

Intro to Bayesian Methods

Rebecca C. Steorts

Bayesian Methods and Modern Statistics: STA 360/601

Lecture 1

- ▶ Course Webpage
- ▶ Syllabus
- ▶ LaTeX reference manual
- ▶ R markdown reference manual
- ▶ Please come to office hours for all questions.
 - ▶ Office hours are not a review period if you cannot come to class.
- ▶ Join Google group
- ▶ Graded on Labs/HWs, Exams.
 - ▶ Labs/HWs and Exams .R markdown format (it must compile).
 - ▶ Nothing late will be accepted.
 - ▶ You're lowest homework will be dropped.
- ▶ Announcements: Emails or in class.
- ▶ All your lab/homework assignments will be uploaded to Sakai.
- ▶ How to reach me and TAs – email or Google.

Expectations

- ▶ Class is optional but you are expected to know everything covered in lecture.
- ▶ Not everything will always be on the slides.
- ▶ 2 Exams: in class, timed. Closed book, closed notes. (Dates are on the syllabus).
- ▶ There are NO make up exams.
- ▶ Late assignments will not be accepted. Don't ask.
- ▶ Final exam: during finals week.
- ▶ You should be reading the book as we go through the material in class.

Expectations for Homework

- ▶ Your write ups should be clearly written.
- ▶ Proofs: show all details.
- ▶ Data analysis: clearly explain.
- ▶ For data analysis questions, don't just turn in code.
- ▶ Code must be well documented.
- ▶ Code style: <https://google.github.io/styleguide/Rguide.xml>
- ▶ For all homeworks, can use Markdown or LaTeX. You must include all files that lead to your solutions (this includes code)!

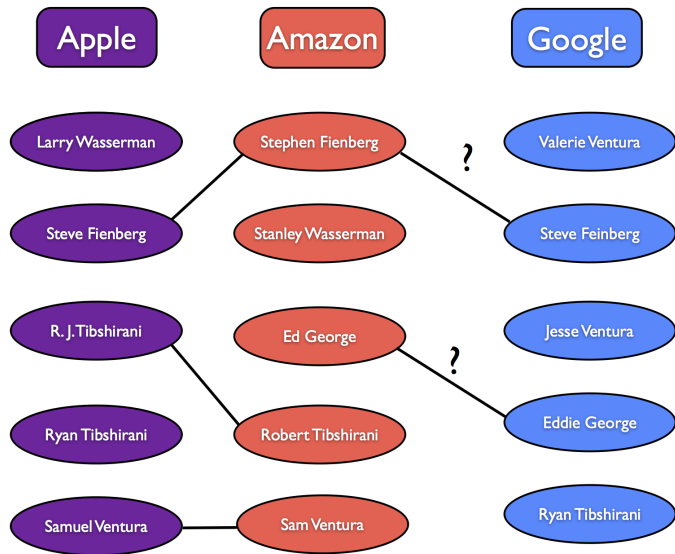
Things available to you!

- ▶ Come to office hours. We want to help you learn!
- ▶ Supplementary reading to go with the notes by yours truly. (Beware of typos).
 - ▶ Undergrad level notes
 - ▶ PhD level notes
 - ▶ Example form of write up in .Rmd on Sakai (Module 0).
 - ▶ You should have your homeworks graded and returned within one week by the TA's!

- ▶ Why should we learn about Bayesian concepts?
- ▶ Natural if thinking about unknown parameters as random.
- ▶ They naturally give a full distribution when we perform an update.
- ▶ We automatically get uncertainty quantification.
- ▶ Drawbacks: They can be slow and inconsistent.

Record linkage

Record linkage is the process of merging together noisy databases to remove duplicate entries.





Steve Feinberg

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Pittsburgh PA 15235

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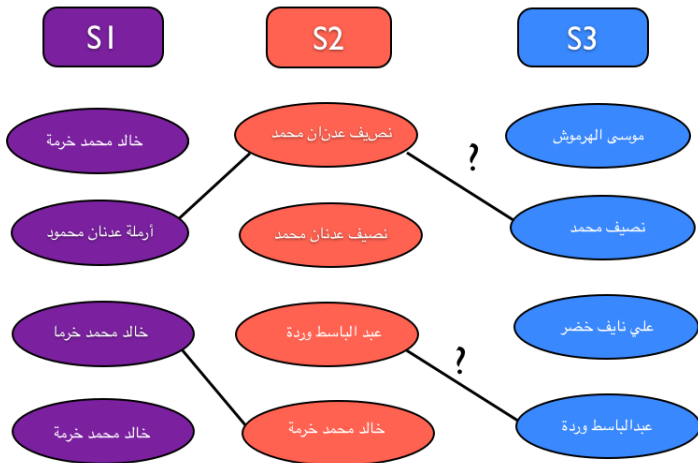
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These are clearly not the *same* Steve Fienberg!

Syrian Civil War



Bayesian Model

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$$X_{ij\ell} \mid \lambda_{ij}, Y_{\lambda_{ij}\ell}, z_{ij\ell} \sim \begin{cases} \delta(Y_{\lambda_{ij}\ell}) & \text{if } z_{ij\ell} = 0 \\ F_\ell(Y_{\lambda_{ij}\ell}) & \text{if } z_{ij\ell} = 1 \text{ and field } \ell \text{ is string-valued} \\ G_\ell & \text{if } z_{ij\ell} = 1 \text{ and field } \ell \text{ is categorical} \end{cases}$$

$$Y_{j'\ell} \sim G_\ell$$

$$z_{ij\ell} \mid \beta_{i\ell} \sim \text{Bernoulli}(\beta_{i\ell})$$

$$\beta_{i\ell} \sim \text{Beta}(a, b)$$

$$\lambda_{ij} \sim \text{DiscreteUniform}(1, \dots, N_{\max}), \quad \text{where } N_{\max} = \sum_{i=1}^k n_i$$

The model I showed you is very complicated.

This course will give you an intro to Bayesian models and methods.

Often Bayesian models are hard to work with, so we'll learn about approximations.

The above record linkage problem is one that needs such an approximation.

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$p(x \theta)$	likelihood
$p(\theta)$	prior

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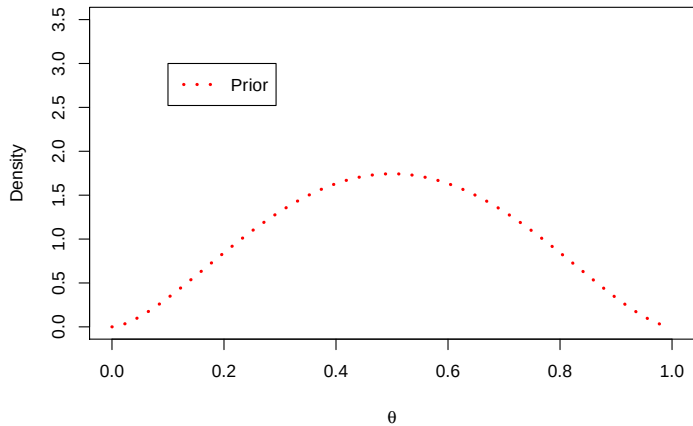
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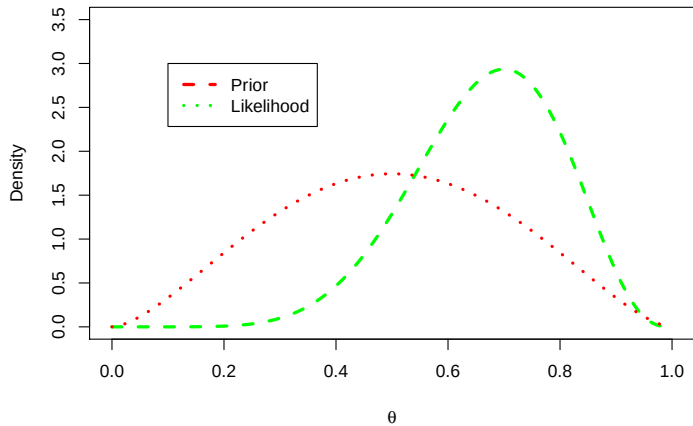
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- ▶ Based on this prior information, we'll use a Beta prior for θ and we'll choose a and b . (Won't get into this here).

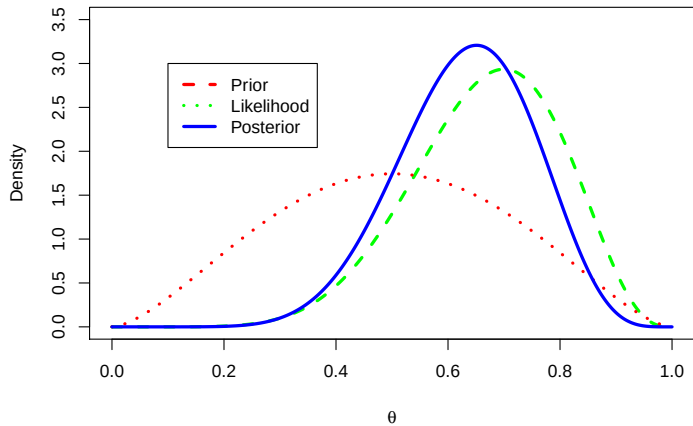
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- ▶ Based on this prior information, we'll use a Beta prior for θ and we'll choose a and b . (Won't get into this here).
- ▶ We can plot the prior and likelihood distributions in R and then see how the two mix to form the posterior distribution.







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- ▶ Bayesians treat an unknown parameter θ as *random*.
- ▶ Frequentists treat θ as unknown but *fixed*.

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- ▶ To perform a frequentist hypothesis test, we must define a random variable to describe the data.
- ▶ The proper way to do this depends on exactly which of the following two experiments was actually performed:

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So we fail to reject H_0 at $\alpha = 0.05$.

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So we reject H_0 at $\alpha = 0.05$.

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- ▶ P-values aren't close—one is 3.5 times as large as the other.
- ▶ The result our hypothesis test depends on whether we would have stopped flipping if we had gotten a tails sooner.
- ▶ The tests are dependent on what we call the *stopping rule*.

- ▶ The likelihood for the actual value of x that was observed is the same for both experiments (up to a constant):

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- ▶ A Bayesian approach would take the data into account only through this likelihood.
- ▶ This would provide the same answers regardless of which experiment was being performed.

The Bayesian analysis is independent of the stopping rule since it only depends on the likelihood (show this at home!).

Hierarchical Bayesian Models

In a hierarchical Bayesian model, rather than specifying the prior distribution as a single function, we specify it as a hierarchy.

Hierarchical Bayesian Models

$$X|\theta \sim f(x|\theta)$$

$$\Theta|\gamma \sim \pi(\theta|\gamma)$$

$$\Gamma \sim \phi(\gamma),$$

where we assume that $\phi(\gamma)$ is known and not dependent on any other unknown *hyperparameters*.

Conjugate Distributions

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- ▶ Then let P denote the class of prior distributions on θ .
- ▶ Then P is said to be conjugate to F if for every $p(\theta) \in P$ and $p(y|\theta) \in F$, $p(\theta | y) \in P$.

Simple definition: A family of priors such that, upon being multiplied by the likelihood, yields a posterior in the same family.

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$$\theta|x \sim \text{Beta}(x+a, n-x+b).$$