

Bayesian Inference

February 14, 2013

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- Randomly (iid) sample y from some data generating process (DGP): $P(y|\theta)$
- Goal is to estimate $P(\theta|y)$ given $P(y|\theta)$ and information about $P(\theta)$

Posterior Inference

Basic Setup

Simple
Example

Normal Model

Model
Features

- Use Bayes' theorem for $p(\theta|y)$:

$$P(\theta|y) = \frac{P(\theta, y)}{P(y)}$$

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- The Prior: $P(\theta)$
- The Sampling (Data): $P(y|\theta)$
- The Posterior: $P(\theta|y)$
- The Marginal: $P(y)$

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The Bayesian Approach

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- Use all available information about the model parameters
 - Evidence from the sampling process
 - All prior information

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- Why formalize prior information?
 - Inference drawn from the probability process
 - Incorporating the 'body of prior knowledge'
 - Integral to iterative data analysis and model checking

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- Use all available information about the model parameters
 - Evidence from the sampling process
 - All prior information
- Why formalize prior information?
 - Inference drawn from the probability process
 - Incorporating the 'body of prior knowledge'
 - Integral to iterative data analysis and model checking
- Why to not formalize prior information?
 - Does not emerge from the randomization process
 - Prior information could be explicitly biased
 - Inferences are not invariant to reparameterization (or even different types of diffuse priors)
 - Computationally and mathematically complex

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- “The posterior is proportional to the prior and the likelihood:”

$$p(\theta|y) \propto p(\theta)p(y|\theta)$$

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- Estimate $\hat{\theta}$ by randomly sampling from the posterior $p(\theta|y)$ distribution
 - Sample from a closed form of $p(\theta|y)$ based on the model and priors
 - Approximate $p(\theta|y)$ numerically or through sampling and monte carlo approaches

A Simple Example: The Prior

Basic Setup

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- Inference about genetic probability (Gelman et al 2004):
 - Hemophilia exhibits X-chromosomal recessive inheritance
 - A male with the carrier X-chromosome is affected, whereas a woman with only one carrier X-chromosome is not

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- Inference about genetic probability (Gelman et al 2004):
 - Hemophilia exhibits X-chromosomal recessive inheritance
 - A male with the carrier X-chromosome is affected, whereas a woman with only one carrier X-chromosome is not
- Consider a woman whose brother is affected, but whose father is not
 - The woman is either a carrier ($\theta = 1$) or is not ($\theta = 0$)
 - Given no other information, what is the $Pr(\theta = 1)$ for the woman?

A Simple Example: The Data

Basic Setup

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- The data we collect to update this prior is the affliction status status of her sons
 - Define y_i be 1 if her i th son is afflicted, and 0 otherwise
 - Assume that y_i and y_j are exchangeable for all $i \neq j$, and are independent conditional on θ

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- We then have the following likelihoods (for two unafflicted sons):
 - $Pr(y_1 = 0, y_2 = 0 | \theta = 1)$?
 - $Pr(y_1 = 0, y_2 = 0 | \theta = 0)$?

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- In this process:
 - What is $Pr(y_1 = 1 | \theta = 1)$?
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 - What is $Pr(y_1 = 1 | \theta = 1) = 1/2$
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 - Define y_i be 1 if her i th son is afflicted, and 0 otherwise
 - Assume that y_i and y_j are exchangeable for all $i \neq j$, and are independent conditional on θ
- We then have the following likelihoods (for two unafflicted sons):
 - $Pr(y_1 = 0, y_2 = 0 | \theta = 1) = (1/2)(1/2) = 1/4$
 - $Pr(y_1 = 0, y_2 = 0 | \theta = 0) = (1)(1) = 1$
- In this process:
 - What is $Pr(y_1 = 1 | \theta = 1) = 1/2$
 - What is $Pr(y_1 = 1 | \theta = 0) = 0$

A Simple Example: The Posterior

Basic Setup

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Normal Model

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Features

- Combine the prior and the evidence about her sons:

$$\begin{aligned}Pr(\theta = 1|y) &= \frac{p(y|\theta = 1)Pr(\theta = 1)}{p(y|\theta = 1)Pr(\theta = 1) + p(y|\theta = 0)Pr(\theta = 0)} \\&= \frac{(0.25)(0.5)}{(0.25)(0.5) + (1.0)(0.5)} = \frac{0.125}{0.625} = 0.2\end{aligned}$$

- A woman with no affected sons is less likely to be afflicted, 'correcting' the prior in the direction of the data

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- A woman with no affected sons is less likely to be afflicted, 'correcting' the prior in the direction of the data
- Add more 'data' with a third unaffected son, using the previous posterior as the new prior:

$$Pr(\theta = 1|y_1, y_2, y_3) = \frac{(0.5)(0.2)}{(0.5)(0.2) + (1.0)(0.8)} = 0.111$$

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- A third son who IS affected? What is the ML estimate?

Normal Data with Known Variance

Basic Setup

Simple
Example

Normal Model

Model
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- We randomly sample (iid) n units, and observe y_i .
Assuming $y \sim N(\beta, \sigma^2)$:

$$P(y|\theta) = \prod_{i=1}^n p(y_i|\theta)$$

$$\begin{aligned} P(y|\beta, \sigma) &= \prod_{i=1}^n N(y_i|\beta, \sigma^2) \\ &= \prod_{i=1}^n \frac{1}{\sigma\sqrt{2\pi}} \exp \left\{ -\frac{1}{2} \left(\frac{y_i - \beta}{\sigma} \right)^2 \right\} \end{aligned}$$

Normal Data with Known Variance

Basic Setup

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- To estimate β using $p(\beta|y)$, we need to choose a prior distribution $p(\beta)$
- A natural choice is a Gaussian prior: $p(\beta) \sim N(\mu_0, \tau_0^2)$, where μ_0, τ_0^2 are known hyperparameters. Thus:

$$p(\beta) \propto \exp \left\{ -\frac{1}{2\tau_0^2} (\beta - \mu_0)^2 \right\}$$

- This is a conjugate prior, i.e., in the same probability distribution family

Normal Data with Known Variance

Basic Setup

Simple
Example

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- Use this prior to identify an analytical form for $p(\beta|y)$

$$\begin{aligned}
 P(\beta|y) &\propto \prod_{i=1}^n \frac{1}{\sigma\sqrt{2\pi}} \exp\left\{-\frac{1}{2\sigma^2} (y_i - \beta)^2\right\} \\
 &\quad \times \exp\left\{-\frac{1}{2\tau_0^2} (\beta - \mu_0)^2\right\} \\
 &\propto \exp\left\{-\frac{1}{2} \left[\frac{\sum_{i=1}^n (y_i - \beta)^2}{\sigma^2} + \frac{(\beta - \mu_0)^2}{\tau_0^2} \right]\right\}
 \end{aligned}$$

Normal Data with Known Variance

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- Combine and factor out terms that do not depend on β

$$\propto \exp \left\{ -\frac{1}{2} \left[\frac{\sum y_i^2 - 2\beta \sum y_i + n\beta^2}{\sigma^2} + \frac{\beta^2 - 2\beta\mu_0 + \mu_0^2}{\tau_0^2} \right] \right\}$$

$$\propto \exp \left\{ -\frac{\beta^2}{2\tau_0^2} + \frac{\beta\mu_0}{\tau_0^2} + \frac{\beta n\bar{y}}{\sigma^2} - \frac{n\beta^2}{2\sigma^2} \right\}$$

$$\propto \exp \left\{ -\frac{\beta^2}{2} \left(\frac{1}{\tau_0^2} + \frac{n}{\sigma^2} \right) + \beta \left(\frac{\mu_0}{\tau_0^2} + \frac{n\bar{y}}{\sigma^2} \right) \right\}$$

$$\propto \exp \left\{ -\frac{\beta^2}{2\tau_*^2} + \frac{2}{2} \cdot \frac{\beta\mu_*}{\tau_*^2} \right\} = \exp \left\{ -\frac{1}{2\tau_*^2} (\beta^2 - 2\beta\mu_*) \right\}$$

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$$\propto \exp \left\{ -\frac{\beta^2}{2\tau_*^2} + \frac{2}{2} \cdot \frac{\beta\mu_*}{\tau_*^2} \right\} = \exp \left\{ -\frac{1}{2\tau_*^2} (\beta^2 - 2\beta\mu_*) \right\}$$

- Completing the square:

$$\propto \exp \left\{ -\frac{1}{2\tau_*^2} (\beta - \mu_*)^2 - \mu_*^2 \right\}$$

$$p(\beta|y) \propto \exp \left\{ -\frac{1}{2\tau_*^2} (\beta - \mu_*)^2 \right\}$$

Normal Data with Known Variance

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- We should notice that this implies $\beta|y \sim N(\mu_*, \tau_*^2)$, where μ_* and τ_*^2 are parameterized as:

$$\begin{aligned}\mu_* &= \tau_*^2 \left(\frac{\mu_0}{\sigma_0^2} + \frac{n\bar{y}}{\sigma^2} \right) \\ \frac{1}{\tau_*^2} &= \left(\frac{1}{\tau_0^2} + \frac{n}{\sigma^2} \right)\end{aligned}$$

- We see the posterior mean μ_* is a weighted average of the prior mean μ_0 and the data $n\bar{y}$, normalized by the posterior *precision* τ_*^2
- Should also observe that as number of samples n gets large, μ_* is increasingly weighted towards \bar{y}

Posterior Estimation of $\hat{\beta}$

- We can estimate $\hat{\beta}$ simply by sampling from $\beta|y$, and taking the expectation $E[\beta|y]$

```
mu0 <- 1.2; tau0 <- 1.3; sigma <- 1.5; N <- 10
set.seed(1004)
y <- rnorm(mean=2.3,sd=sqrt(sigma),n=N)

tau_star <- function(n,tau0,sigma){
  return(1/((1/tau0^2)+(n/sigma^2)))}

mu_star <- function(y,n,mu0,tau0,sigma,tau_star){
  return(tau_star*((mu0/tau0^2)+
    (n*mean(y)/sigma^2)))}

ts <- tau_star(n=N,tau0,sigma)
ms <- mu_star(y,n=N,mu0,tau0,sigma,tau_star=ts)
bhat <- mean(rnorm(mean=ms,sd=sqrt(ts),n=1000))
```

Conjugacy

Basic Setup

Simple
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- Recall that we said:

$$\begin{aligned} y|\beta &\sim N(\beta, \sigma^2) \\ \beta &\sim N(\mu_0, \tau_0^2) \end{aligned}$$

- whose product is the posterior:

$$\beta|y \sim N(\mu_*, \tau_*^2)$$

- so the evidence, prior and posterior are all Normal
- This feature is called *conjugacy*: all model components are from the same class of probability distributions
 - Analytic and computational gains from considering conjugate models
 - Sampling techniques and powerful computers lessen the need for conjugacy

Diffuse Priors

- Consider the Normal model with a vague (uniform within a deviation) prior about β :

$$p(\beta, \sigma^2) \propto \frac{1}{\sigma^2}$$

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- The joint posterior distribution is proportional to the product of the likelihood and $1/\sigma^2$:

$$\begin{aligned} p(\beta, \sigma^2 | y) &\propto \sigma^{-n-2} \exp \left\{ -\frac{1}{2\sigma^2} \sum_{i=1}^n (y_i - \beta)^2 \right\} \\ &\propto \sigma^{-n-2} \exp \left\{ -\frac{1}{2\sigma^2} \left[\sum_{i=1}^n (y_i - \bar{y}) + n(\bar{y} - \beta) \right] \right\} \\ &\propto \sigma^{-n-2} \exp \left\{ -\frac{1}{2\sigma^2} [(n-1)s^2 + n(\bar{y} - \beta)] \right\} \end{aligned}$$

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- with $s^2 = \frac{1}{n-1} \sum_{i=1}^n (y_i - \bar{y})^2$, $\bar{y} = \frac{1}{n} \sum_{i=1}^n y_i$

Diffuse Priors

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- We can show the conditional posterior is $\beta|\sigma^2, y \sim N(\bar{y}, \frac{\sigma^2}{n})$
- Summing over β in the joint posterior, we can find the marginal posterior $\sigma^2|y \sim \text{Inv-}\chi^2(n-1, s^2)$

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- We can show the conditional posterior is $\beta|\sigma^2, y \sim N(\bar{y}, \frac{\sigma^2}{n})$
- Summing over β in the joint posterior, we can find the marginal posterior $\sigma^2|y \sim \text{Inv-}\chi^2(n-1, s^2)$
- Thus an alternative form of $p(\beta, \sigma^2|y)$ is:

$$N(\bar{y}, \frac{\sigma^2}{n}) \times \text{Inv-}\chi^2(n-1, s^2)$$

- Suggests we can iteratively sample σ^2 from the marginal posterior $\sigma^2|y$, then using this σ^2 draw, can sample from $\beta|\sigma^2, y$ to obtain a posterior estimates for β and σ^2

Estimation of $\hat{\beta}$ with Diffuse Priors

- The above suggests we sample σ_t^2 from $p(\sigma^2|y)$, and then sample from $p(\beta|\sigma_t^2, y)$ using this sample value of σ_t^2 to find β_t

```
sigma <- 1.5; N <- 10
set.seed(1004)
y <- rnorm(mean=2.3, sd=sqrt(sigma), n=N)
s2 <- 1/(N-1)*sum((y-mean(y))^2)

invchi=((N-1)*s2)/(rchisq(1000, df=N-1))

bhat <- mean(rnorm(mean=mean(y),
                    sd=sqrt(invchi/N), n=1000))
```

- We can sample independently since σ^2 is independent of β in the joint posterior
- (Also, note how the Inv- χ^2 is nicely sampled from the χ^2 distribution)

Prediction on \tilde{y}

Basic Setup

Simple
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- We can make predictions on future observations of y using posterior inference
- We make use of the posterior predictive distribution:

$$\begin{aligned} p(\tilde{y}|y) &= \int p(\tilde{y}, \theta|y) d\theta \\ &= \int p(\tilde{y}|\theta) p(\theta|y) d\theta \end{aligned}$$

- For our above example:

$$p(\tilde{y}|y) \propto \int \exp\left\{-\frac{1}{2\sigma^2} (\tilde{y} - \beta)^2\right\} \exp\left\{-\frac{1}{2\tau_*^2} (\beta - \mu_*)^2\right\} d\theta$$

- Can show that $E[\tilde{y}|y] = \mu_*$ and $\text{Var}(\tilde{y}|y) = \sigma^2 + \tau_*^2$
- Useful for model checking with out-of-sample y