

Hierarchical Bayesian modeling

Tom Loredo

Cornell Center for Astrophysics and Planetary Science

<http://www.astro.cornell.edu/staff/loredo/bayes/>

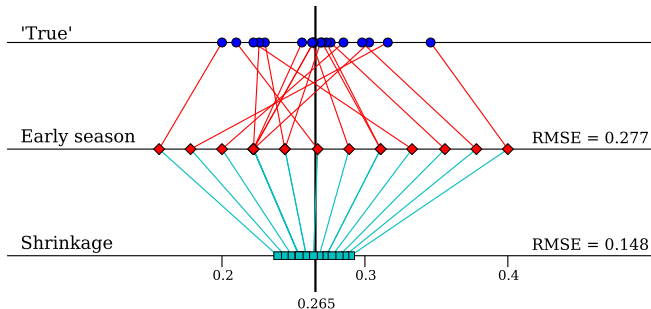
SAMSI ASTRO — 19 Oct 2016

1970 baseball averages

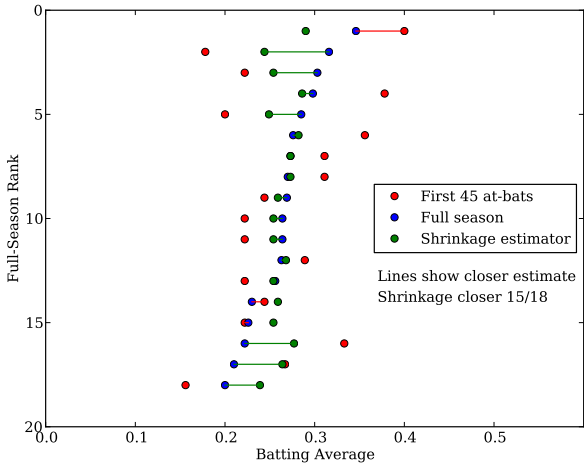
Efron & Morris looked at batting averages of baseball players who had $N = 45$ at-bats in May 1970 — ‘large’ N & includes Roberto Clemente (outlier!)

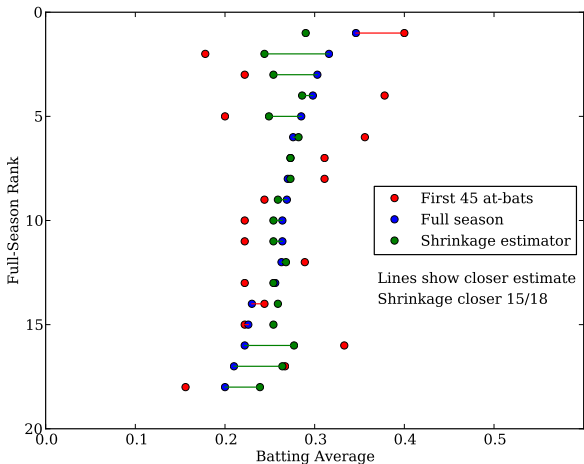
Red = n/N maximum likelihood estimates of true averages

Blue = Remainder of season, $N_{\text{rmldr}} \approx 9N$



Cyan = James-Stein estimator: nonlinear, correlated, biased
But *better!*





Theorem (independent Gaussian setting): In dimension $\gtrsim 3$, shrinkage estimators always beat independent MLEs in terms of expected RMS error

“The single most striking result of post-World War II statistical theory”
 — Brad Efron

All 18 players are *humans playing baseball*—they are members of a population, not arbitrary, unrelated binomial random number generators!

In the absence of data about player i , we may use the performance of the other players to guide a guess about that player's performance—they provide *indirect evidence* (Efron) about player i

But information that is relevant in the absence of data for i remains relevant when we additionally obtain that data; shrinkage estimators account for this

There is “mustering and *borrowing of strength*” (Tukey) across the population

Hierarchical Bayesian modeling is the most flexible framework for generalizing this lesson; *empirical Bayes* is an approximate version with a straightforward frequentist interpretation

Agenda

① Basic Bayes recap

② Key idea in a nutshell

③ Going deeper

Joint distributions and DAGs

Conditional dependence/independence

Example: Binomial prediction

Beta-binomial model

Point estimation and shrinkage

Gamma-Poisson model & Stan

Algorithms

Bayesian inference in one slide

Probability as generalized logic

Probability quantifies the *strength of arguments*

To appraise hypotheses, calculate probabilities for arguments from data and modeling assumptions to each hypothesis

Use *all* of probability theory for this

Bayes's theorem

$$p(\text{Hypothesis} \mid \text{Data}) \propto p(\text{Hypothesis}) \times p(\text{Data} \mid \text{Hypothesis})$$

Data *change* the support for a hypothesis \propto ability of hypothesis to *predict* the data

Law of total probability

$$p(\text{Hypotheses} \mid \text{Data}) = \sum p(\text{Hypothesis} \mid \text{Data})$$

The support for a *compound/composite* hypothesis must account for all the ways it could be true

Bayes's theorem

\mathcal{C} = context, initial set of premises

Consider $P(H_i, D_{\text{obs}}|\mathcal{C})$ using the product rule:

$$\begin{aligned}P(H_i, D_{\text{obs}}|\mathcal{C}) &= P(H_i|\mathcal{C}) P(D_{\text{obs}}|H_i, \mathcal{C}) \\ &= P(D_{\text{obs}}|\mathcal{C}) P(H_i|D_{\text{obs}}, \mathcal{C})\end{aligned}$$

Solve for the *posterior probability* (expands the premises!):

$$P(H_i|D_{\text{obs}}, \mathcal{C}) = P(H_i|\mathcal{C}) \frac{P(D_{\text{obs}}|H_i, \mathcal{C})}{P(D_{\text{obs}}|\mathcal{C})}$$

Theorem holds for any propositions, but for hypotheses & data the factors have names:

posterior \propto *prior* \times *likelihood*

norm. const. $P(D_{\text{obs}}|\mathcal{C}) =$ *prior predictive*

Law of Total Probability (LTP)

Consider exclusive, exhaustive $\{B_i\}$ (\mathcal{C} asserts one of them must be true),

$$\begin{aligned}\sum_i P(A, B_i | \mathcal{C}) &= \sum_i P(B_i | A, \mathcal{C}) P(A | \mathcal{C}) = P(A | \mathcal{C}) \\ &= \sum_i P(B_i | \mathcal{C}) P(A | B_i, \mathcal{C})\end{aligned}$$

If we do not see how to get $P(A | \mathcal{P})$ directly, we can find a set $\{B_i\}$ and use it as a “basis”—*extend the conversation*:

$$P(A | \mathcal{C}) = \sum_i P(B_i | \mathcal{C}) P(A | B_i, \mathcal{C})$$

If our problem already has B_i in it, we can use LTP to get $P(A | \mathcal{C})$ from the joint probabilities—*marginalization*:

$$P(A | \mathcal{C}) = \sum_i P(A, B_i | \mathcal{C})$$

Example: Take $A = D_{\text{obs}}$, $B_i = H_i$; then

$$\begin{aligned} P(D_{\text{obs}}|\mathcal{C}) &= \sum_i P(D_{\text{obs}}, H_i|\mathcal{C}) \\ &= \sum_i P(H_i|\mathcal{C})P(D_{\text{obs}}|H_i, \mathcal{C}) \end{aligned}$$

prior predictive for $D_{\text{obs}} =$ Average likelihood for H_i
(a.k.a. *marginal likelihood*)

Parameter Estimation

Problem statement

\mathcal{C} = Model M with parameters θ (+ any add'l info)

H_i = statements about θ ; e.g. " $\theta \in [2.5, 3.5]$," or " $\theta > 0$ "

Probability for any such statement can be found using a *probability density function* (PDF) for θ :

$$\begin{aligned} P(\theta \in [\theta, \theta + d\theta] | \dots) &= f(\theta)d\theta \\ &= p(\theta | \dots)d\theta \end{aligned}$$

Posterior probability density

$$p(\theta | D, M) = \frac{p(\theta | M) \mathcal{L}(\theta)}{\int d\theta p(\theta | M) \mathcal{L}(\theta)}$$

Summaries of posterior

- “Best fit” values:
 - ▶ Mode, $\hat{\theta}$, maximizes $p(\theta|D, M)$
 - ▶ Posterior mean, $\langle \theta \rangle = \int d\theta \theta p(\theta|D, M)$
- Uncertainties:
 - ▶ Credible region Δ of probability C :
 $C = P(\theta \in \Delta|D, M) = \int_{\Delta} d\theta p(\theta|D, M)$
Highest Posterior Density (HPD) region has $p(\theta|D, M)$ higher inside than outside
 - ▶ Posterior standard deviation, variance, covariances
- Marginal distributions
 - ▶ Interesting parameters ϕ , nuisance parameters η
 - ▶ Marginal dist'n for ϕ : $p(\phi|D, M) = \int d\eta p(\phi, \eta|D, M)$

Many Roles for Marginalization

Eliminate nuisance parameters

$$p(\phi|D, M) = \int d\eta p(\phi, \eta|D, M)$$

Propagate uncertainty

Model has parameters θ ; what can we infer about $F = f(\theta)$?

$$\begin{aligned} p(F|D, M) &= \int d\theta p(F, \theta|D, M) = \int d\theta p(\theta|D, M) p(F|\theta, M) \\ &= \int d\theta p(\theta|D, M) \delta[F - f(\theta)] \quad [\textit{single-valued case}] \end{aligned}$$

Prediction

Given a model with parameters θ and present data D , predict future data D' (e.g., for *experimental design*):

$$p(D'|D, M) = \int d\theta p(D', \theta|D, M) = \int d\theta p(\theta|D, M) p(D'|\theta, M)$$

Model comparison

Marginal likelihood for model M_i :

$$Z_i \equiv p(D|M_i) = \int d\theta_i p(\theta_i|M) \mathcal{L}_i(\theta_i)$$

Bayes factor $B_{ij} \equiv Z_i/Z_j$

Can write $Z_i = \mathcal{L}_i(\hat{\theta}_i) \cdot \Omega_i$ with Ockham factor

$\Omega_i \approx \delta\theta/\Delta\theta = (\text{posterior volume})/(\text{prior volume})$

Hierarchical modeling, aka...

- Graphical models — Hierarchical and other structures
- Multilevel models — In regression, linear model settings
- Bayesian networks (Bayes nets) — In AI/ML settings

Agenda

① Basic Bayes recap

② Key idea in a nutshell

③ Going deeper

Joint distributions and DAGs

Conditional dependence/independence

Example: Binomial prediction

Beta-binomial model

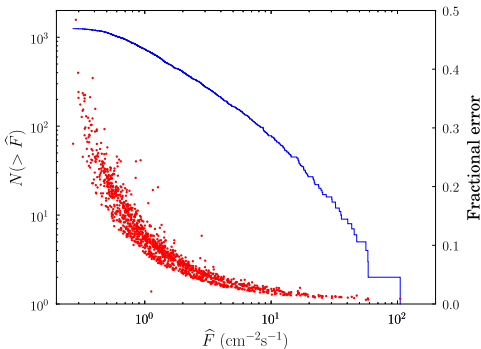
Point estimation and shrinkage

Gamma-Poisson model & Stan

Algorithms

Motivation: Measurement error in surveys

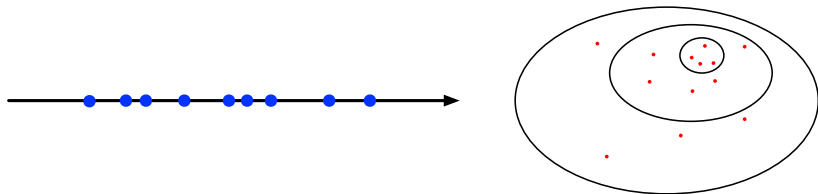
BATSE GRB peak flux estimates



- *Selection effects* (truncation, censoring) — *obvious* (usually)
Typically treated by “correcting” data
Most sophisticated: product-limit estimators
- *“Scatter” effects* (measurement error, etc.) — *insidious*
Typically ignored (average out??? — *No!*)

Accounting For Measurement Error

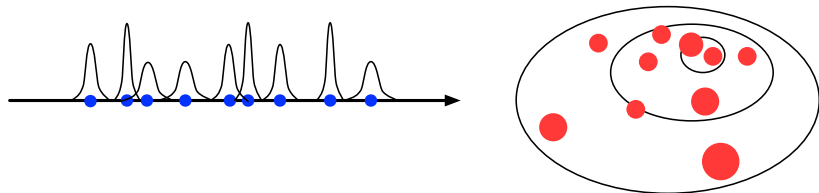
Suppose $f(x|\theta)$ is a distribution for an observable, x (scalar or vector, $\vec{x} = (x, y, \dots)$); and θ is unknown



From N precisely measured samples, $\{x_i\}$, we can infer θ from

$$\mathcal{L}(\theta) \equiv p(\{x_i\}|\theta) = \prod_i f(x_i|\theta)$$
$$p(\theta|\{x_i\}) \propto p(\theta)\mathcal{L}(\theta) = p(\theta, \{x_i\})$$

But what if the x data are *noisy*, $D_i = \{x_i + \epsilon_i\}$?



$\{x_i\}$ are now *uncertain (latent/hidden/incidental) parameters*

We should somehow incorporate $\ell_i(x_i) = p(D_i|x_i)$

The joint PDF for *everything* is

$$\begin{aligned} p(\theta, \{x_i\}, \{D_i\}) &= p(\theta) p(\{x_i\}|\theta) p(\{D_i\}|\{x_i\}) \\ &= p(\theta) \prod_i f(x_i|\theta) \ell_i(x_i) \end{aligned}$$

The conditional (posterior) PDF for the unknowns is

$$p(\theta, \{x_i\}|\{D_i\}) = \frac{p(\theta, \{x_i\}, \{D_i\})}{p(\{D_i\})} \propto p(\theta, \{x_i\}, \{D_i\})$$

$$\begin{aligned} p(\theta, \{x_i\} | \{D_i\}) &\propto p(\theta, \{x_i\}, \{D_i\}) \\ &= p(\theta) \prod_i f(x_i | \theta) \ell_i(x_i) \end{aligned}$$

Marginalize over $\{x_i\}$ to summarize inferences for θ

Marginalize over θ to summarize inferences for $\{x_i\}$

Key point: *Maximizing over x_i (i.e., just using best-fit \hat{x}_i) and integrating over x_i can give very different results!*

(See Loredo (2004) for tutorial examples)

To estimate x_1 :

$$\begin{aligned} p(x_1 | \{x_2, \dots\}) &= \int d\theta p(\theta) f(x_1 | \theta) l_1(x_1) \times \prod_{i=2}^N \int dx_i f(x_i | \theta) l_i(x_i) \\ &= l_1(x_1) \int d\theta p(\theta) f(x_1 | \theta) \mathcal{L}_{m, \mathbf{I}}(\theta) \\ &\approx l_1(x_1) f(x_1 | \hat{\theta}_{\mathbf{I}}) \end{aligned}$$

with $\hat{\theta}_{\mathbf{I}}$ determined by the remaining data

$f(x_1 | \hat{\theta}_{\mathbf{I}})$ behaves like a “prior” that shifts the x_1 estimate away from the peak of $l_1(x_1)$; each member’s prior depends on all of the rest of the data \rightarrow shrinkage

[For astronomers: This generalizes the corrections derived by Eddington, Malmquist and Lutz-Kelker (sans selection effects)]

Agenda

① Basic Bayes recap

② Key idea in a nutshell

③ **Going deeper**

- Joint distributions and DAGs

- Conditional dependence/independence

- Example: Binomial prediction

- Beta-binomial model

- Point estimation and shrinkage

- Gamma-Poisson model & Stan

- Algorithms

Joint and conditional distributions

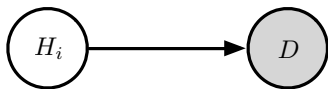
Bayesian inference is largely about the interplay between *joint* and *conditional* distributions for related quantities

Ex: Bayes's theorem relating hypotheses and data ($||C$):

$$P(H_i|D) = \frac{P(H_i)P(D|H_i)}{P(D)} = \frac{P(H_i, D)}{P(D)} = \frac{\text{joint for everything}}{\text{marginal for knowns}}$$

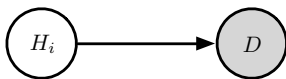
The usual form identifies an available factorization of the joint

Express this via a *directed acyclic graph* (DAG):

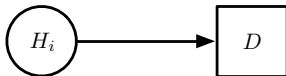


Joint distribution structure as a graph

- Graph = *nodes/vertices* connected by *edges/links*
- Circular/square nodes/vertices = a priori uncertain quantities (gray/square = becomes known as data)
- Directed edges specify conditional dependence
- Absence of an edge indicates conditional *in*dependence
→ *the most important edges are the missing ones*



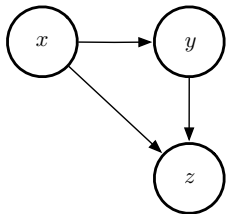
OR



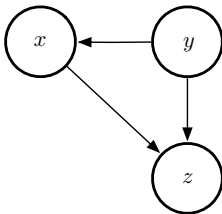
$$P(H_i, D) = P(H_i) \times P(D|H_i)$$

$$p(x, y, z)$$

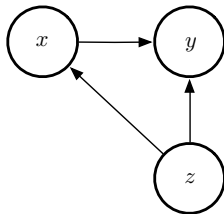
$$p(x)p(y|x)p(z|x, y)$$



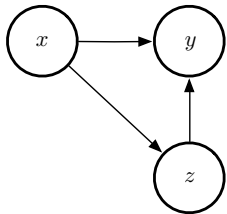
$$p(y)p(x|y)p(z|y, x)$$



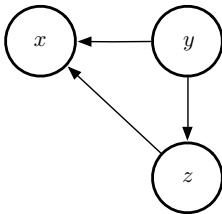
$$p(z)p(x|z)p(y|z, x)$$



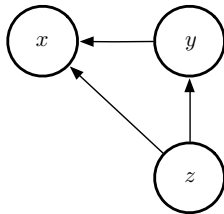
$$p(x)p(z|x)p(y|x, z)$$



$$p(y)p(z|y)p(x|y, z)$$

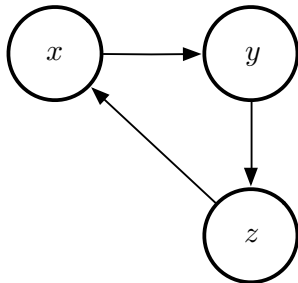


$$p(z)p(y|z)p(x|z, y)$$



Cycles not allowed

$$p(x|z) \times p(y|x) \times p(z|y)?$$

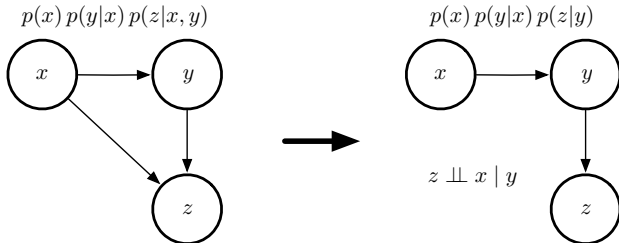


Conditional independence

Suppose for the problem at hand z is independent of x when y is known:

$$p(z|x, y) = p(z|y)$$

“ z is *conditionally independent* of x , given y ”: $z \perp\!\!\!\perp x \mid y$



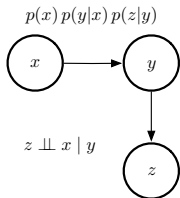
Absence of an edge indicates conditional *independence*

Missing edges indicate simplification in structure

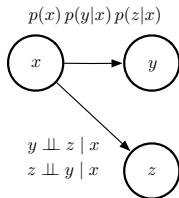
→ *the most important edges are the missing ones*

DAGs with missing edges

Conditional independence

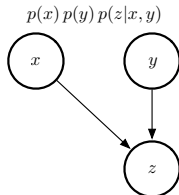


“Causal chain”



“Common cause”

Conditional dependence



“Common effects”

Conditional vs. complete independence

“z is *conditionally* independent of x, given y”

≠

“z is independent of x”

(Complete) independence between z and x (“z $\perp\!\!\!\perp$ x”) would imply:

$$p(z|x) = p(z) \quad (\text{i.e., not a function of } x)$$

Conditional independence *given y* (“z $\perp\!\!\!\perp$ x | y”) is weaker:

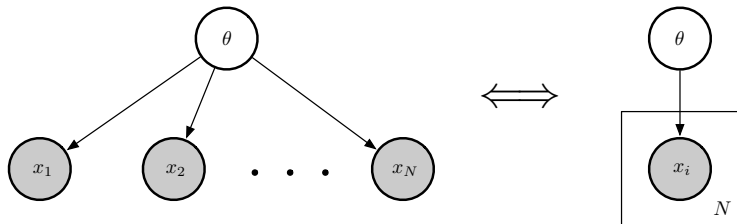
$$\begin{aligned} p(z|x) &= \int dy p(z, y|x) \\ &= \int dy p(y|x)p(z|x, y) \\ &= \int dy p(y|x)p(z|y) \quad \text{since } z \perp\!\!\!\perp x | y \end{aligned}$$

Although x drops out of the last factor, x dependence remains in $p(y|x)$

x *does* provide information about z, but it only does so through the information it provides about y (which directly influences z)

Bayes's theorem with IID samples

For model with parameters θ predicting data $D = \{x_i\}$ that are IID given θ :



$$p(\theta, D) = p(\theta)p(\{x_i\}|\theta) = p(\theta) \prod_{i=1}^N p(x_i|\theta)$$

To find the posterior for the unknowns (θ), divide the joint by the marginal for the knowns ($\{x_i\}$):

$$p(\theta|\{x_i\}) = \frac{p(\theta) \prod_{i=1}^N p(x_i|\theta)}{p(\{x_i\})} \quad \text{with} \quad p(\{x_i\}) = \int d\theta p(\theta) \prod_{i=1}^N p(x_i|\theta)$$

Binomial counts



■ ■ ■ n_1 heads in N flips



■ ■ ■ n_2 heads in N flips

Suppose we know n_1 and want to predict n_2

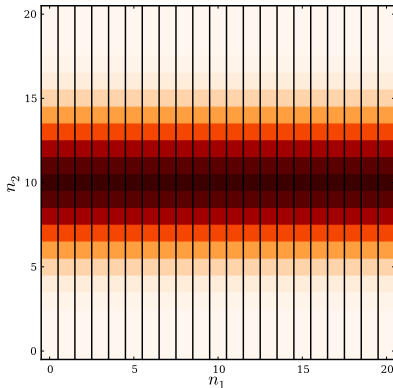
Predicting binomial counts — known α

Success probability $\alpha \rightarrow p(n|\alpha) = \frac{N!}{n!(N-n)!} \alpha^n (1-\alpha)^{N-n} \quad || N$

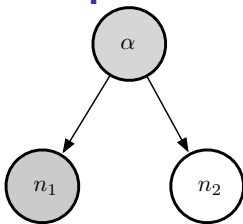
Consider two successive runs of $N = 20$ trials, *known* $\alpha = 0.5$

$$p(n_2 | n_1, \alpha) = p(n_2 | \alpha) \quad || N$$

n_1 and n_2 are *conditionally independent*



DAG for binomial prediction — known α



$$p(\alpha, n_1, n_2) = p(\alpha)p(n_1|\alpha)p(n_2|\alpha)$$

$$\begin{aligned} p(n_2|\alpha, n_1) &= \frac{p(\alpha, n_1, n_2)}{p(\alpha, n_1)} \\ &= \frac{p(\alpha)p(n_1|\alpha)p(n_2|\alpha)}{p(\alpha)p(n_1|\alpha)\sum_{n_2} p(n_2|\alpha)} \\ &= p(n_2|\alpha) \end{aligned}$$

Knowing α lets you predict each n_i , independently

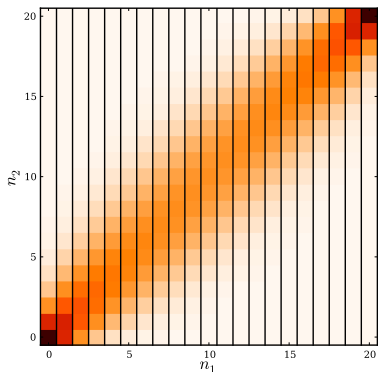
Predicting binomial counts — uncertain α

Consider the same setting, but with α *uncertain*

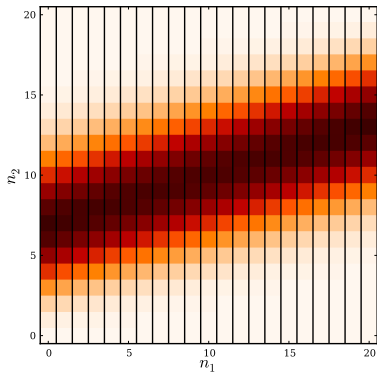
Outcomes are *physically* independent, but n_1 tells us about $\alpha \rightarrow$ outcomes are *marginally dependent* (see Lec 12 for calculation):

$$p(n_2|n_1, N) = \int d\alpha p(\alpha, n_2|n_1, N) = \int d\alpha p(\alpha|n_1, N) p(n_2|\alpha, N)$$

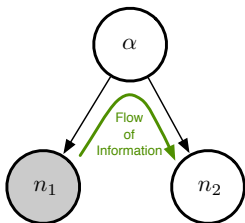
Flat prior on α



Prior: $\alpha = 0.5 \pm 0.1$



DAG for binomial prediction



$$p(\alpha, n_1, n_2) = p(\alpha)p(n_1|\alpha)p(n_2|\alpha)$$

From joint to conditionals:

$$p(\alpha|n_1, n_2) = \frac{p(\alpha, n_1, n_2)}{p(n_1, n_2)} = \frac{p(\alpha)p(n_1|\alpha)p(n_2|\alpha)}{\int d\alpha p(\alpha)p(n_1|\alpha)p(n_2|\alpha)}$$

$$p(n_2|n_1) = \frac{\int d\alpha p(\alpha, n_1, n_2)}{p(n_1)}$$

Observing n_1 lets you learn about α

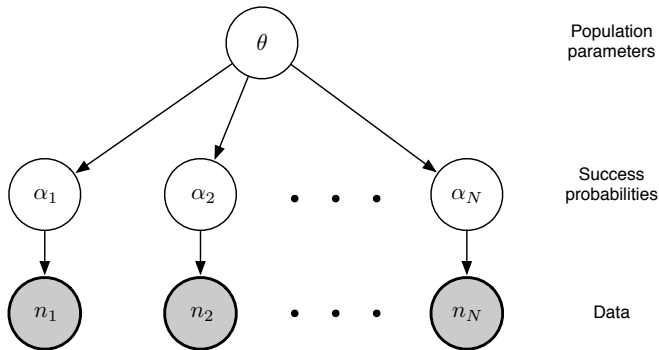
Knowledge of α affects predictions for $n_2 \rightarrow$ dependence on n_1

A population of coins/flippers



Each flipper+coin flips different number of times

- What do we learn about the *population* of coins—the distribution of α s?
- How does population membership effect inference for a single coin's α ?



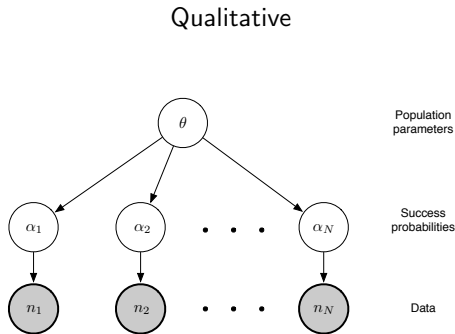
$$\begin{aligned}
 p(\theta, \{\alpha_i\}, \{n_i\}) &= \pi(\theta) \prod_i p(\alpha_i | \theta) p(n_i | \alpha_i) \\
 &= \pi(\theta) \prod_i p(\alpha_i | \theta) \ell_i(\alpha_i)
 \end{aligned}$$

Terminology: θ are *hyperparameters*, $\pi(\theta)$ is the *hyperprior*

A simple multilevel model: beta-binomial

Goals:

- Learn a population-level “prior” by pooling data
- Account for population membership in member inferences



Quantitative

$$\theta = (a, b) \text{ or } (\mu, \sigma)$$

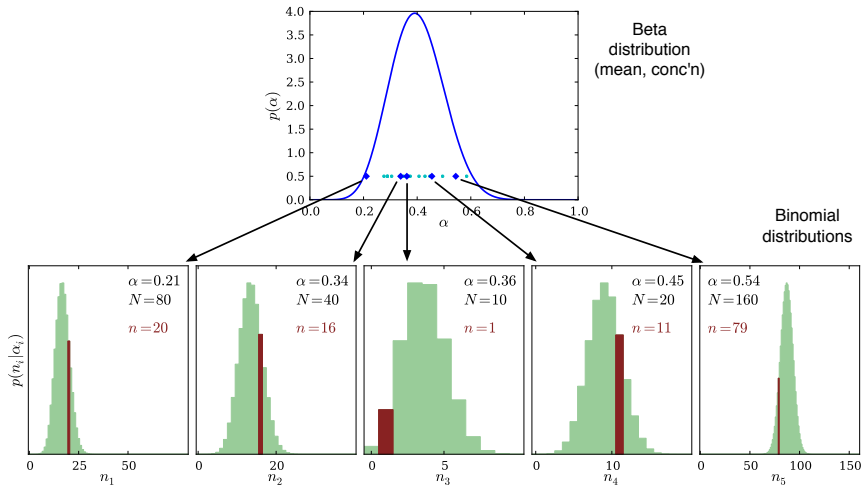
$$\pi(\theta) = \text{Flat}(\mu, \sigma)$$

$$p(\alpha_i | \theta) = \text{Beta}(\alpha_i | \theta)$$

$$p(n_i | \alpha_i) = \binom{N_i}{n_i} \alpha_i^{n_i} (1 - \alpha_i)^{N_i - n_i}$$

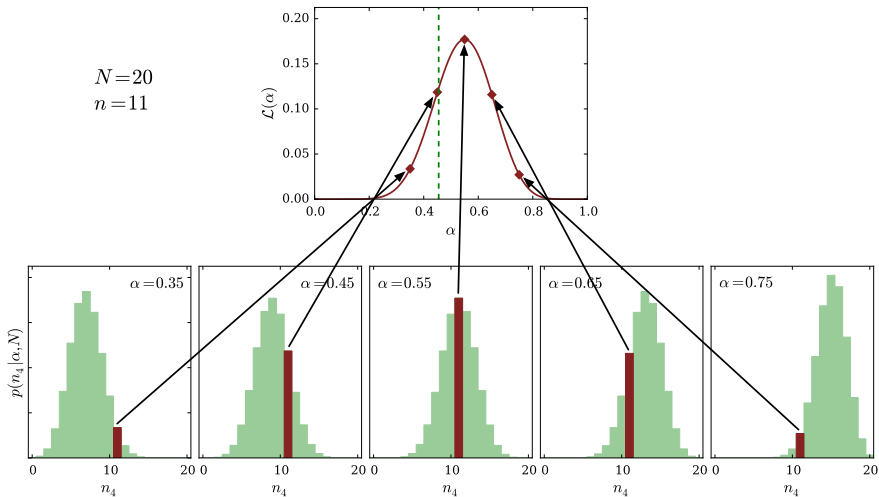
$$\begin{aligned} p(\theta, \{\alpha_i\}, \{n_i\}) &= \pi(\theta) \prod_i p(\alpha_i | \theta) p(n_i | \alpha_i) \\ &= \pi(\theta) \prod_i p(\alpha_i | \theta) \ell_i(\alpha_i) \end{aligned}$$

Generating the population & data

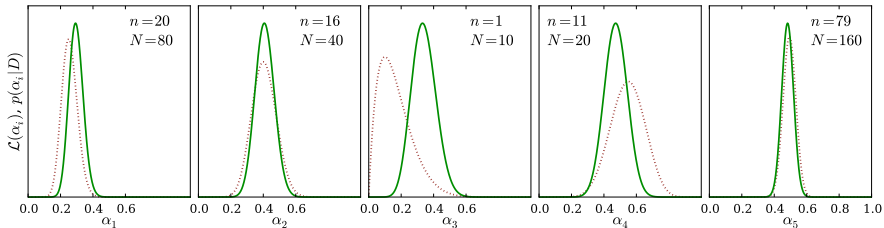
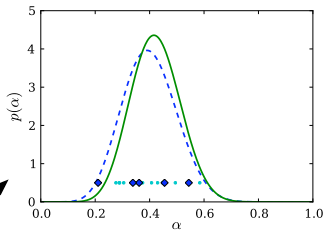


Likelihood function for one member's α

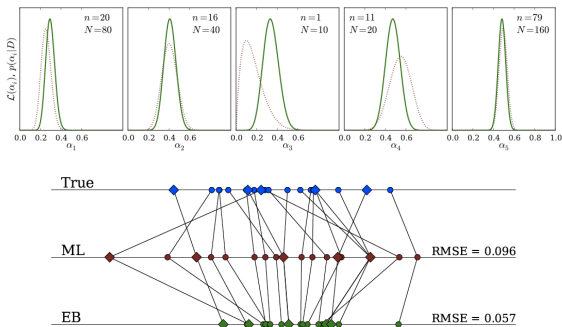
$N=20$
 $n=11$



Learning the population distribution



Lower level estimates



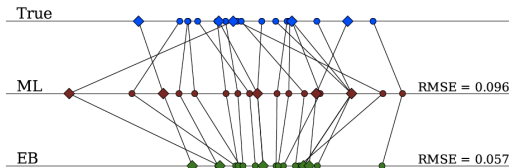
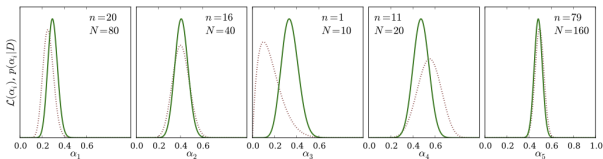
Two approaches

- **Hierarchical Bayes (HB):** Calculate marginals

$$p(\alpha_j|\{n_i\}) \propto \int d\theta \pi(\theta) \prod_{i \neq j} \int d\alpha_i p(\alpha_i|\theta) p(n_i|\alpha_i)$$

- **Empirical Bayes (EB):** Plug in an optimum $\hat{\theta}$ and estimate $\{\alpha_i\}$
View as approximation to HB, or a frequentist procedure that estimates a prior from the data

Lower level estimates



Bayesian outlook

- Marginal posteriors are *narrower* than likelihoods
- Point estimates tend to be closer to true values than MLEs (averaged across the population)
- Joint distribution for $\{\alpha_i\}$ is *dependent*

Frequentist outlook

- Point estimates are biased
- Reduced variance → estimates are closer to truth on average (lower MSE in repeated sampling)
- Bias for one member estimate depends on data for all other members

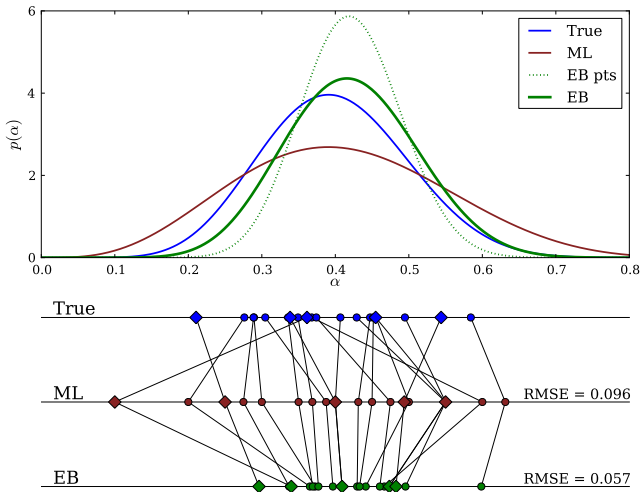
Lingo

- Estimates *shrink* toward prior/population mean
- Estimates “muster and *borrow strength*” across population (Tukey’s phrase); increases accuracy and precision of estimates
- Efron* describes shrinkage as a consequence of accounting for *indirect evidence*

*Bradley Efron (2010): “The Future of Indirect Evidence”

Beware of point estimates!

Population and member estimates



Competing data analysis goals

“Shrunken” member estimates provide improved & reliable estimate for population member properties

But they are *under-dispersed* in comparison to the true values → not optimal for estimating *population* properties*

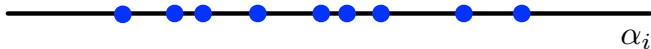
No point estimates of member properties are good for all tasks!

We should view population data tables/catalogs as providing
descriptions of member likelihood functions,
not “estimates with errors”

*Louis (1984); Eddington noted this in 1940!

Measurement error perspective

If the data provided *precise* $\{\alpha_i\}$ values (coin measurements, flip physics), we could easily model them as points drawn from a (beta) population PDF with params θ :

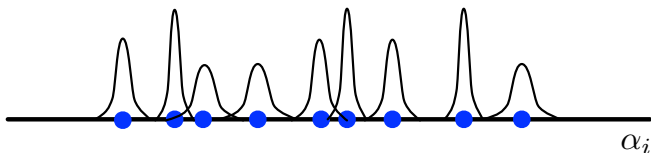


$$D = \{\alpha_i\}$$

$$\begin{aligned} p(D|\theta) &= \prod_i p(\alpha_i|\theta) \\ &= \prod_i \text{Beta}(\alpha_i|\theta) \end{aligned}$$

(A *binomial point process*)

Here the finite number of flips provide *noisy measurements of each* α_i , described by the member likelihood functions $\ell_i(\alpha_i)$;



$$D = \{n_i\}$$

$$\begin{aligned} p(D|\theta) &= \prod_i \int d\alpha_i p(D, \{\alpha_i\}|\theta) \\ &= \prod_i \int d\alpha_i p(\alpha_i|\theta) p(n_i|\theta) \\ &= \prod_i \int d\alpha_i \text{Beta}(\alpha_i|\theta) \text{Binom}(n_i|\theta) \end{aligned}$$

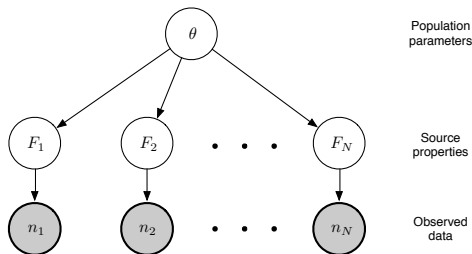
This is a prototype for *measurement error problems*

Another conjugate MLM: Gamma-Poisson

Goal: Learn a rate dist'n from count data
(E.g., learn a star or galaxy brightness dist'n from photon counts)

Qualitative

Quantitative



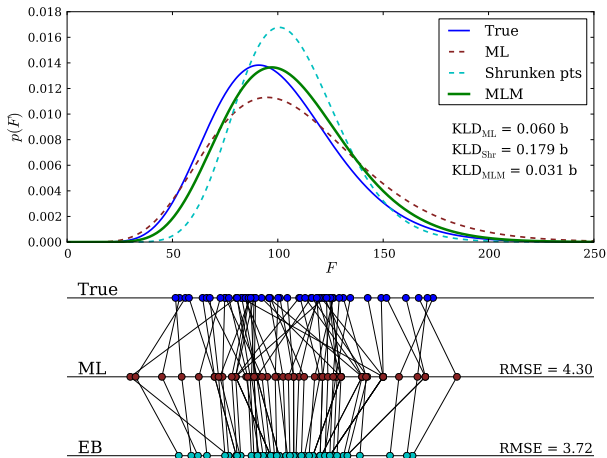
$$\theta = (\alpha, s) \text{ or } (\mu, \sigma)$$

$$\pi(\theta) = \text{Flat}(\mu, \sigma)$$

$$p(F_i|\theta) = \text{Gamma}(F_i|\theta)$$

$$p(n_i|F_i) = \text{Pois}(n_i|\epsilon_i F_i)$$

Gamma-Poisson population and member estimates



Simulations: $N = 60$ sources from gamma with $\langle F \rangle = 100$ and $\sigma_F = 30$;
exposures spanning dynamic range of $\times 16$

Algorithms

Consider the posterior PDF for θ and $\{\alpha_i\}$ in the beta-binomial MLM:

$$p(\theta, \{\alpha_i\} | \{n_i\}) \propto \pi(\theta) \prod_{i=1}^{N_{\text{mem}}} \text{Beta}(\alpha_i | \theta) \text{Binom}(n_i | \alpha_i)$$

For each member, the Beta \times Binom factor is \propto a beta distribution for α_i ; but as a function of θ (e.g., (a, b) or (μ, σ)) it is not simple

The full posterior has a product of N_{mem} such factors specifying its θ dependences \Rightarrow *even for a conjugate model for the lower levels, the overall model is typically analytically intractable*

Two approaches exploit *conditional independence of lower-level parameters*

Member marginalization

- Analytically or numerically integrate over $\{x_i\}$ \rightarrow explore the reduced-dimension marginal for θ via MCMC
 $\rightarrow \{\theta_i\} \sim p(\theta|D)$
- If x_i are of interest, sample them from their conditionals, conditioned on θ_i :
 - ▶ Pick a θ from $\{\theta_i\}$
 - ▶ Draw $\{x_i\}$ by *independent* sampling from their conditionals (give θ)
 - ▶ Iterate

GPUs can accelerate this for application to large datasets

Only useful for low-dimensional latent parameters x_i

Metropolis-within-Gibbs algorithm

Block the full parameter space:

- Block of m population parameters, θ
- N blocks of lower level (latent) parameters, x_i

Get posterior samples by iterating back and forth between:

- m -D Metropolis-Hastings sampling of θ from $p(\theta|\{x_i\}, D)$

This requires a problem-specific proposal distribution

- N *independent* samples of x_i from the conditional $p(x_i|\theta, D_i)$

This can often exploit conjugate structure

E.g., Beta-binomial: $\alpha_i \sim \text{Beta}(\alpha_i|\theta)$ $\text{Binom}(n_i|\alpha_i)$,
which is just a Beta for α_i

MWG explicitly displays the feedback between population and member inference