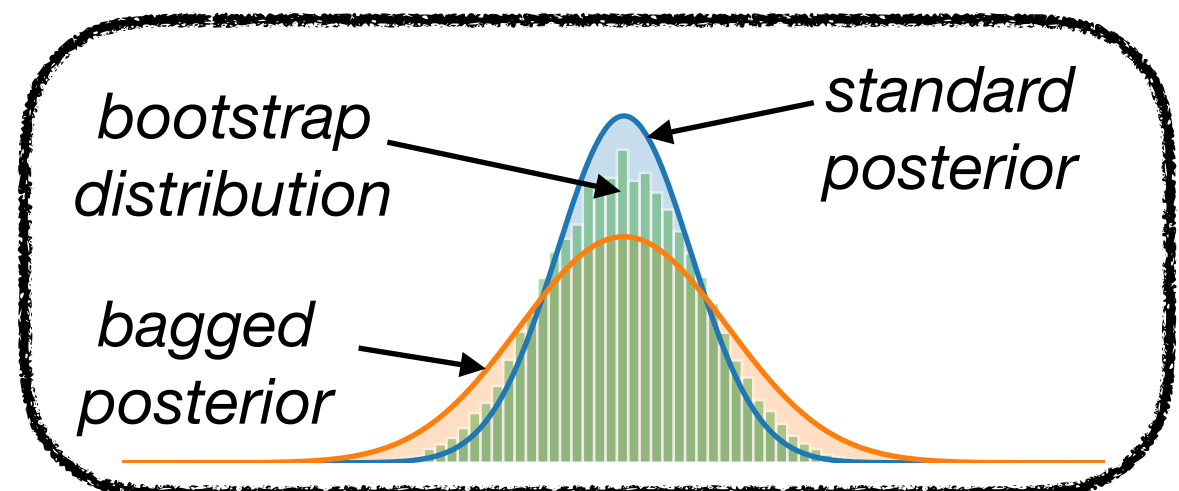
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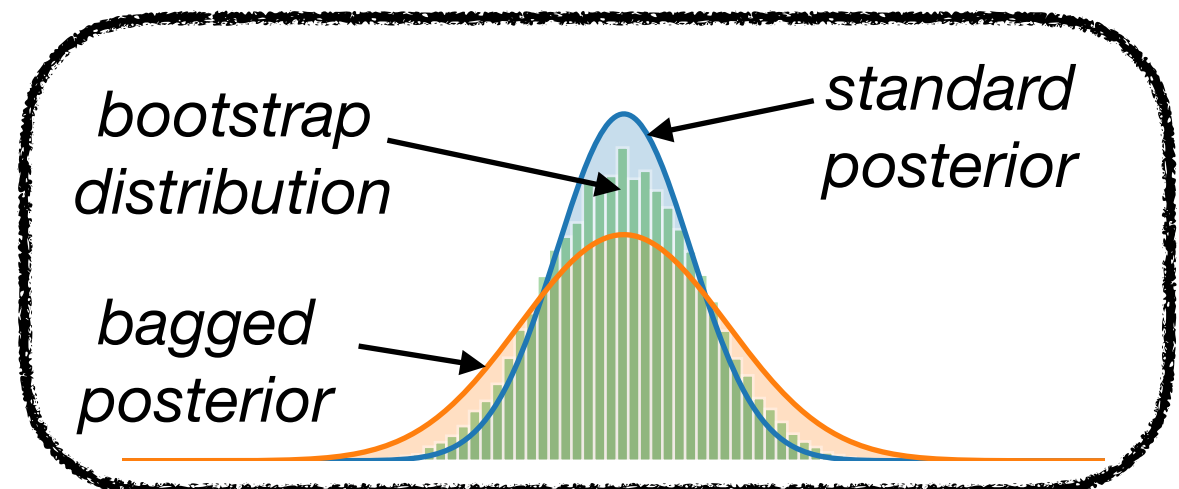
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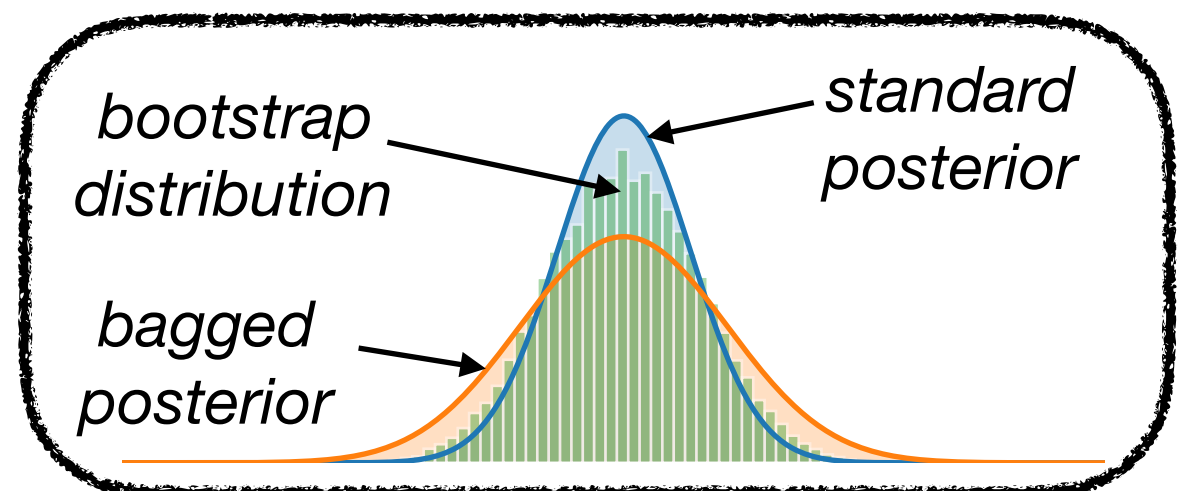
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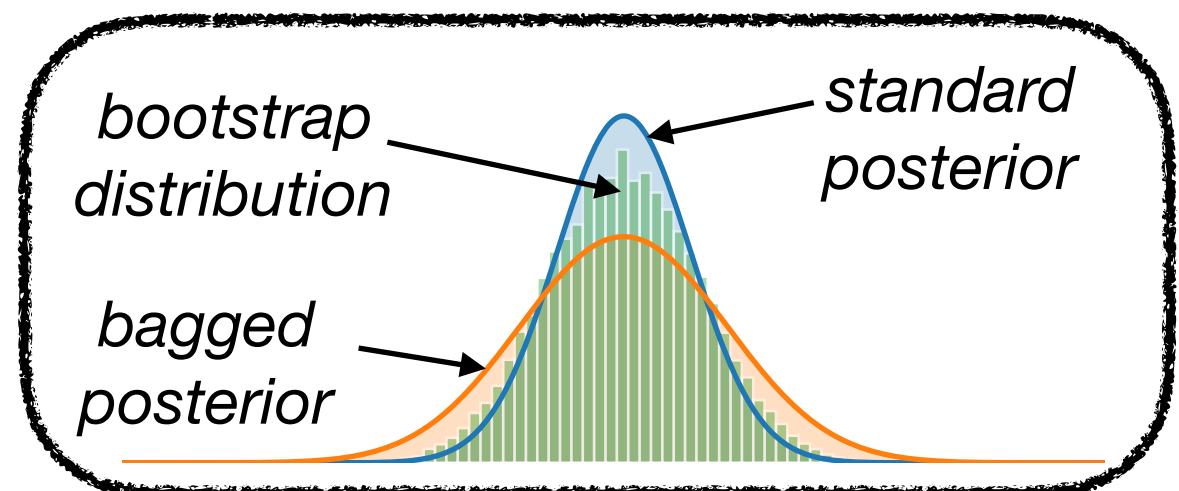
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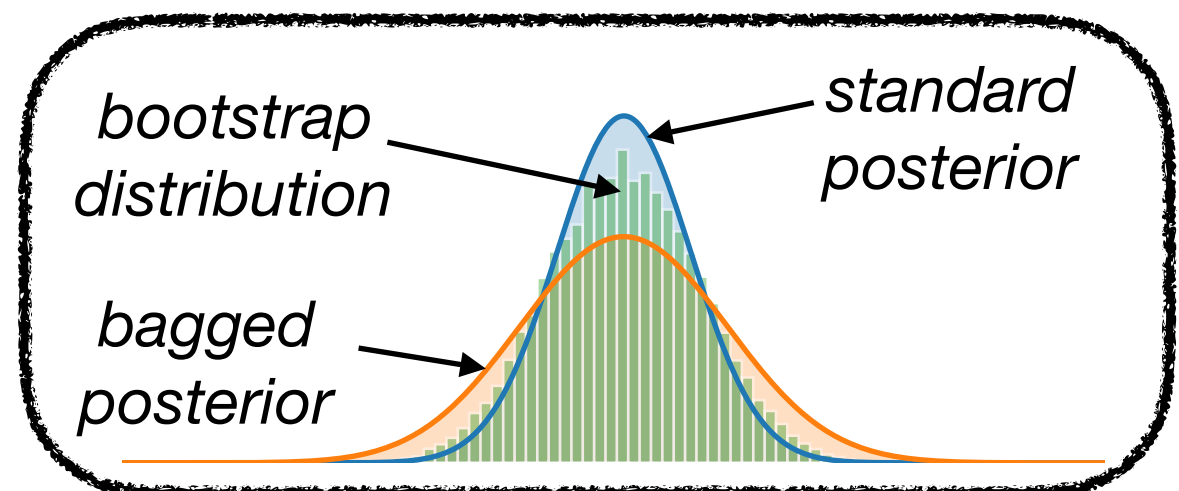
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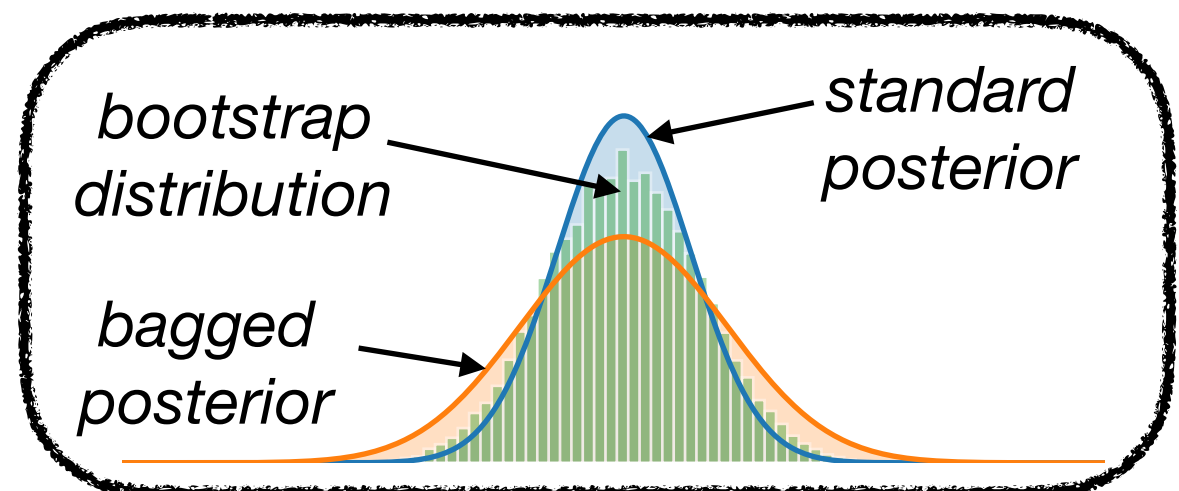
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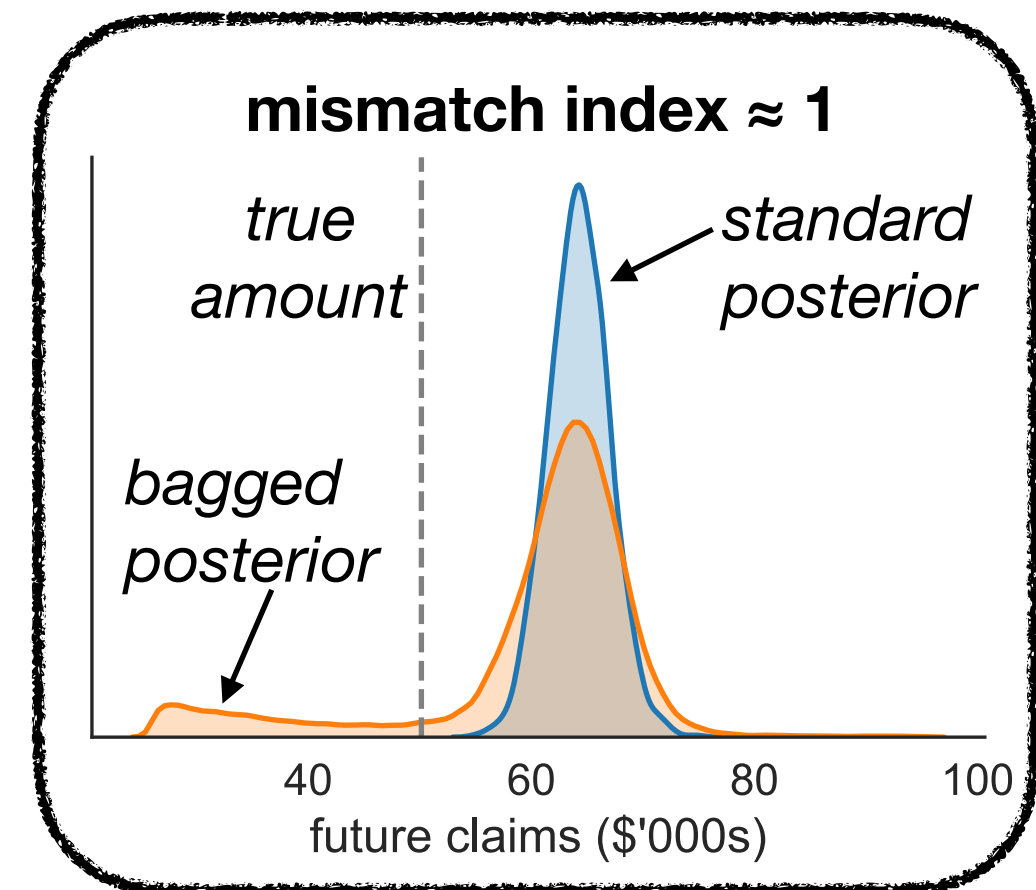
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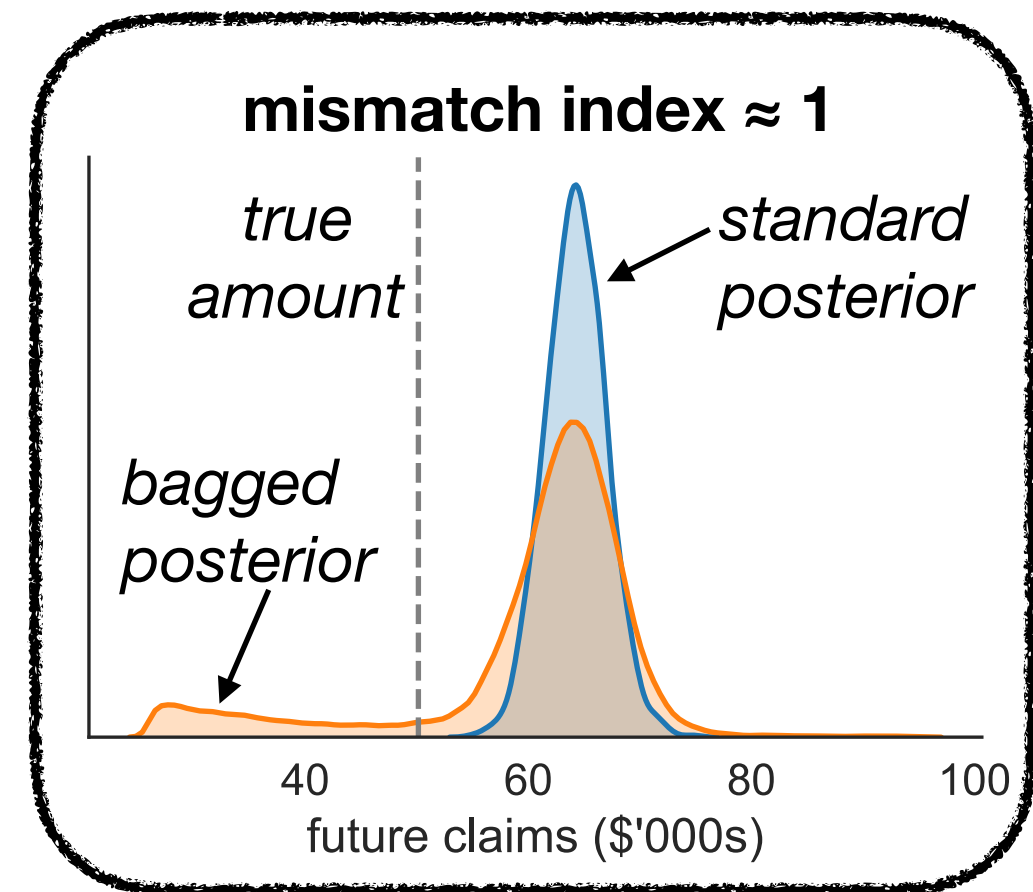


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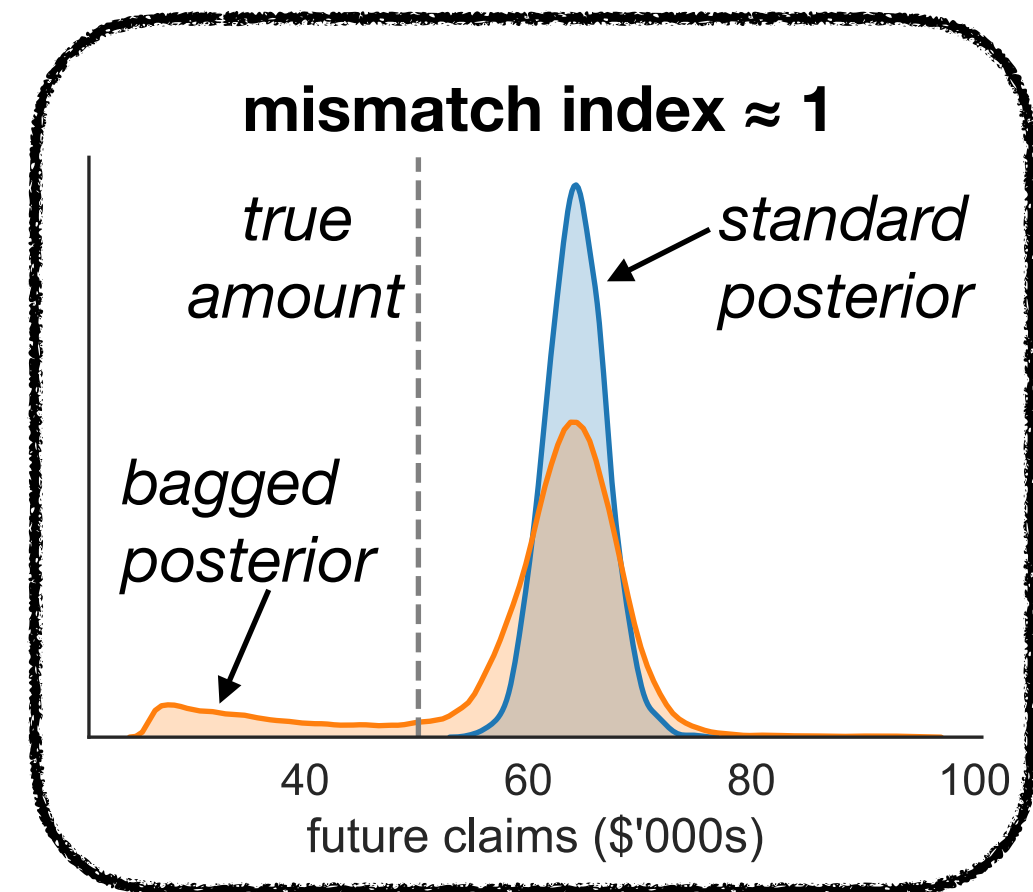
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- 4) output bagged posterior computed in step 2

Agenda

- BayesBag for parameter inference (and prediction)
- BayesBag theory and methodology
- ➡ **BayesBag for model selection**

Bayesian model selection

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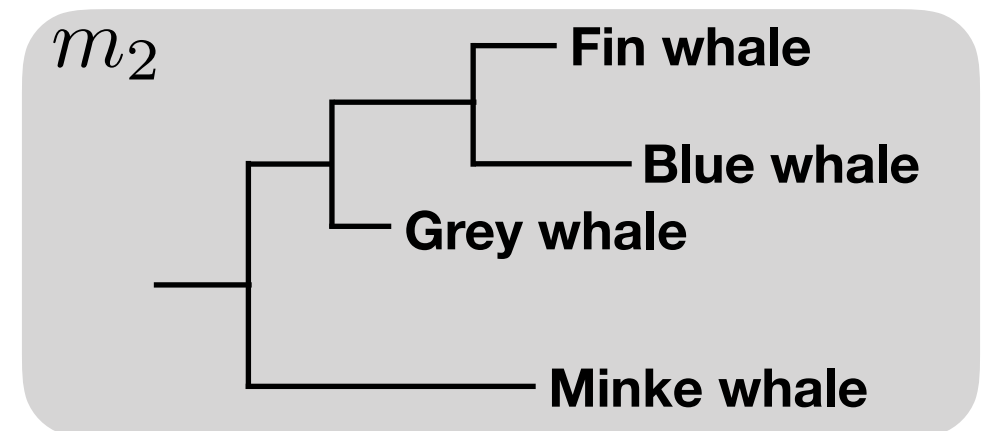
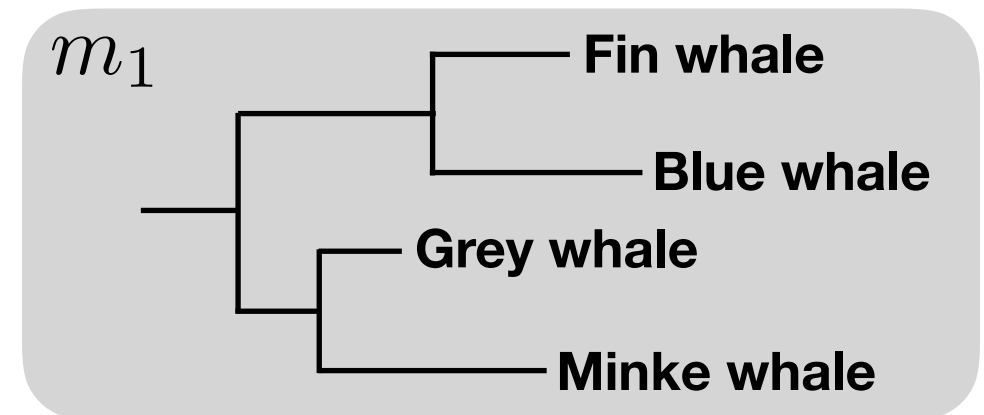
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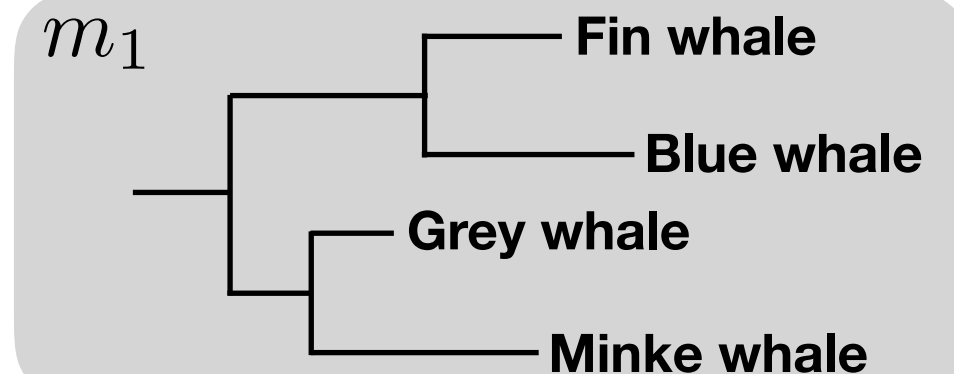


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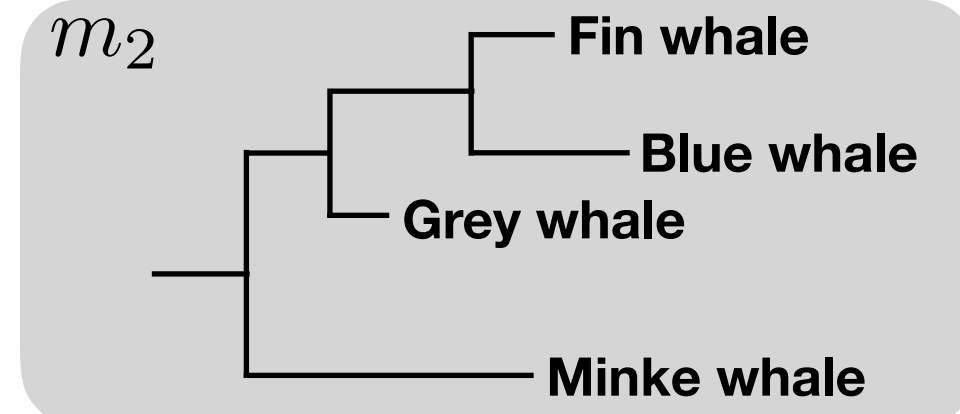
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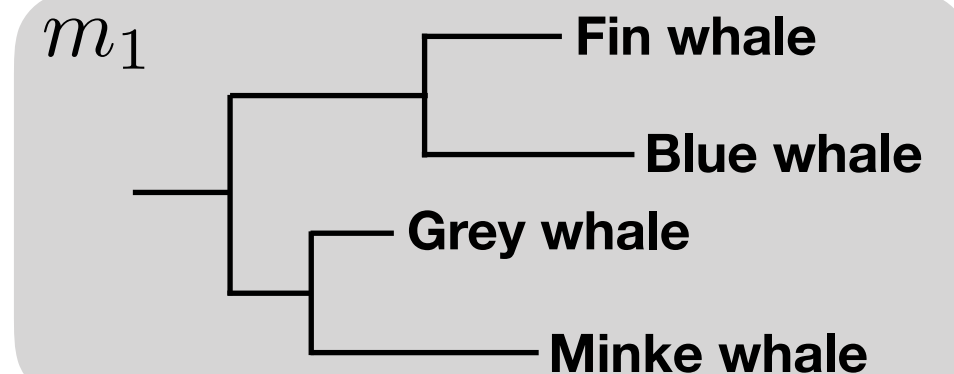


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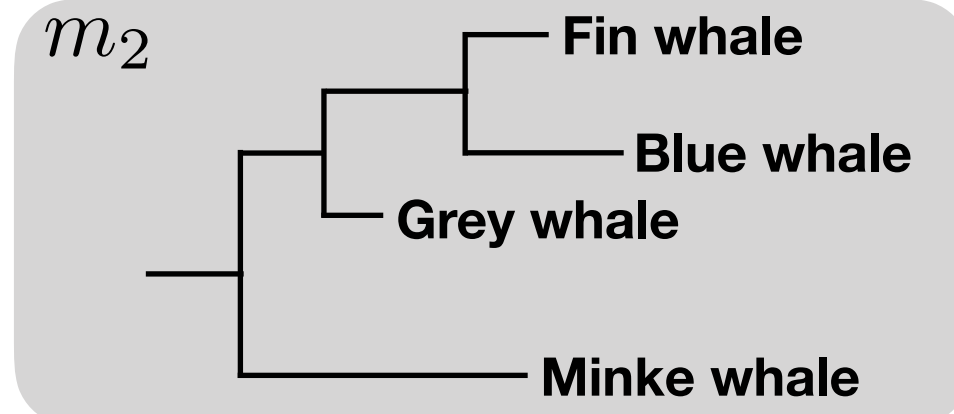
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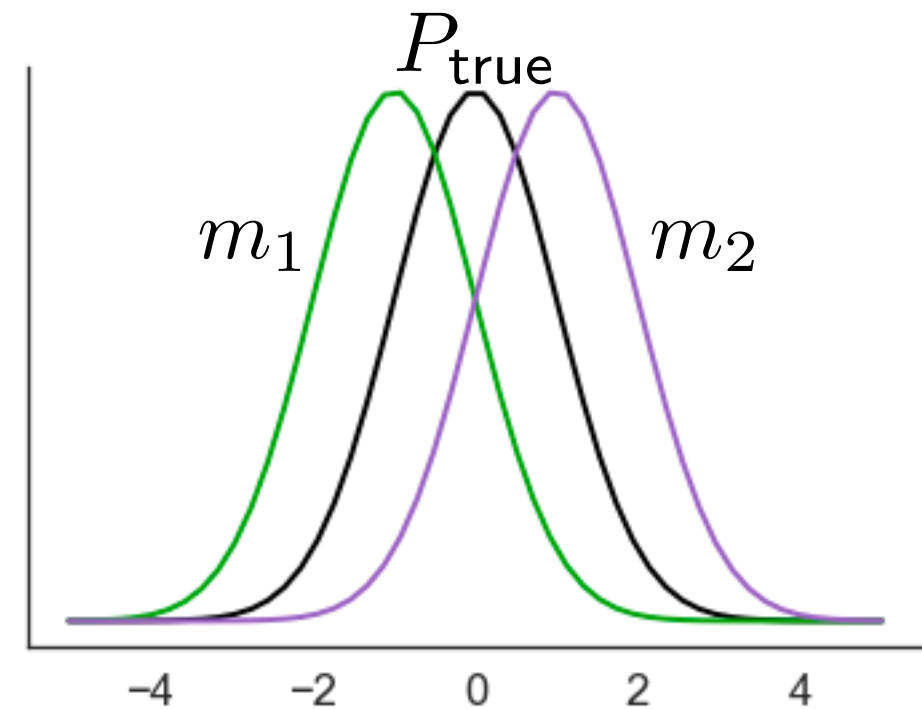


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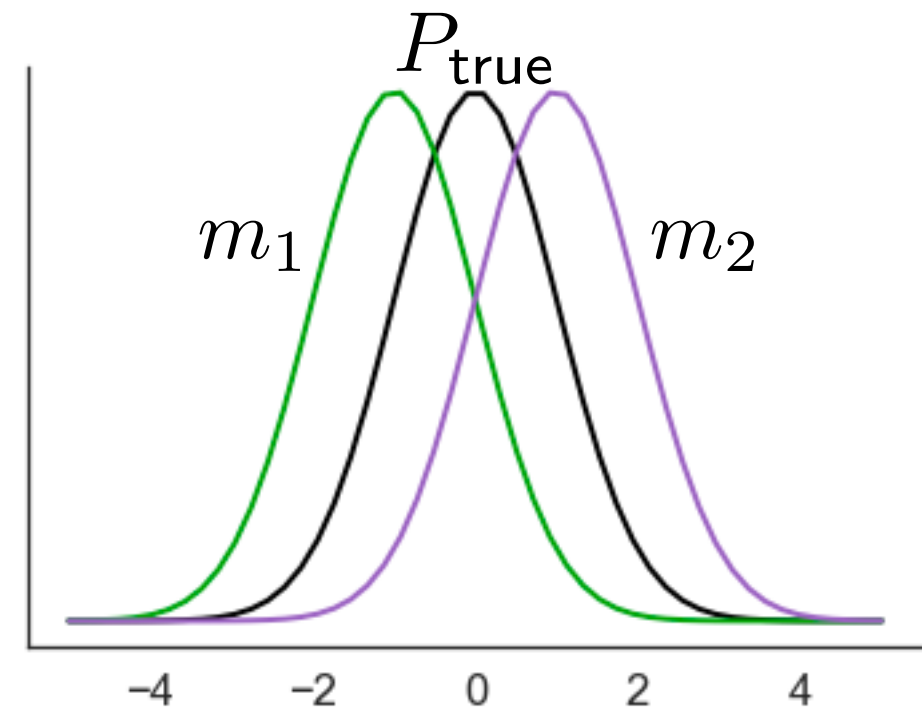


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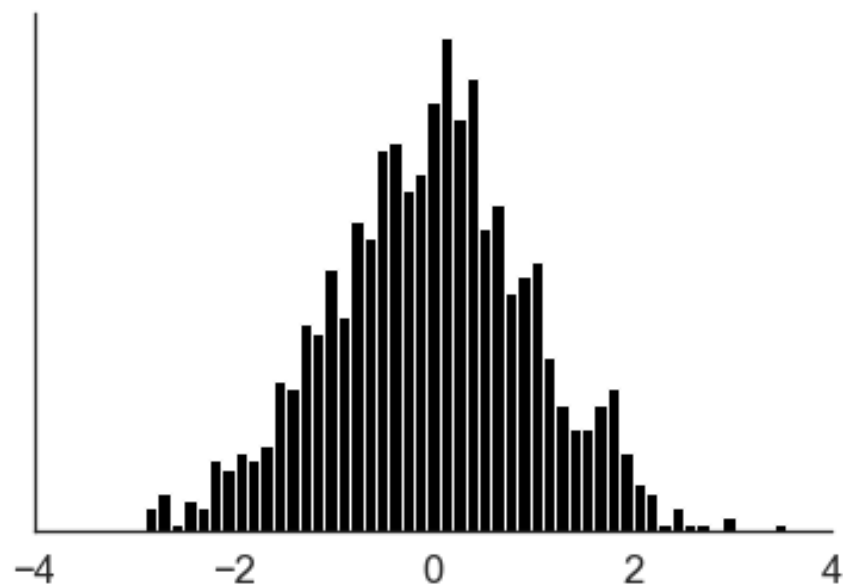
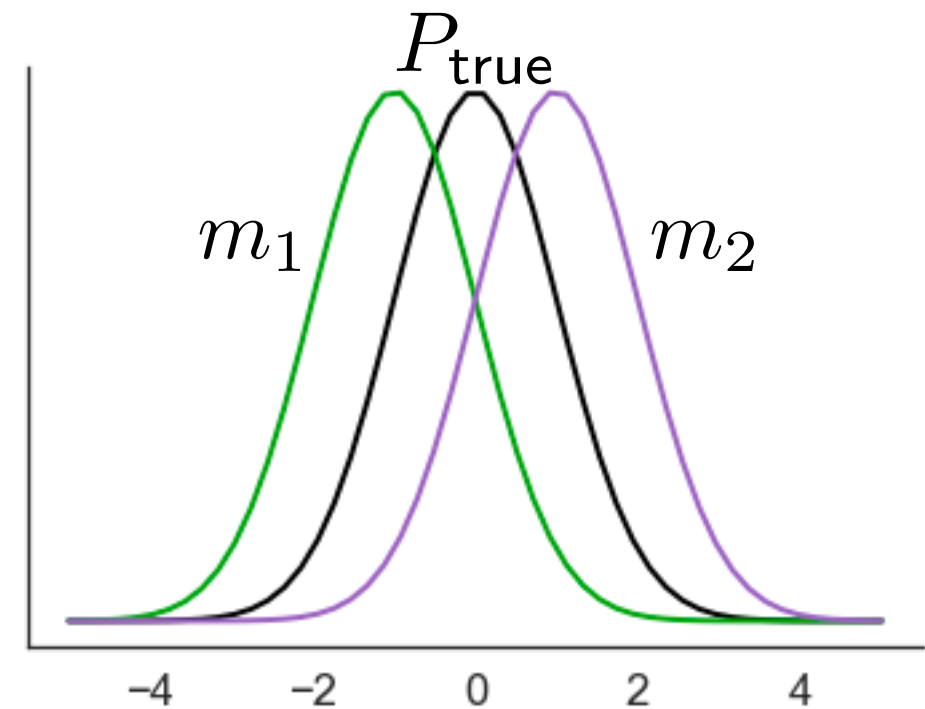
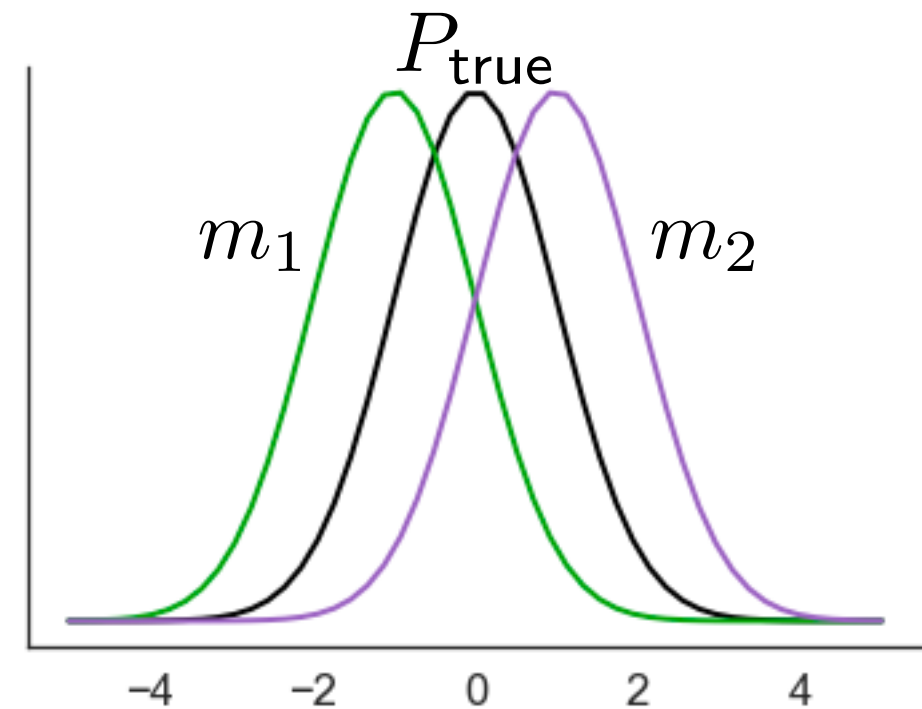


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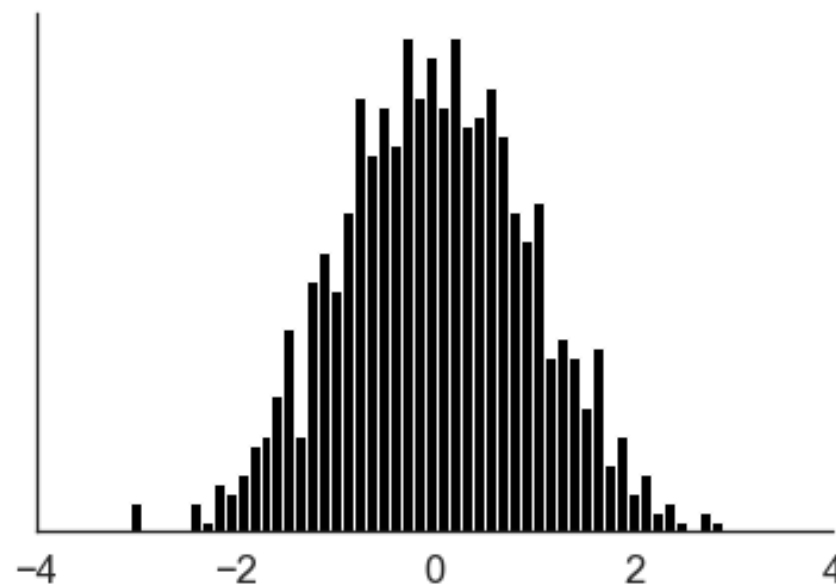
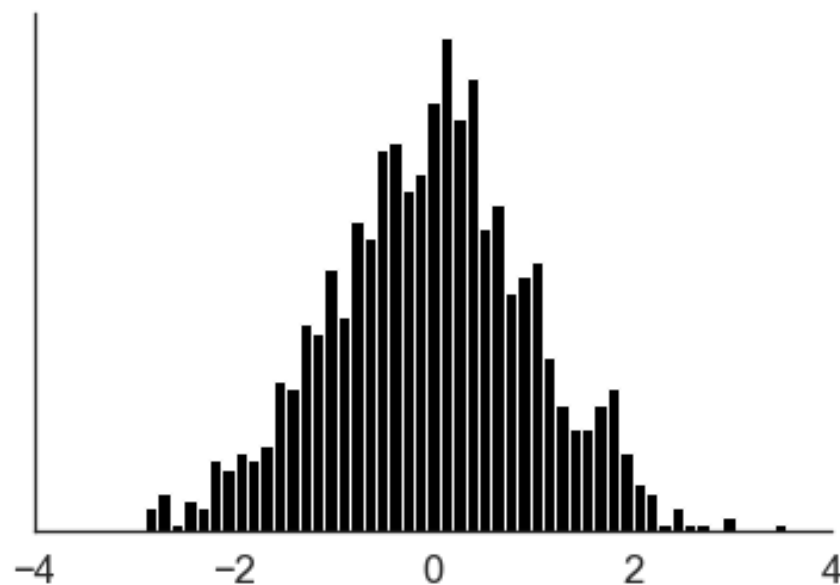
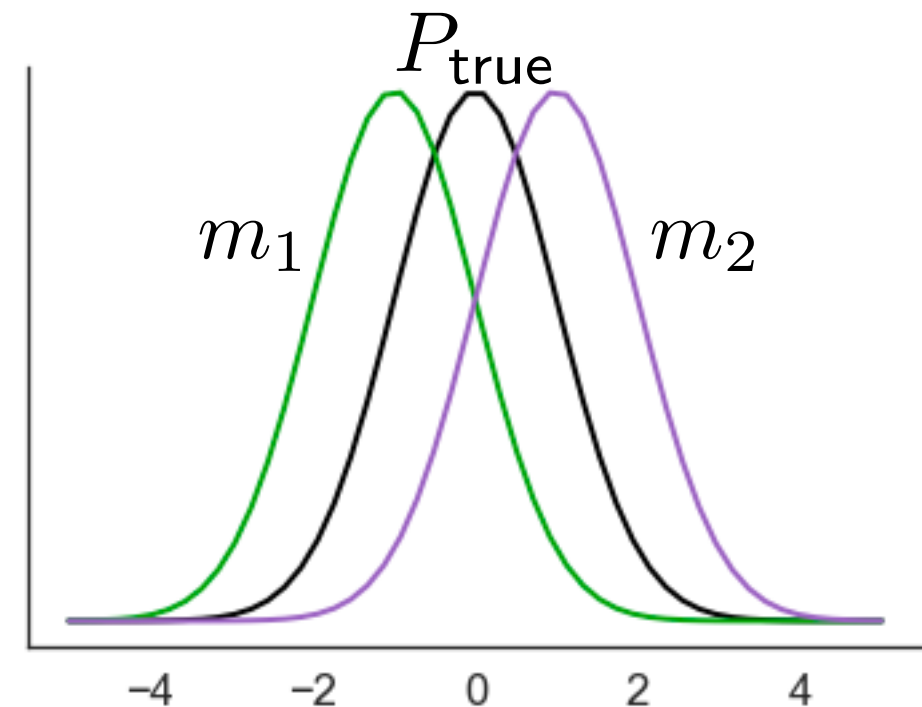


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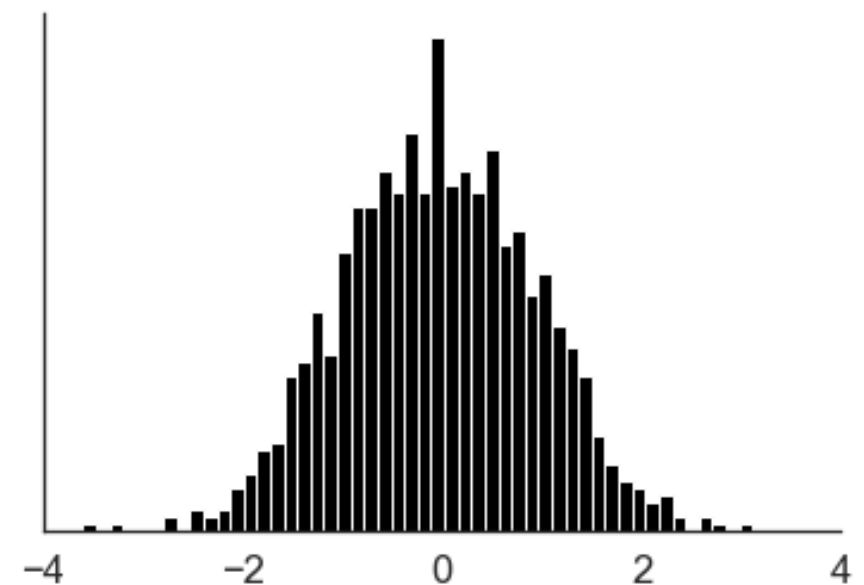
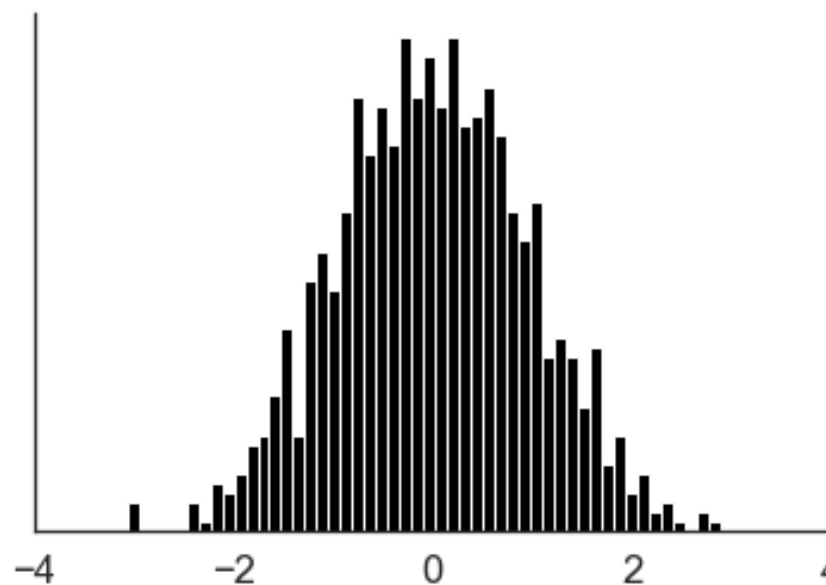
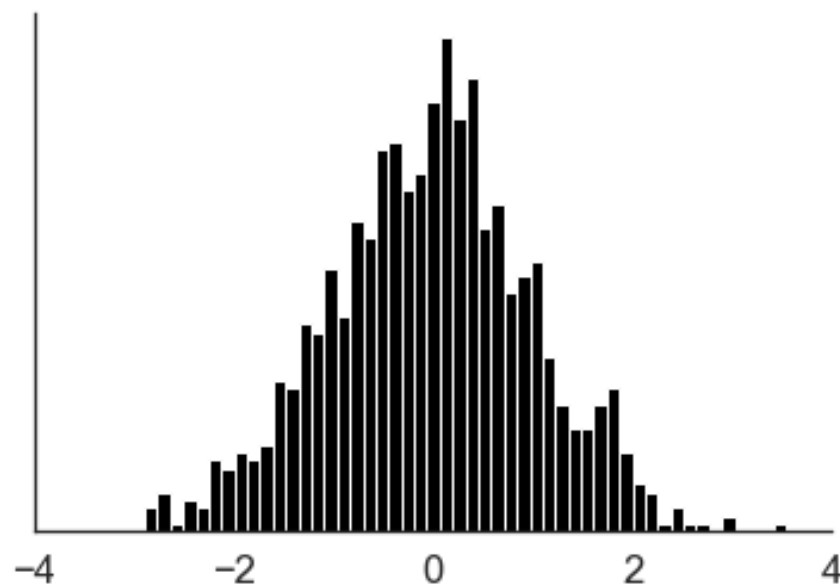
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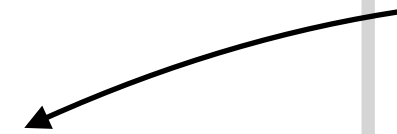
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$\pi_{BB}(m_1 | Y) \sim \text{Uniform}[0,1]$

All posterior mass
on a single,
arbitrary model

bagged
posterior mass
more evenly
distributed

Reproducible phylogenetic inference

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- **Goal:** infer phylogeny of 13 whale species from mitochondrial coding DNA

all

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Blue	CACCCCCCGTACTAT...TGAGTCCGAATTGGAA
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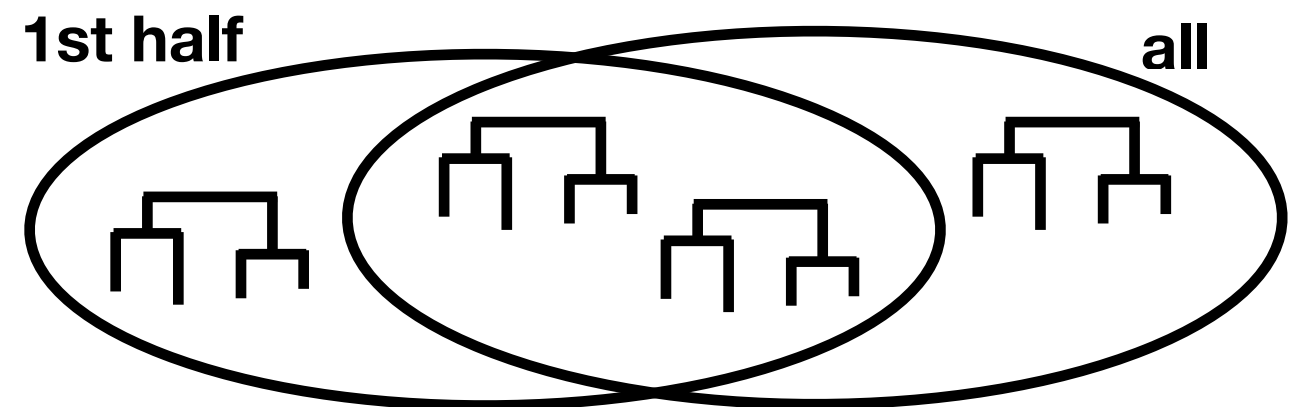
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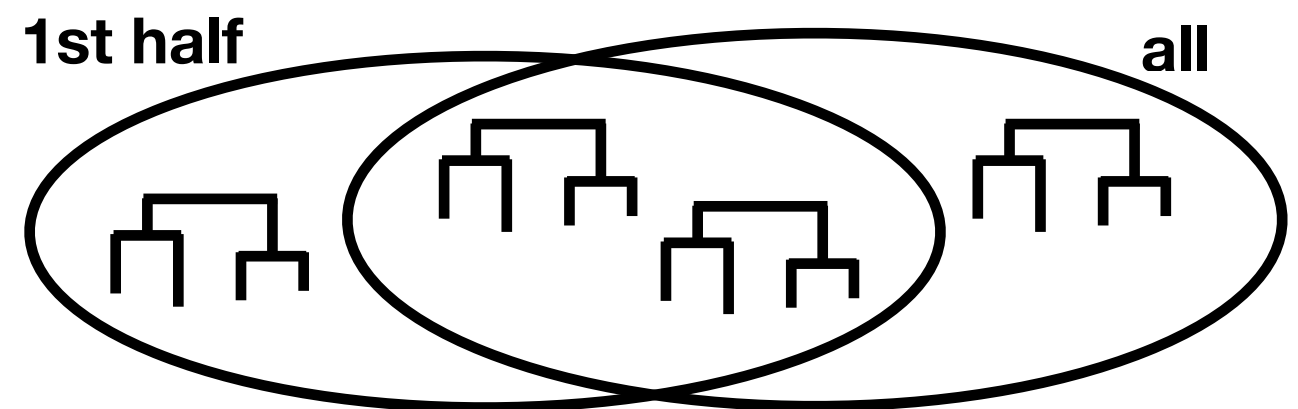
all	1st half	2nd half
Minke	GACCCGAACGTAATAA	ATCCGTTCCCATACTC
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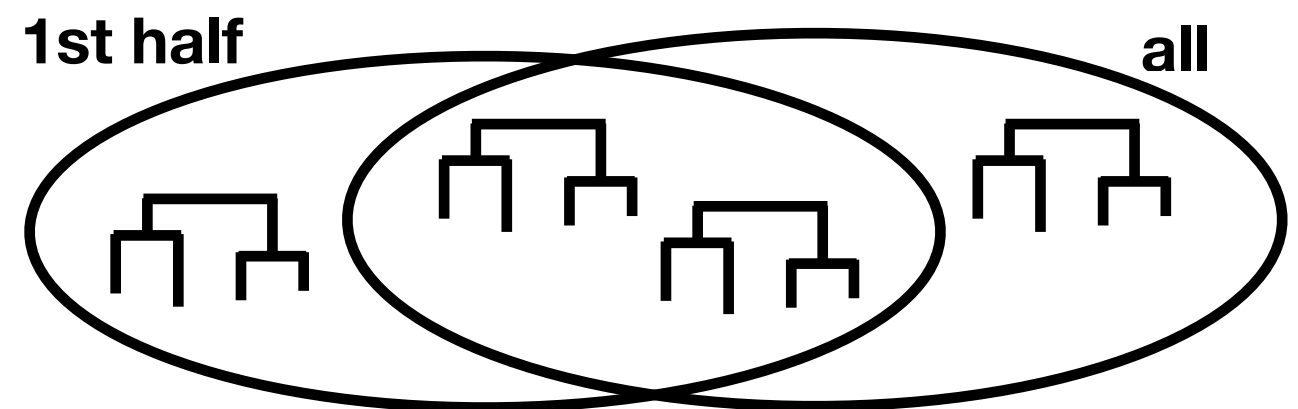
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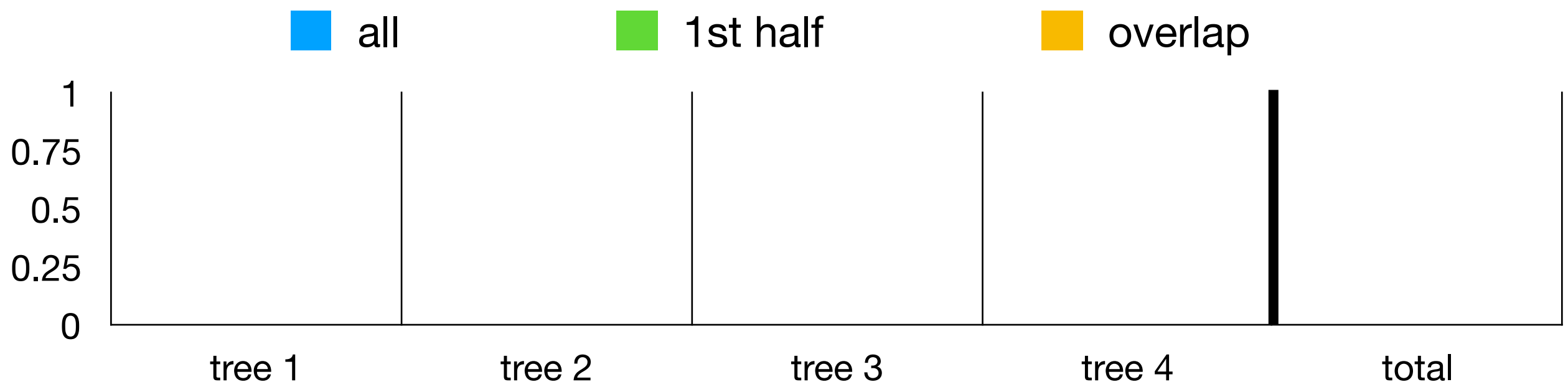
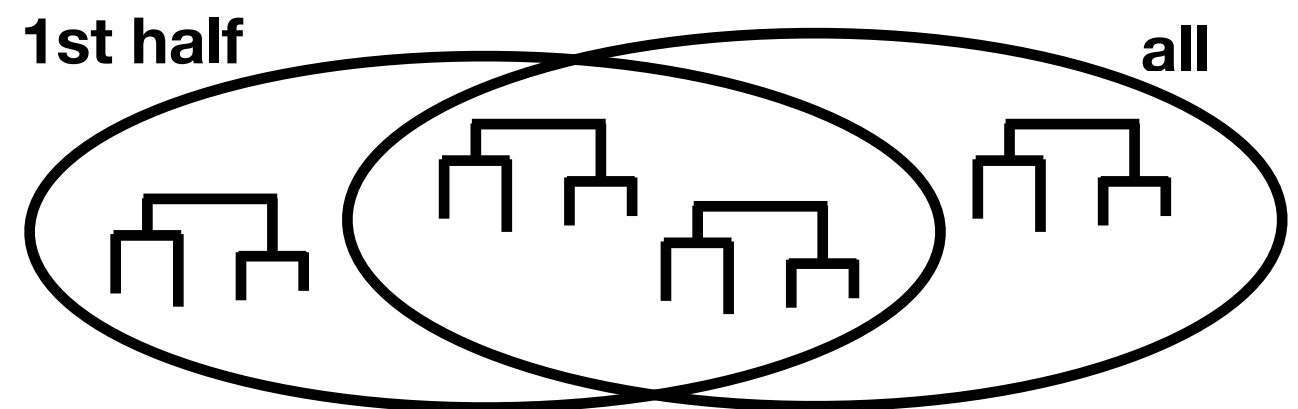
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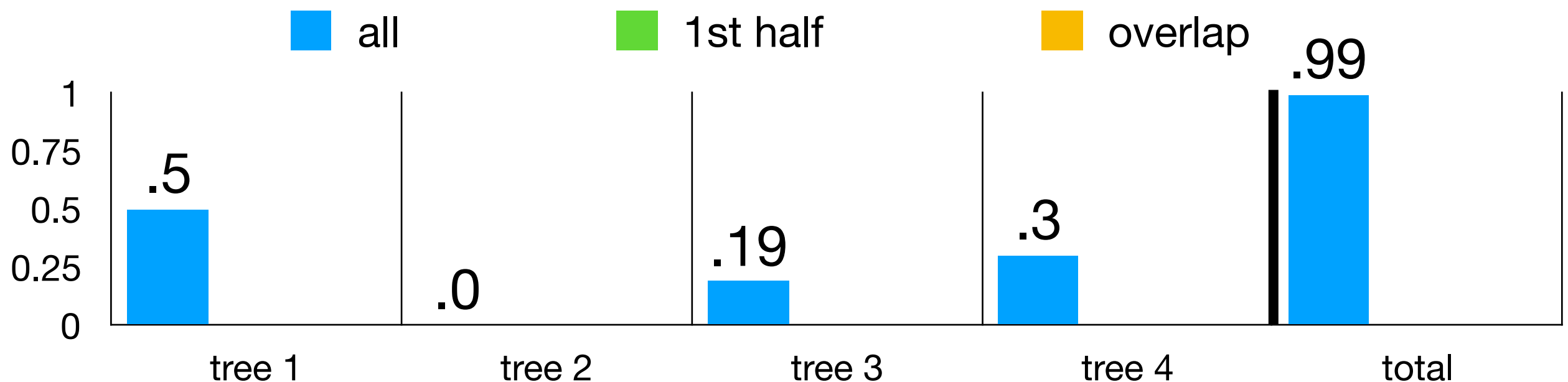
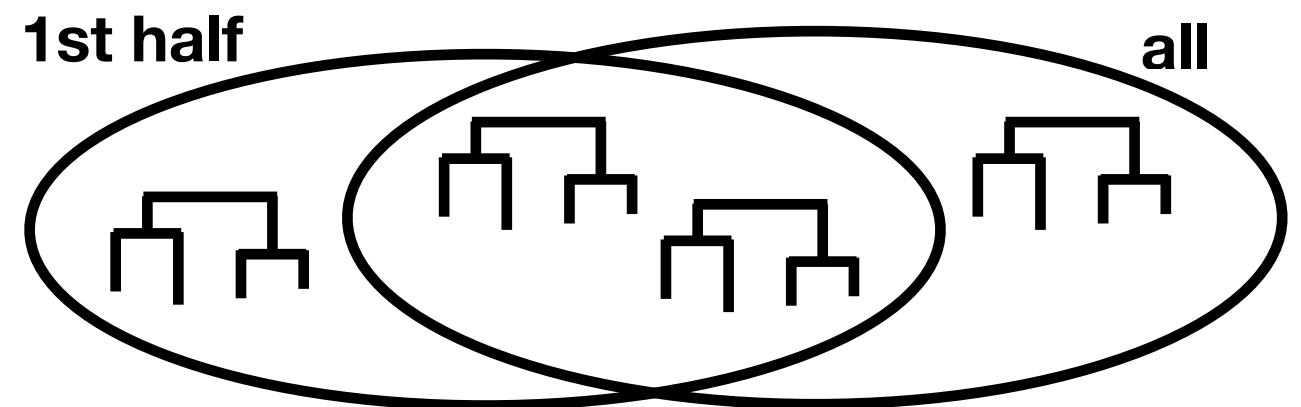
all	1st half	2nd half
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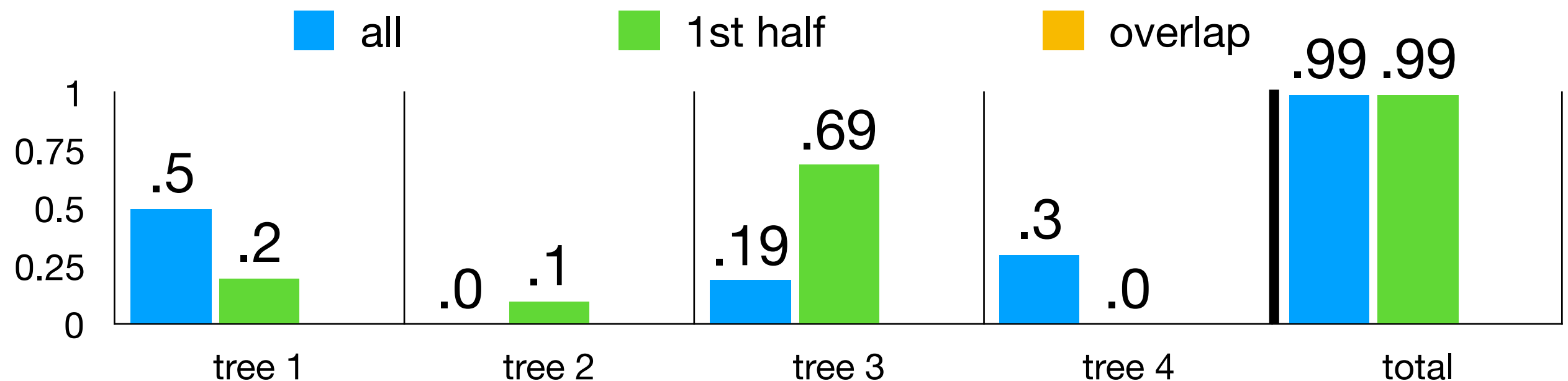
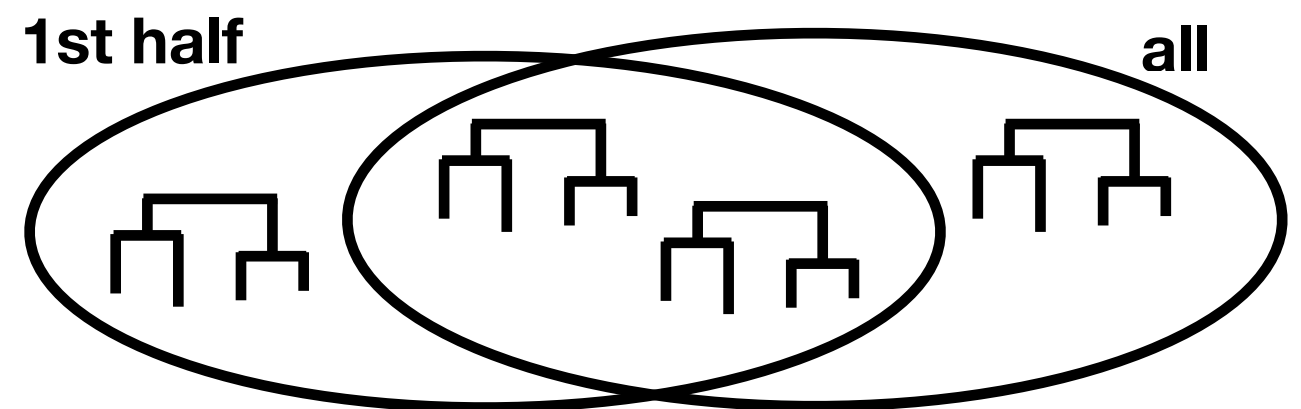
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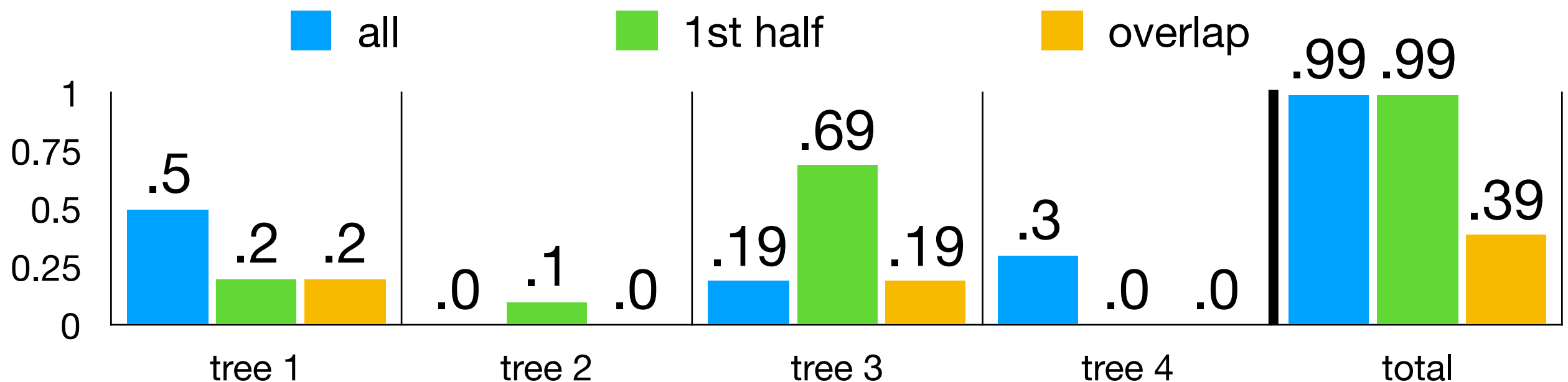
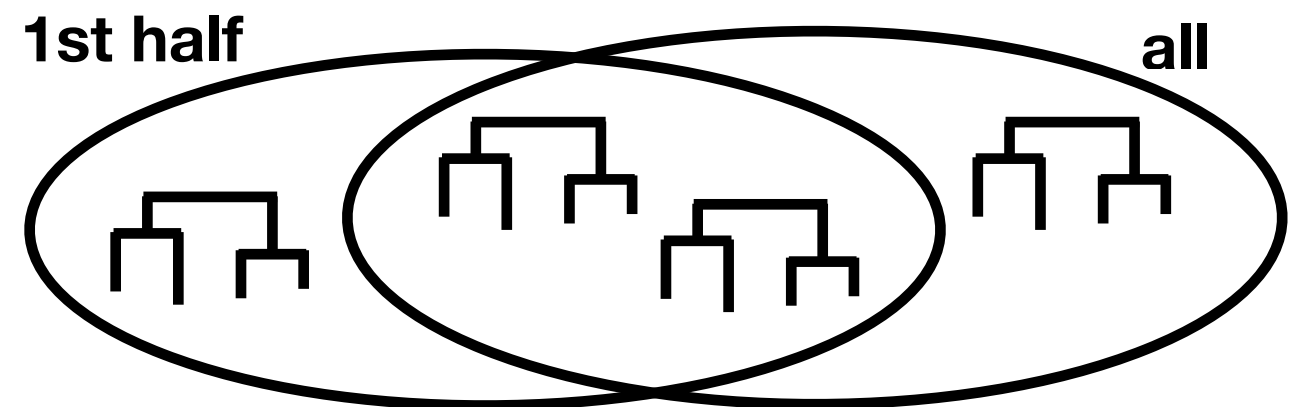
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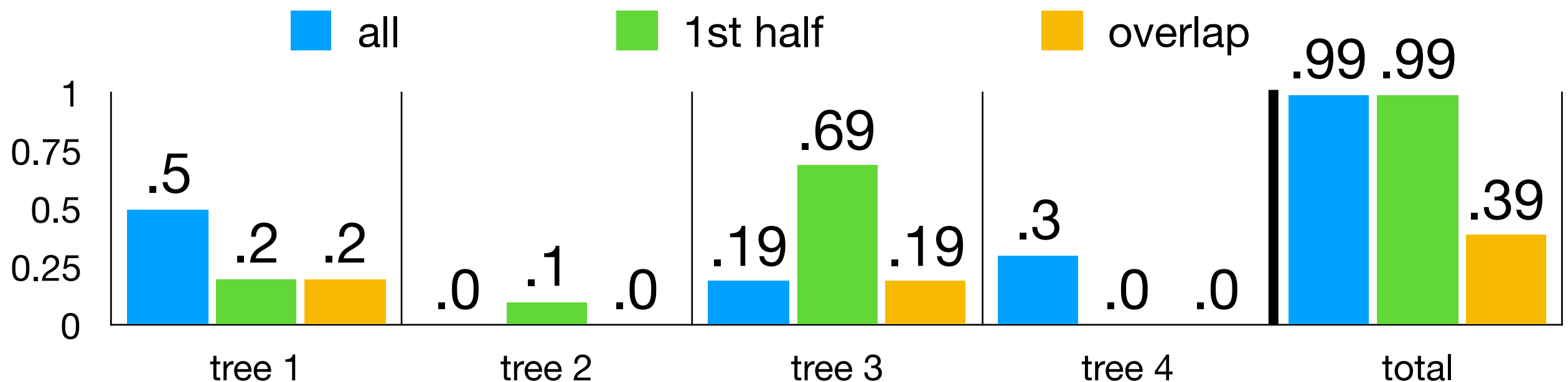
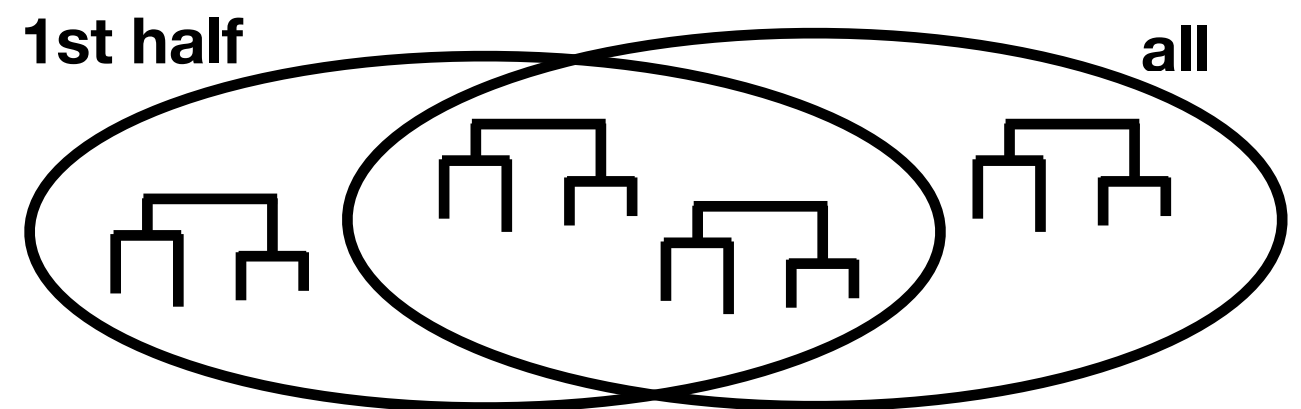
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- 0% overlap = contradiction

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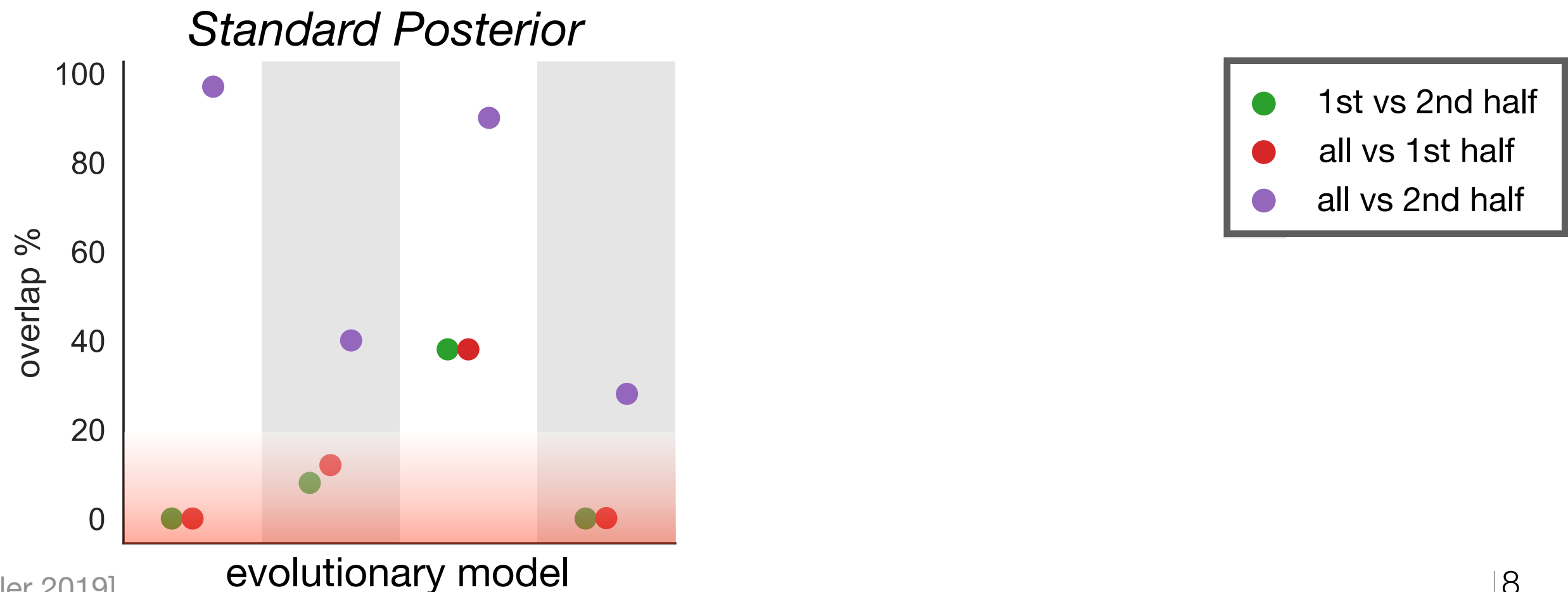


Stable, reproducible phylogenetic inference with BayesBag

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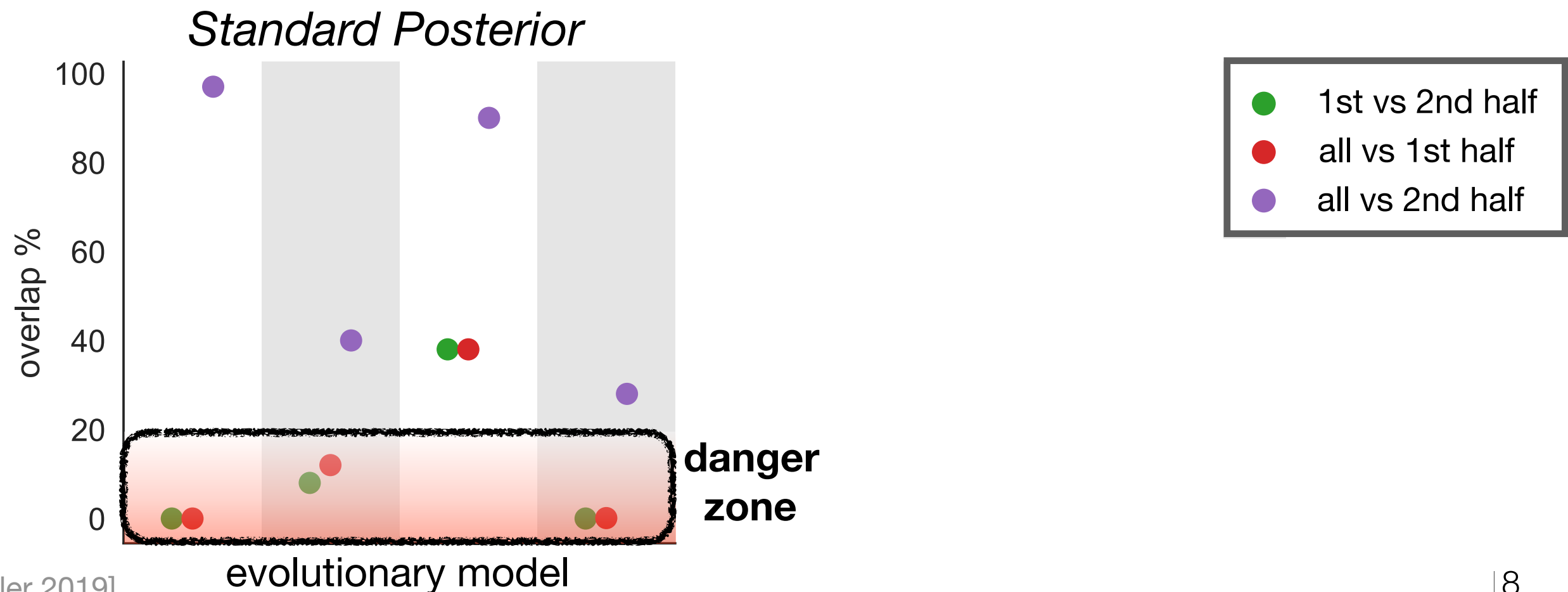
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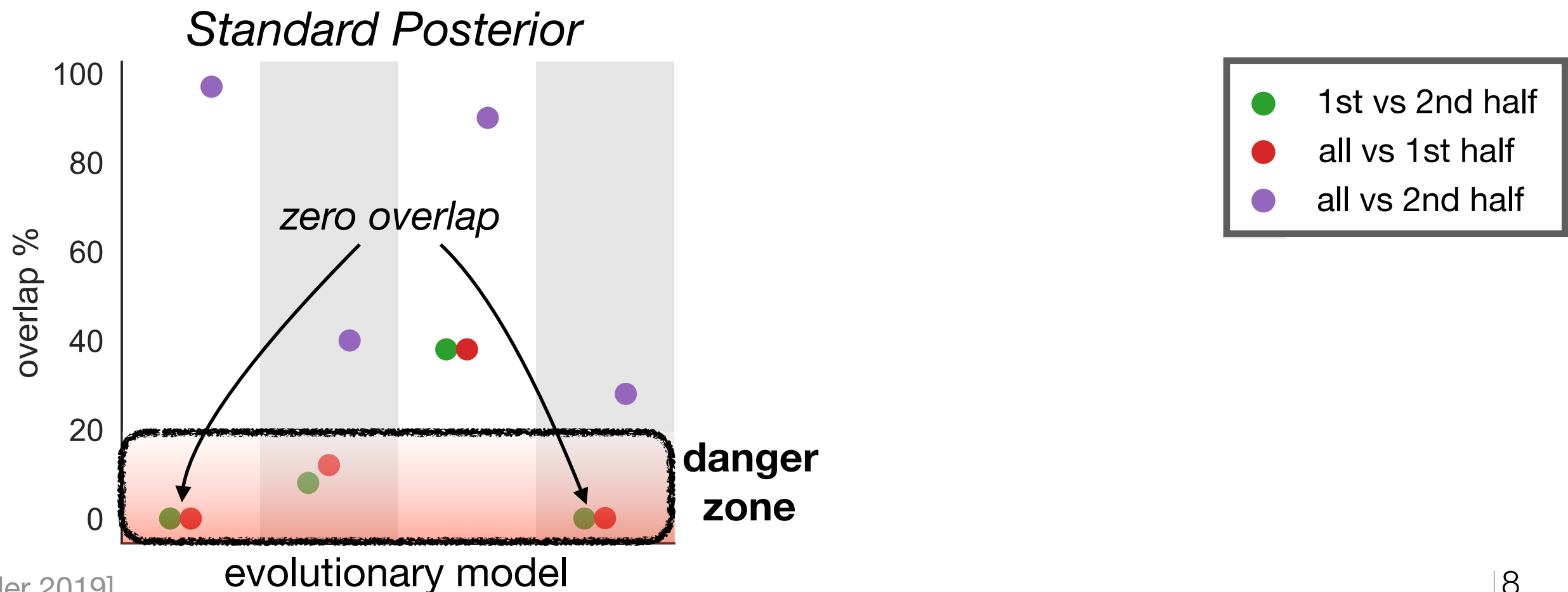
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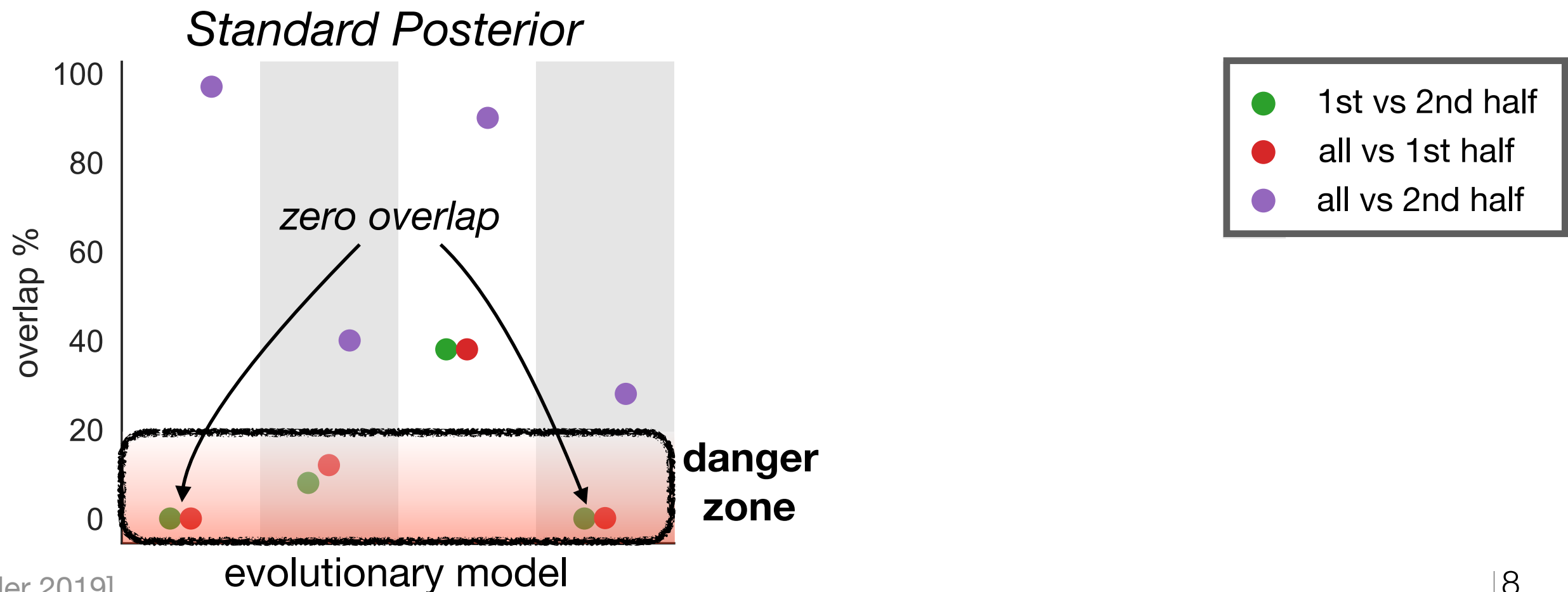
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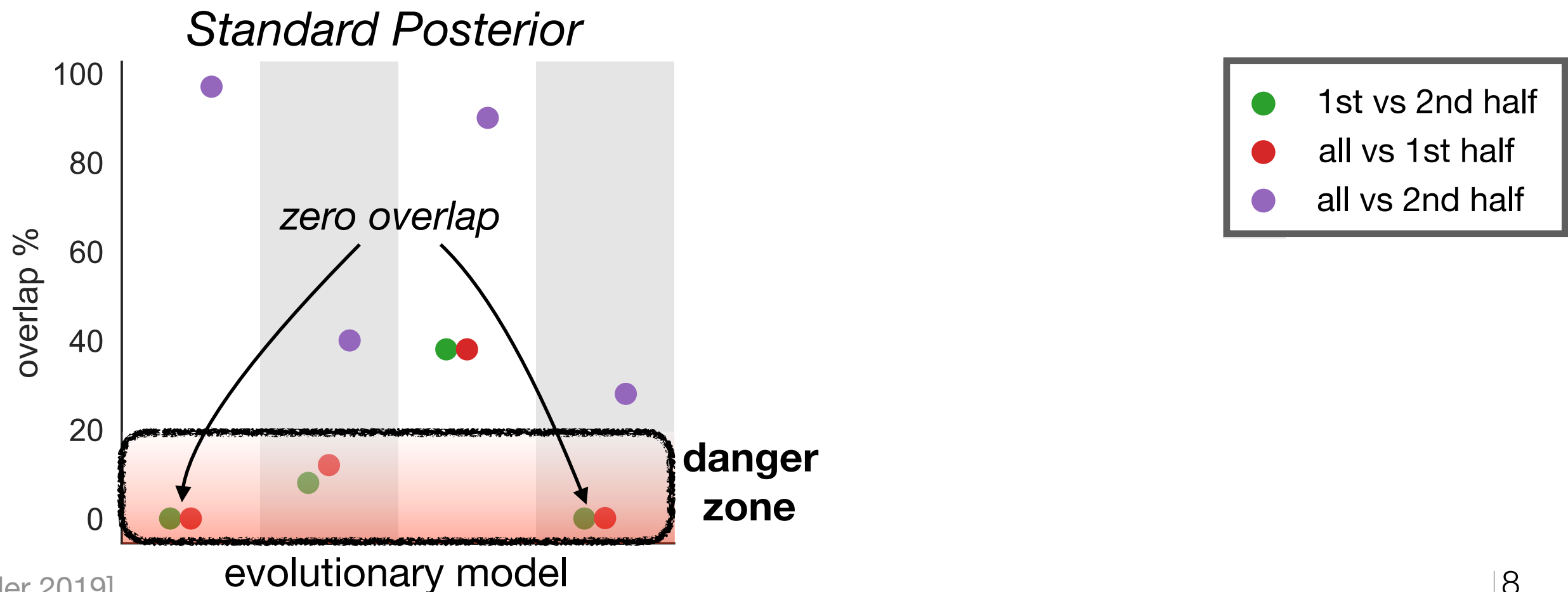
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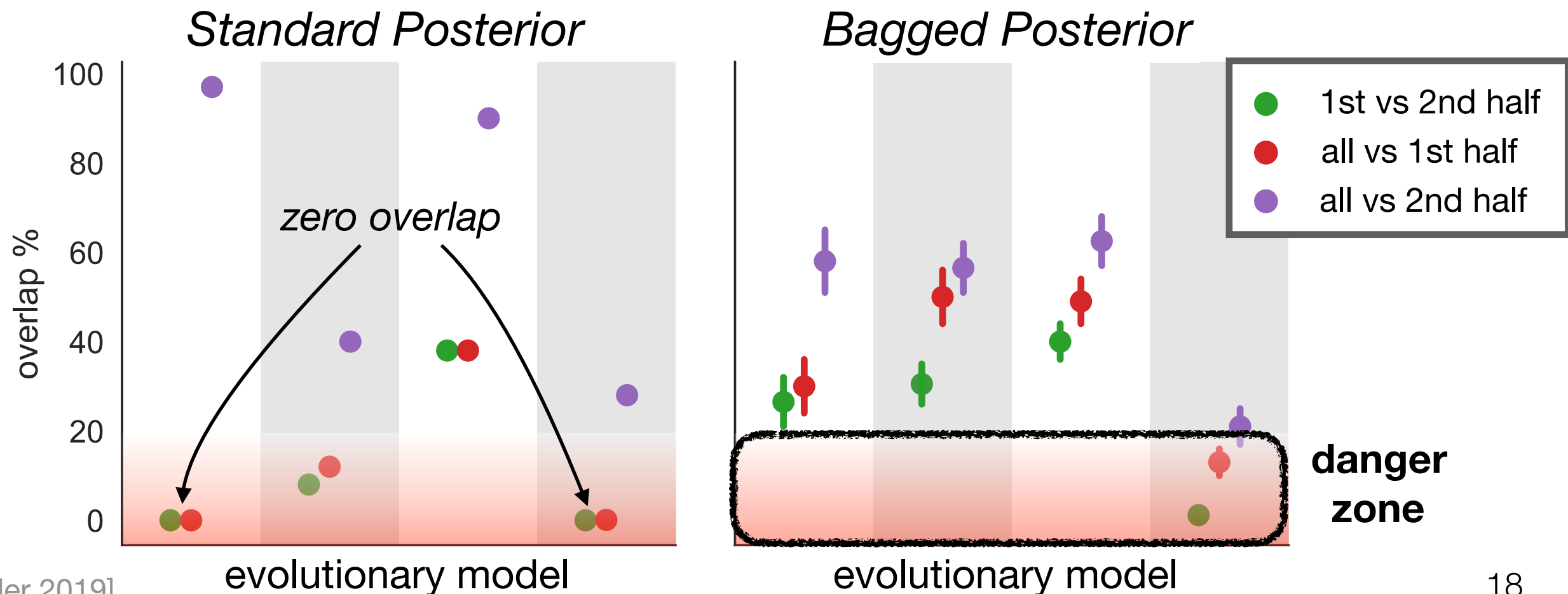
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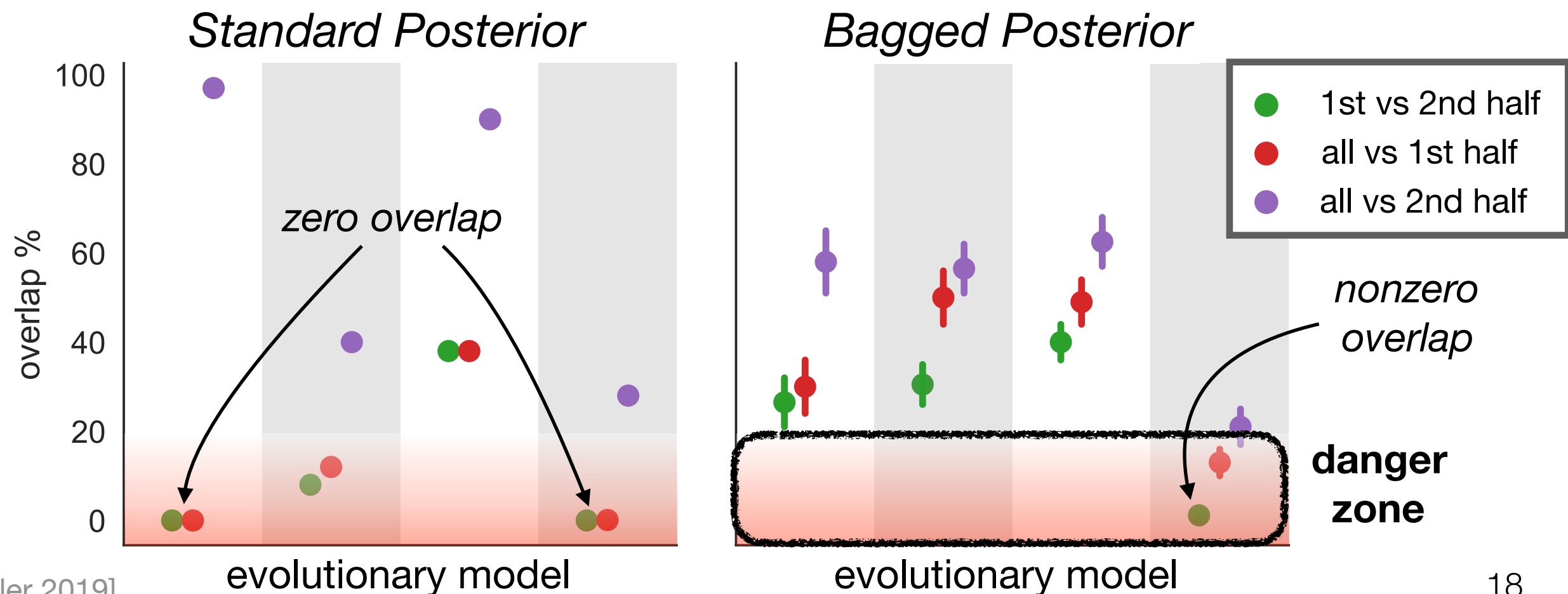
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Thank you

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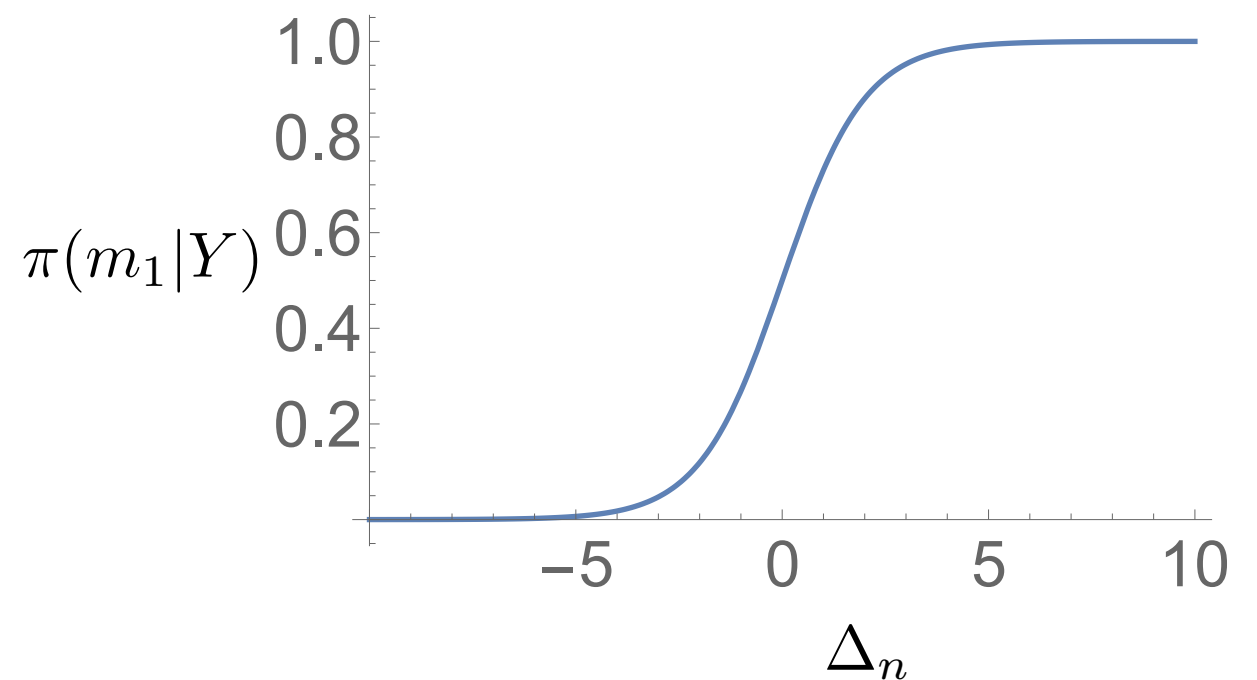
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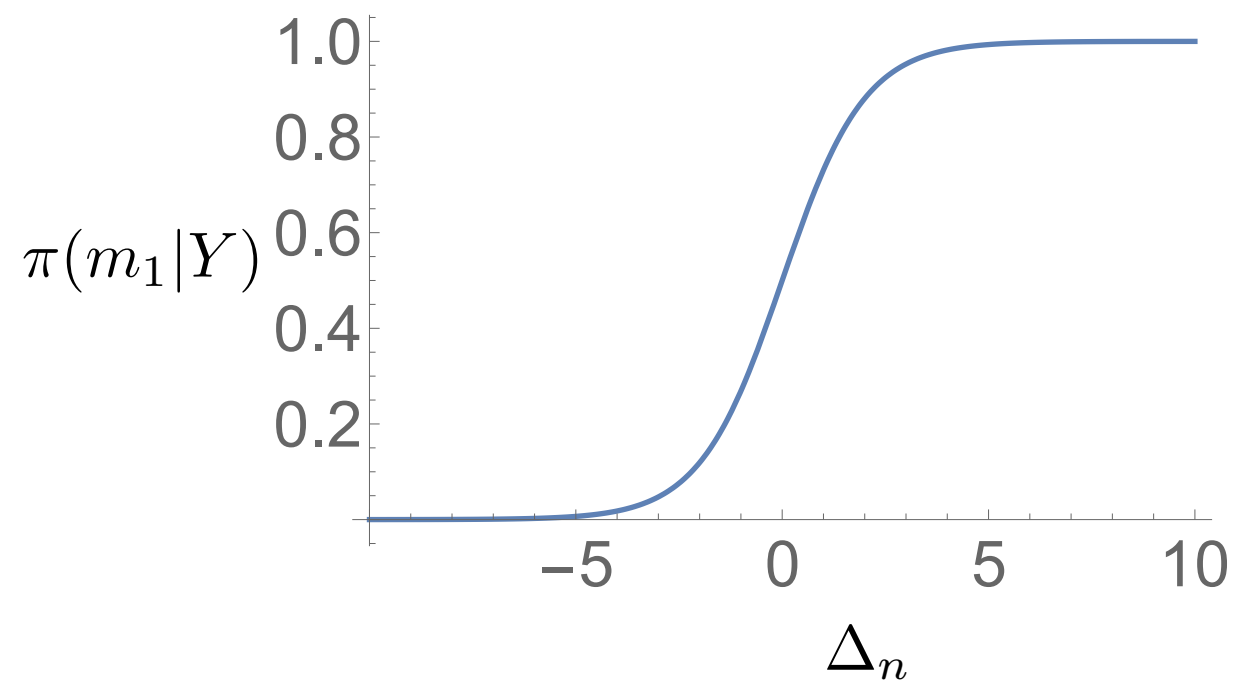


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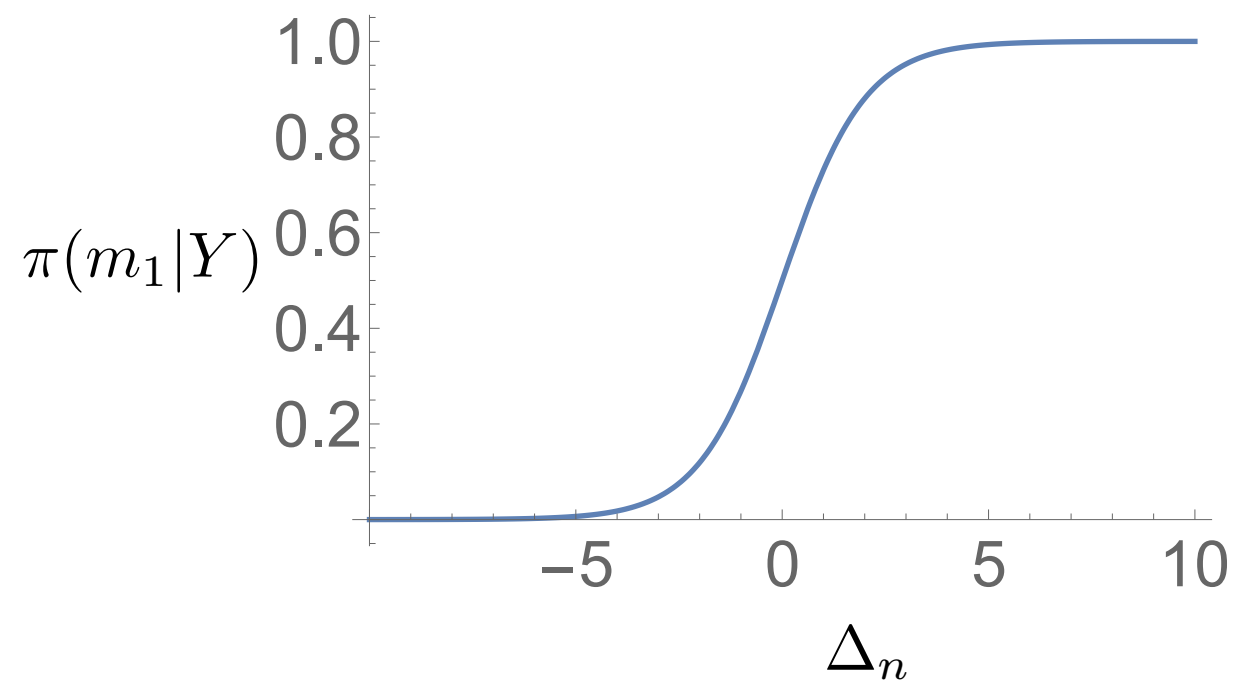
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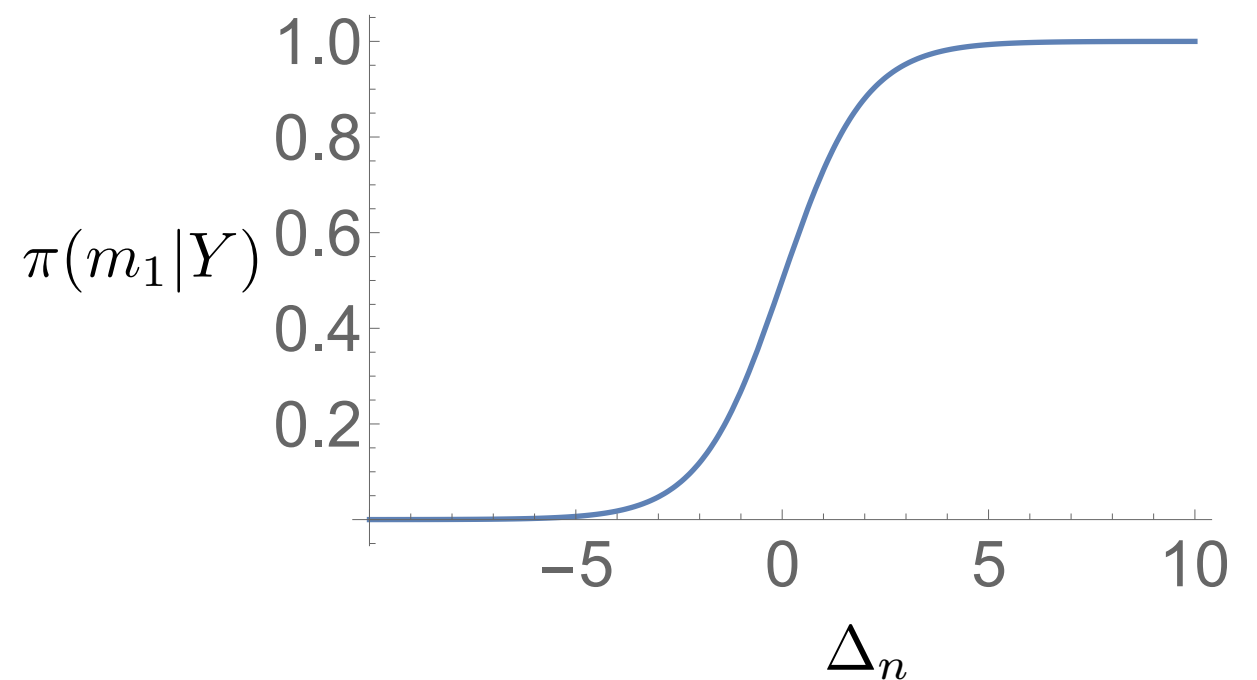
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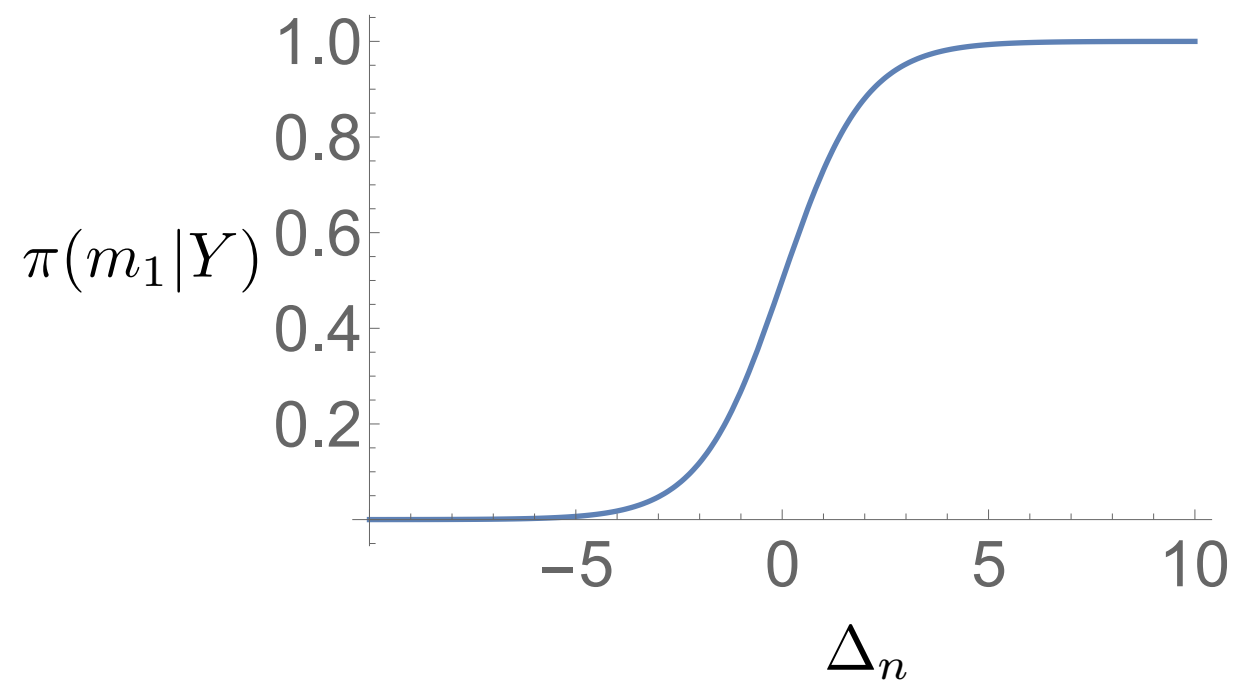
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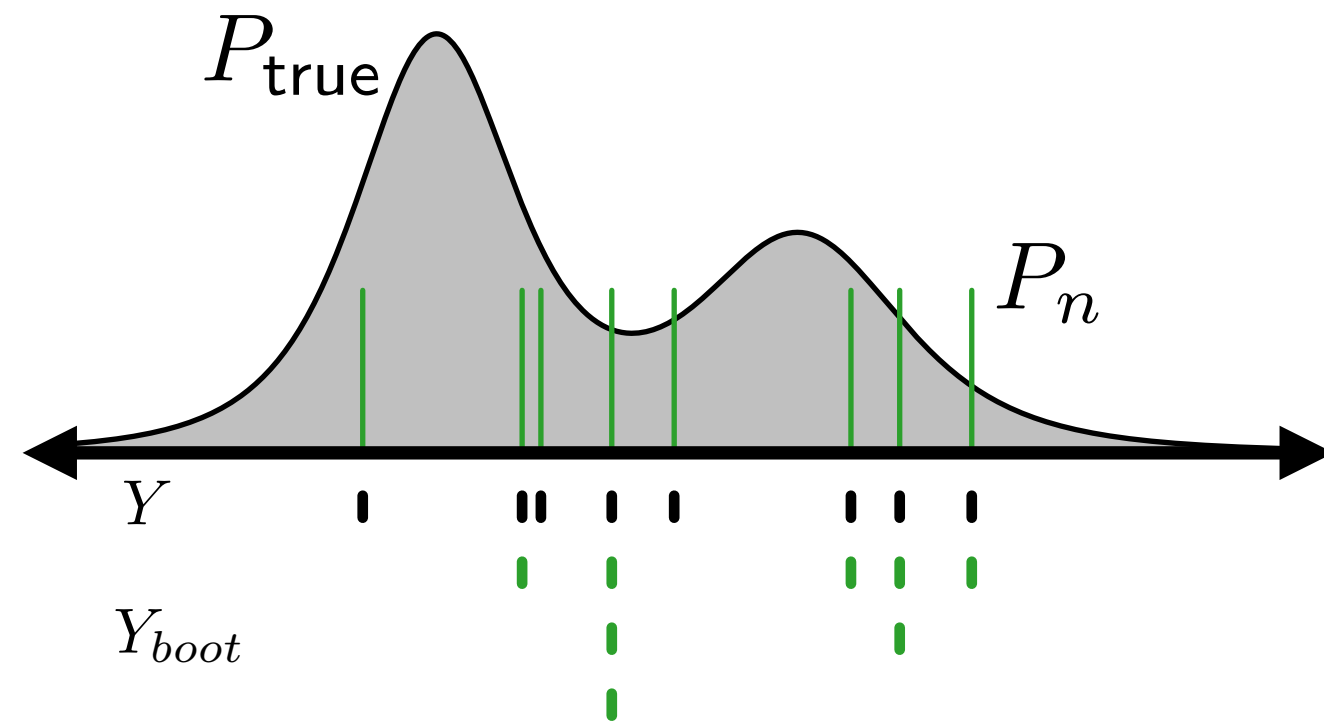
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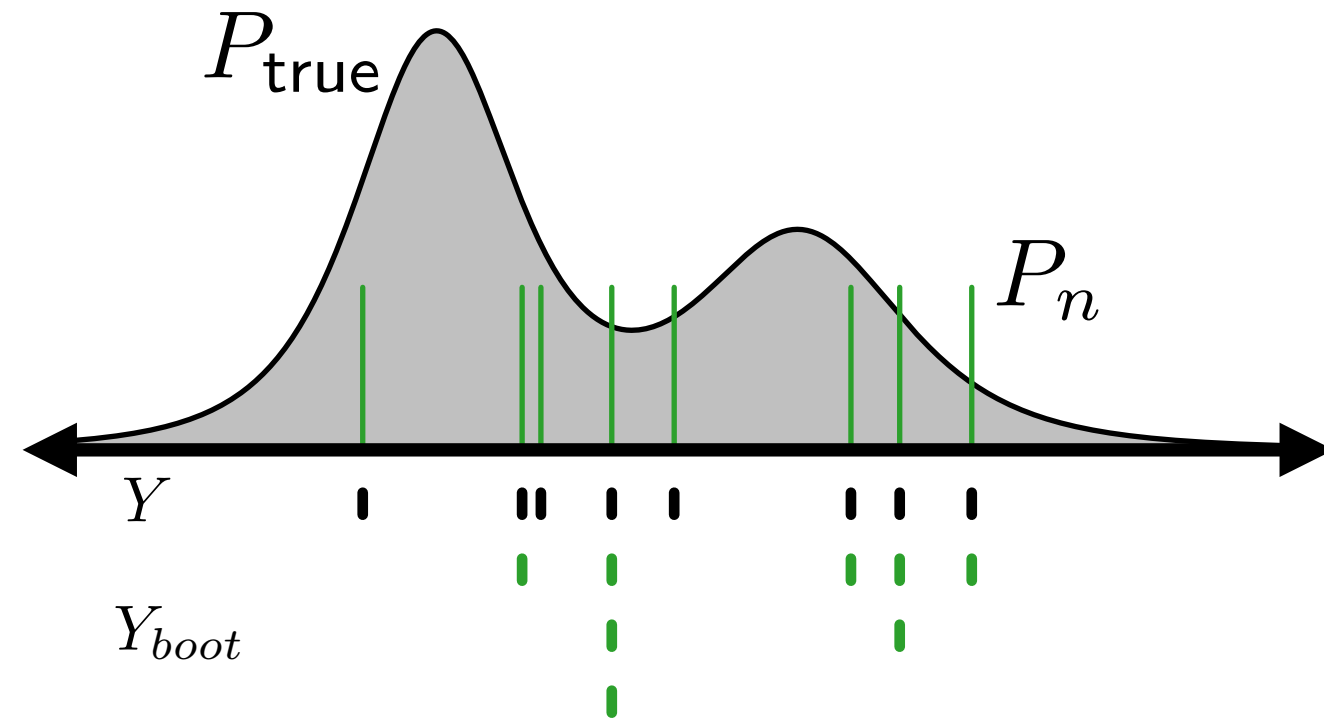


Bootstrap aggregating (bagging)



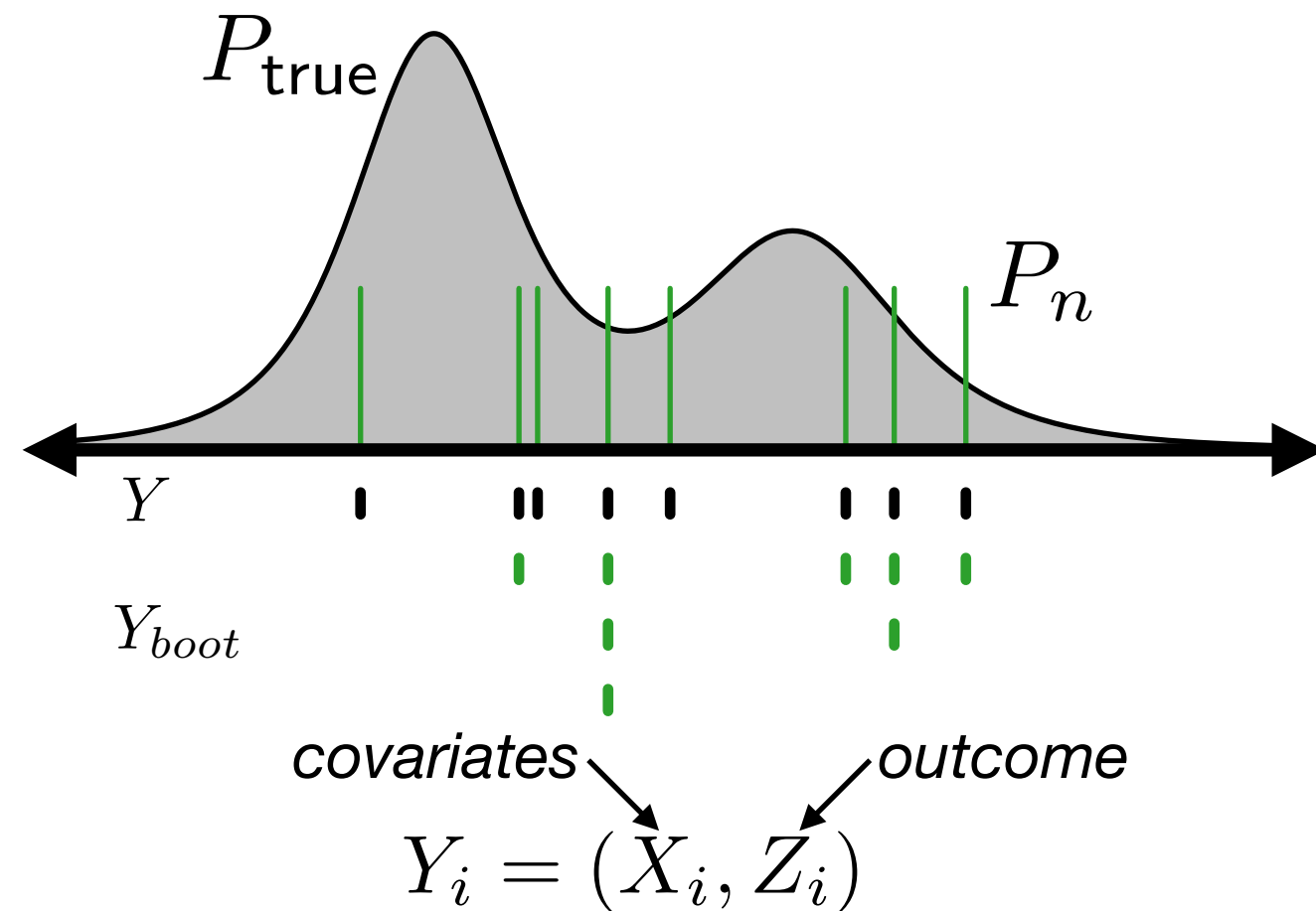
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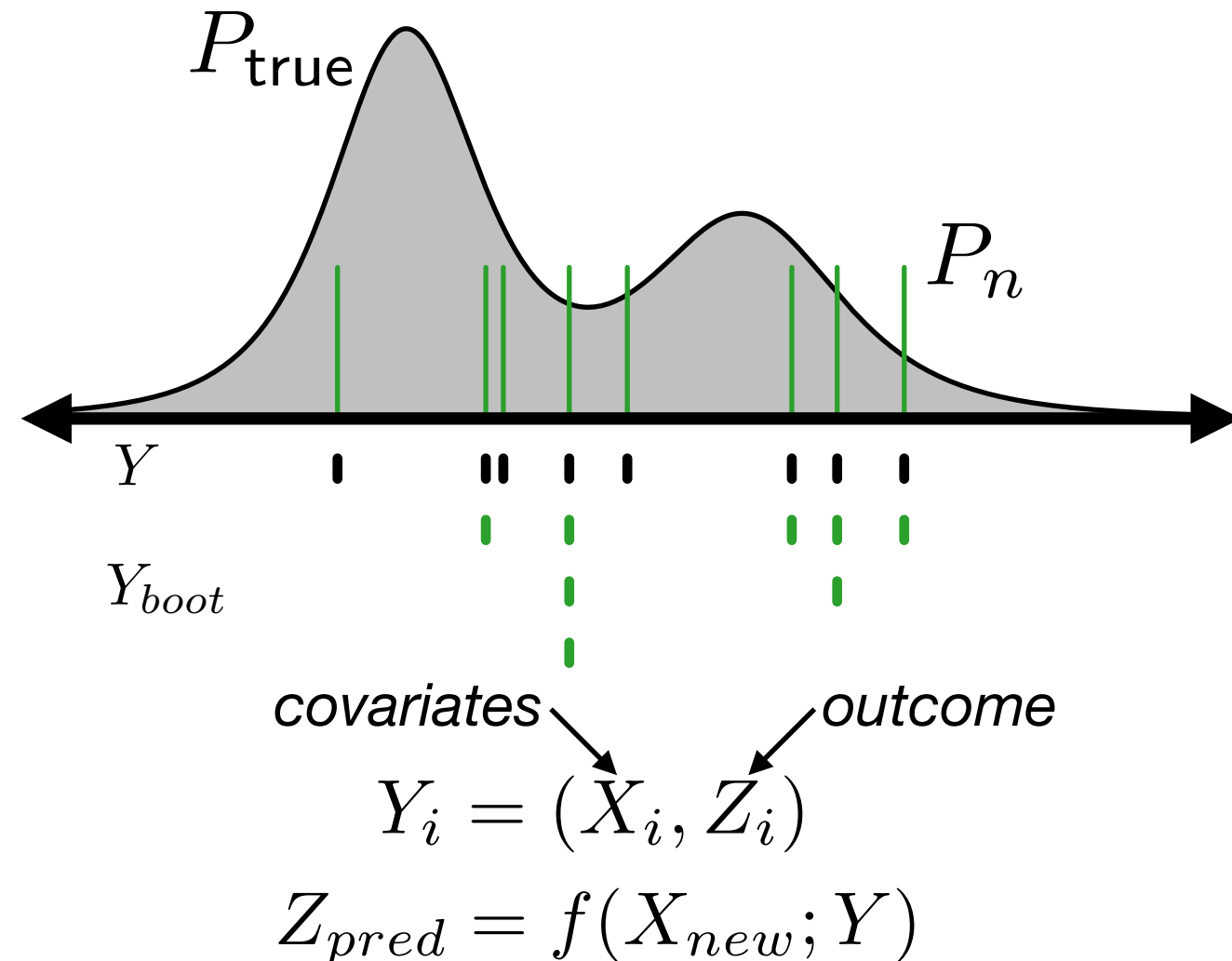
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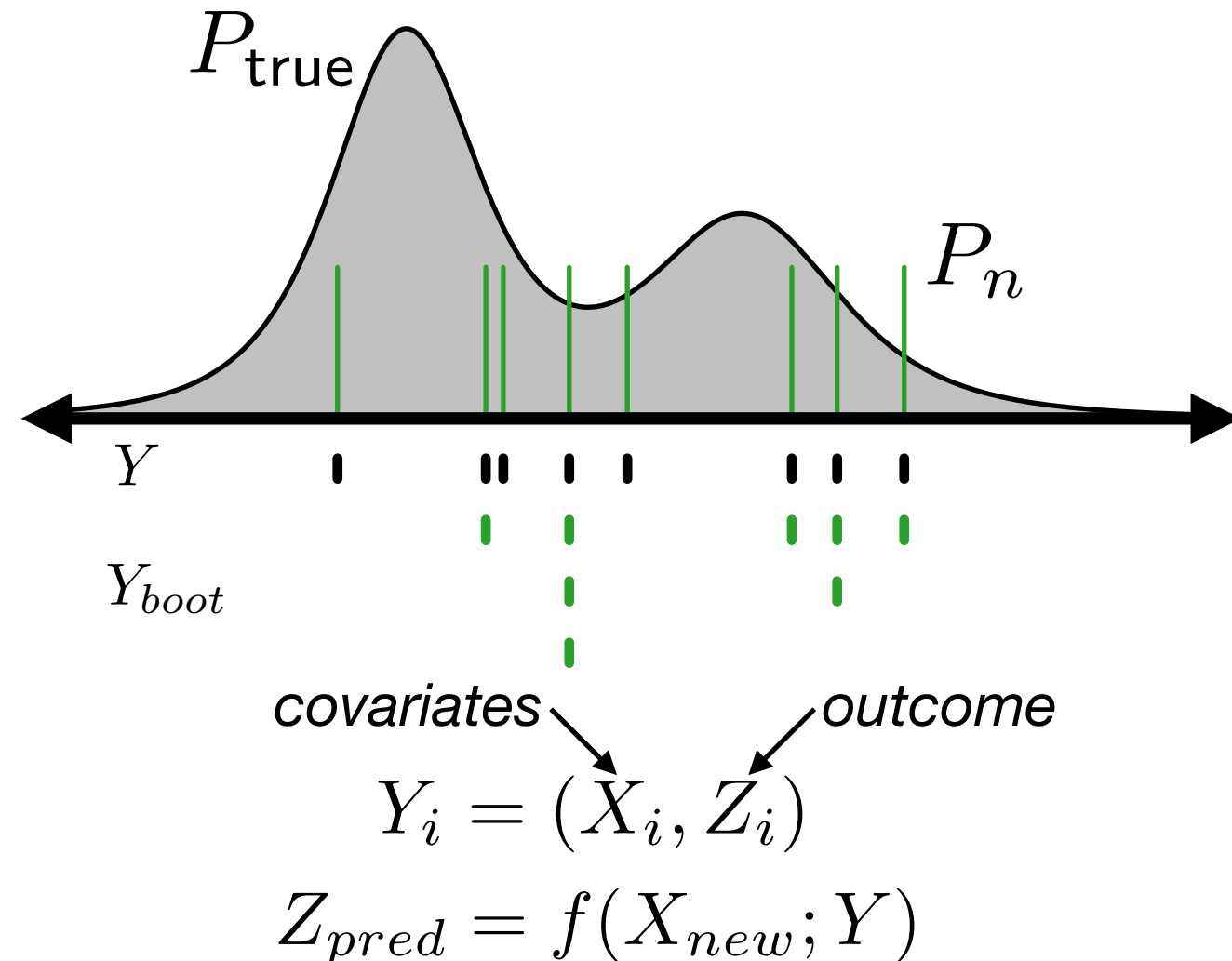
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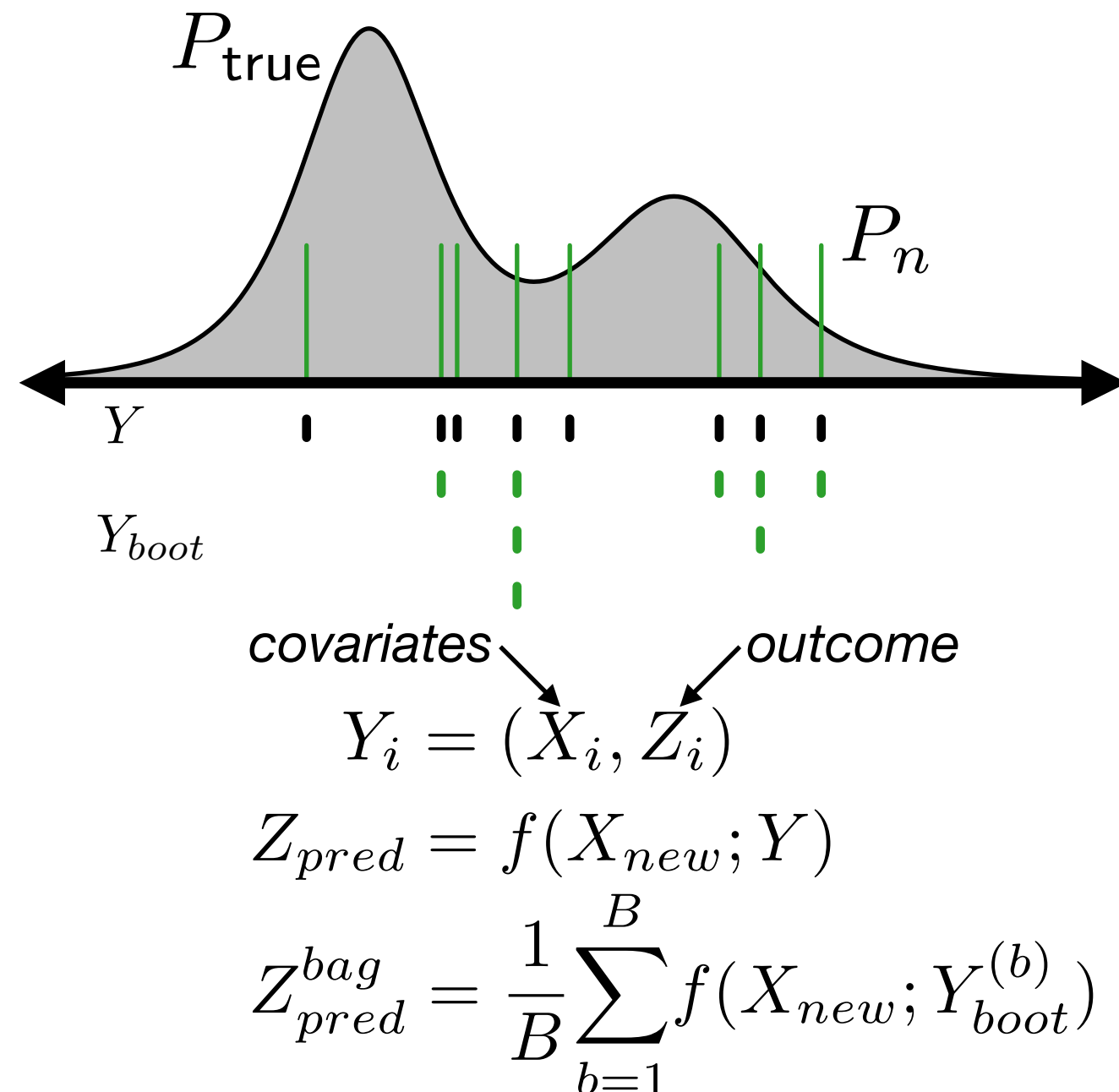
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- Like bagging, BayesBag seems to work well with $B = 50$ or 100

