

A 250-Year Argument

Belief, Behavior, and the Bootstrap

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Bayes Rule (1763)

- 250th anniversary in 2013
- Always influential, usually controversial

$$\pi(\theta|\mathbf{x}) = \pi(\theta)f(\mathbf{x}|\theta)/c(\mathbf{x})$$

\mathbf{x} = observed data

$\pi(\theta)$ = prior belief for parameter θ

$f(\mathbf{x}|\theta)$ = likelihood

$c(\mathbf{x})$ = scaling constant

$\pi(\theta|\mathbf{x})$ = posterior belief

- How to choose $\pi(\theta)$?

The Physicist's Twins

- **Sonogram** “twin boys” ● Prob. *identical*, not *fraternal*?
- **Doctor** “population probability *identical* = 1/3”

- **Bayes Rule**

$$\frac{\pi(\text{ident} | \text{sono})}{\pi(\text{frat} | \text{sono})} = \left[\frac{\pi(\text{ident})}{\pi(\text{frat})} \right] \left[\frac{f(\text{sono} | \text{ident})}{f(\text{sono} | \text{frat})} \right]$$

- **Prior Odds** = $\frac{1/3}{2/3} = \frac{1}{2}$ ● **Likelihood Ratio** = $\frac{1}{1/2} = 2$

- **Posterior Odds** = $\left[\frac{1}{2} \right] \cdot [2] = 1$ ● $\pi(\text{ident} | \text{sono}) = 50\%$

The Complete Population of Twins

		Sexes	
		<i>Same</i>	<i>Different</i>
Twins	<i>Identical</i>	1/3	0
	<i>Fraternal</i>	1/3	1/3

sonogram

1/3 }
2/3 } Doctor

The Doctor's Beliefs

- Bayes Prior $\pi(\theta)$: Infinite list of relevant past cases (millions of prior twins)
- What if only 3 cases, one *identical* and two *fraternal*?
- **Harold Jeffreys** (1930s): theory of **uninformative priors**
- Dominates current Bayesian practice
- *Statistician can “always be a Bayesian”*

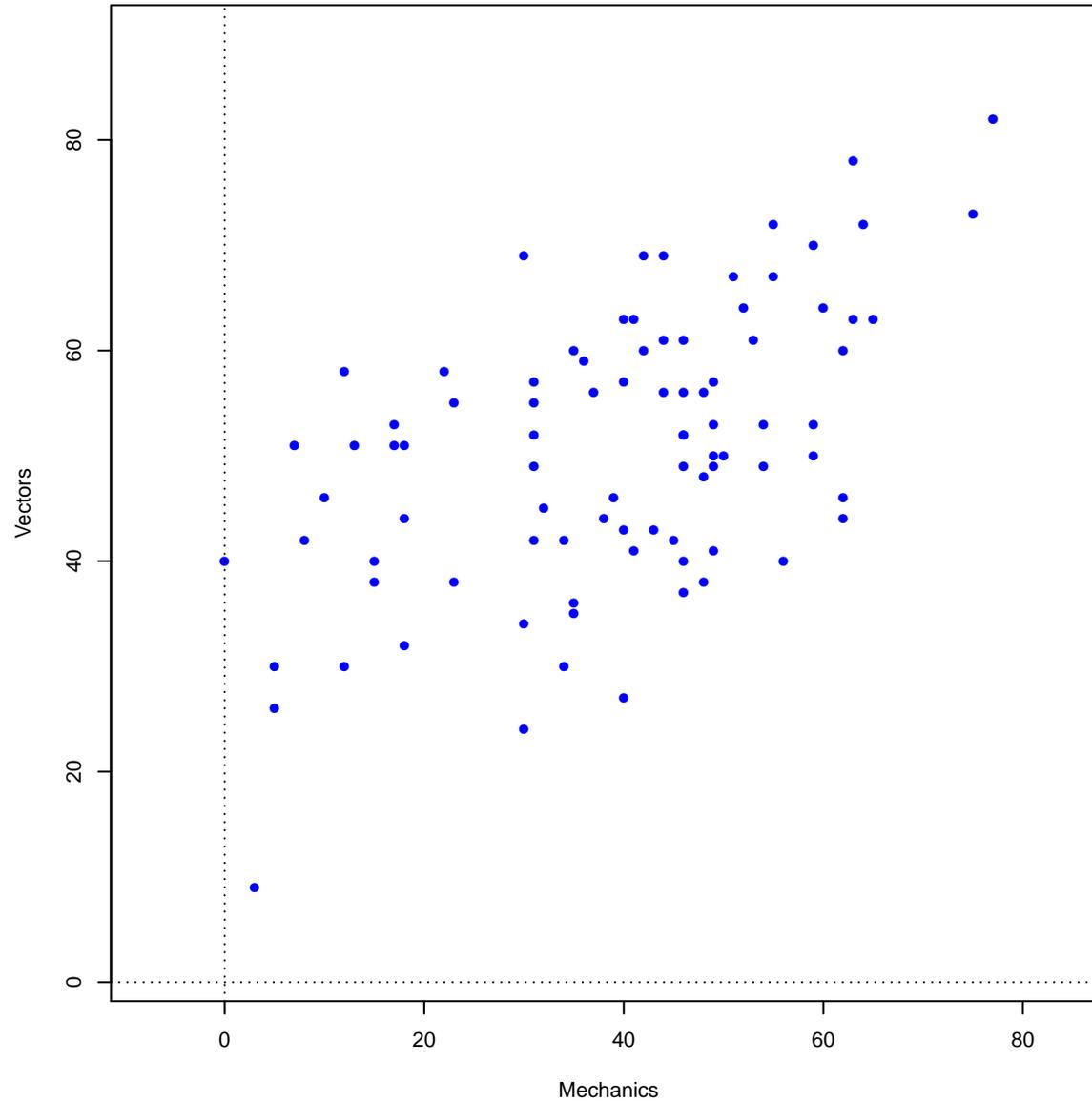
Jeffreys and the Twins

- θ = population proportion of *identical* twins
[Doctor: “ $\theta = 1/3$ ”]
- Uninformative hyperprior $\pi(\theta) \propto \theta^{-1}(1 - \theta)^{-1}$
for θ in $(0,1)$
- *Doctor should believe*: $\pi(\theta|\text{his experience}) = 2 \cdot (1 - \theta)$
[“Beta(1,2)”]
- Then $\pi(\text{ident} | \text{sono}, \text{Doc's experience}) = 50\%$
- Same “50%”?

Frequentist Inference (“Behavior”)

- No prior beliefs “ $\pi(\theta)$ ”
- Specific estimation rule $\hat{\theta} = s(\boldsymbol{x})$
[test, prediction, model selection, etc.]
- Judge $s(\boldsymbol{x})$ by its behavior in repeated use
 - Bayes infinite past experience
 - Frequentist infinite future usage

Scores on two tests for 88 students;
Sample correlation coefficient = .553



Correlation Example (Mardia, Kent, and Bobby)

- **Data** $n = 88$ students:

Scores on two exams “mechanics” and “vectors”

$$\mathbf{x} = (x_1, x_2, \dots, x_{88}) \quad \text{with} \quad x_i = (x_{i1}, x_{i2})$$

- *Parameter of interest:* correlation coefficient $\rho = \text{cor}(\text{mec}, \text{vec})$

- *Estimate* $\hat{\rho} = s(\mathbf{x})$

- Sample correlation coefficient = $0.553 \pm ?$

- **Bivariate normal assumption**

$$x_i \stackrel{\text{ind}}{\sim} \mathcal{N}_2(\mu, \Sigma)$$

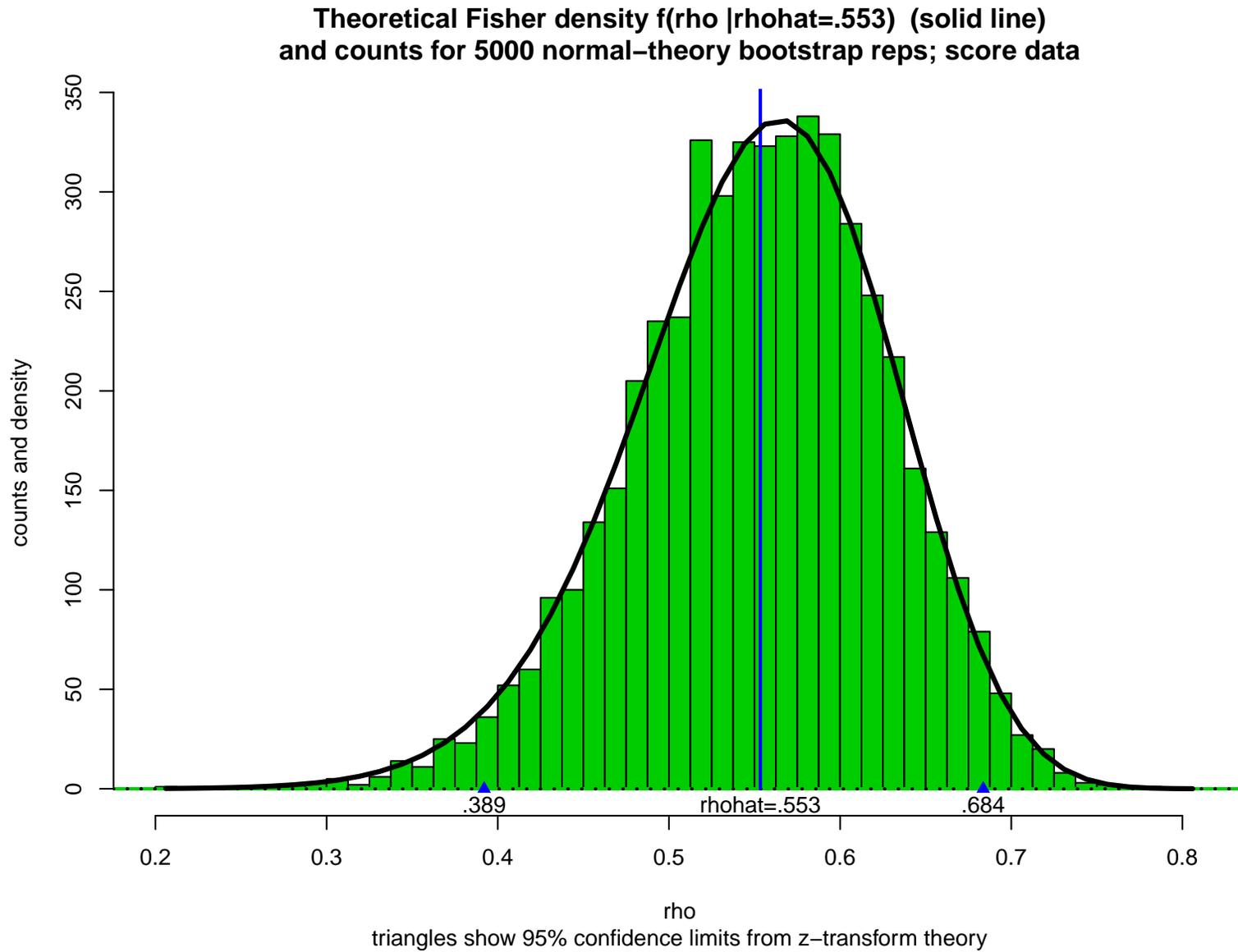
for $i = 1, 2, \dots, 88$

Fisher: Normal Theory for $\hat{\rho}$ (1915)

- *Density* $f_{\rho}(\hat{\rho})$
- $\text{sd}_{\rho}(\hat{\rho}) \doteq (1 - \rho^2) / \sqrt{n - 3}$ [= 0.075 for $\rho = 0.553$]
- **Transformation** $\hat{\tau} = \frac{1}{2} \log \frac{1 + \hat{\rho}}{1 - \hat{\rho}}$ • $\hat{\tau} \sim \mathcal{N}\left(\tau, \frac{1}{n - 3}\right)$
- 95% confidence interval for ρ : transform $\hat{\tau} \pm 1.96 \frac{1}{\sqrt{n - 3}}$
back to ρ scale:

$$\rho \in [0.389, 0.684]$$

- **Bayes** single probability distribution
- **Frequentist** many distributions



Parametric Bootstrap Calculations

- Assume $x_i \stackrel{\text{ind}}{\sim} \mathcal{N}_2(\mu, \Sigma)$ • Estimate $\hat{\mu}$ and $\hat{\Sigma}$ by MLE
 - *Bootstrap sample* $x_i^* \stackrel{\text{ind}}{\sim} \mathcal{N}_2(\hat{\mu}, \hat{\Sigma})$
for $i = 1, 2, \dots, n = 88$
 - *Bootstrap replication* $\hat{\rho}^* = \text{cor coef for } \mathbf{x}^* = (x_1^*, \dots, x_{88}^*)$
- $B = 5000$ replications gave $\widehat{\text{sd}} = 0.074$
- Bootstrap 95% confidence interval $[0.393, 0.682]$ (“BCa”)

Bayes-Frequentist Conversion Factor

- **Bayes** $E \{t(\rho)|\hat{\rho}\} = \frac{\int t(\rho)\pi(\rho)f_{\rho}(\hat{\rho}) d\rho}{\int \pi(\rho)f_{\rho}(\hat{\rho}) d\rho} = \frac{\int t(\rho)\pi(\rho)R(\rho)f_{\hat{\rho}}(\rho) d\rho}{\int \pi(\rho)f_{\hat{\rho}}(\rho) d\rho}$

where $R(\rho) = \frac{f_{\rho}(\hat{\rho})}{f_{\hat{\rho}}(\rho)}$ is “conversion factor.”

- *Simulation* Parametric bootstrap $f_{\hat{\rho}}(\cdot) \rightarrow B$ bootreps
 $\rho_1, \rho_2, \dots, \rho_B \quad [\rho_i = \hat{\rho}_i^*]$

- $\hat{E} \{t(\rho)|\hat{\rho}\} = \frac{\sum_{i=1}^B t_i \pi_i R_i}{\sum \pi_i R_i}$

where $t_i = t(\rho_i)$, $\pi_i = \pi(\rho_i)$, $R_i = R(\rho_i)$.

- Newton–Raftery *JRSS-B* (1994)

Bayes Posterior Density for Flat Prior: $\pi(\rho) = 1$

- Bin the $B = 5000$ bootstrap replications, as for histogram

- $y_k = \text{count in } k\text{th bin} = \sum_1^B I_k(\rho_i)$

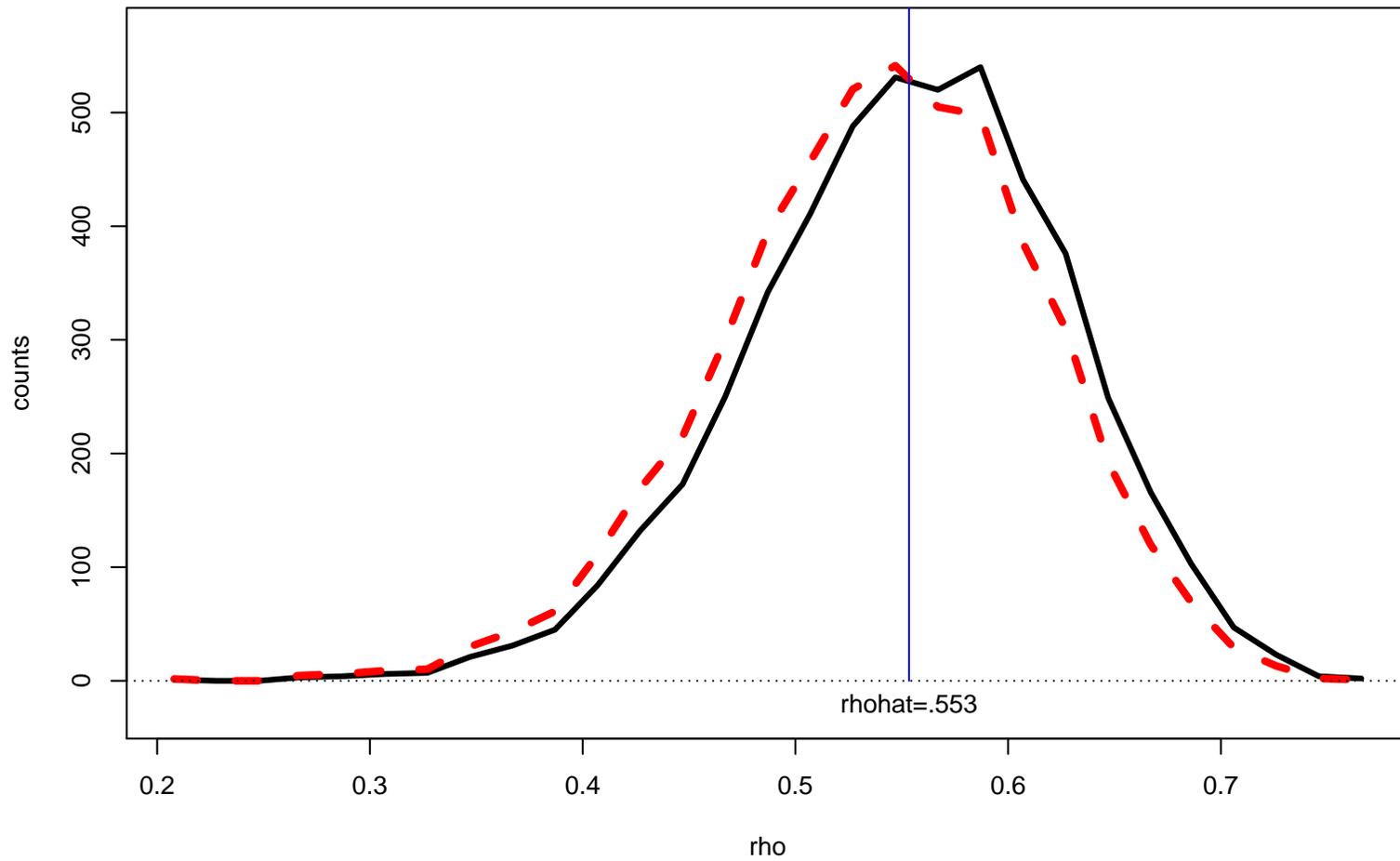
[$I_k(\cdot)$ indicator for ρ in k th bin]

- Bayes counts

$$\tilde{y}_k = \sum_1^B R(\rho_i) I_k(\rho_i) / \bar{R}$$

- Plot y_k and \tilde{y}_k versus bin centers to see Bayes-frequentist differences (i.e., usual bootstrap histogram compared to Bayes posterior dist.)

Counts for 5000 parametric bootstrap replications of rho (black)
compared to Bayes posterior distribution (red) for flat prior



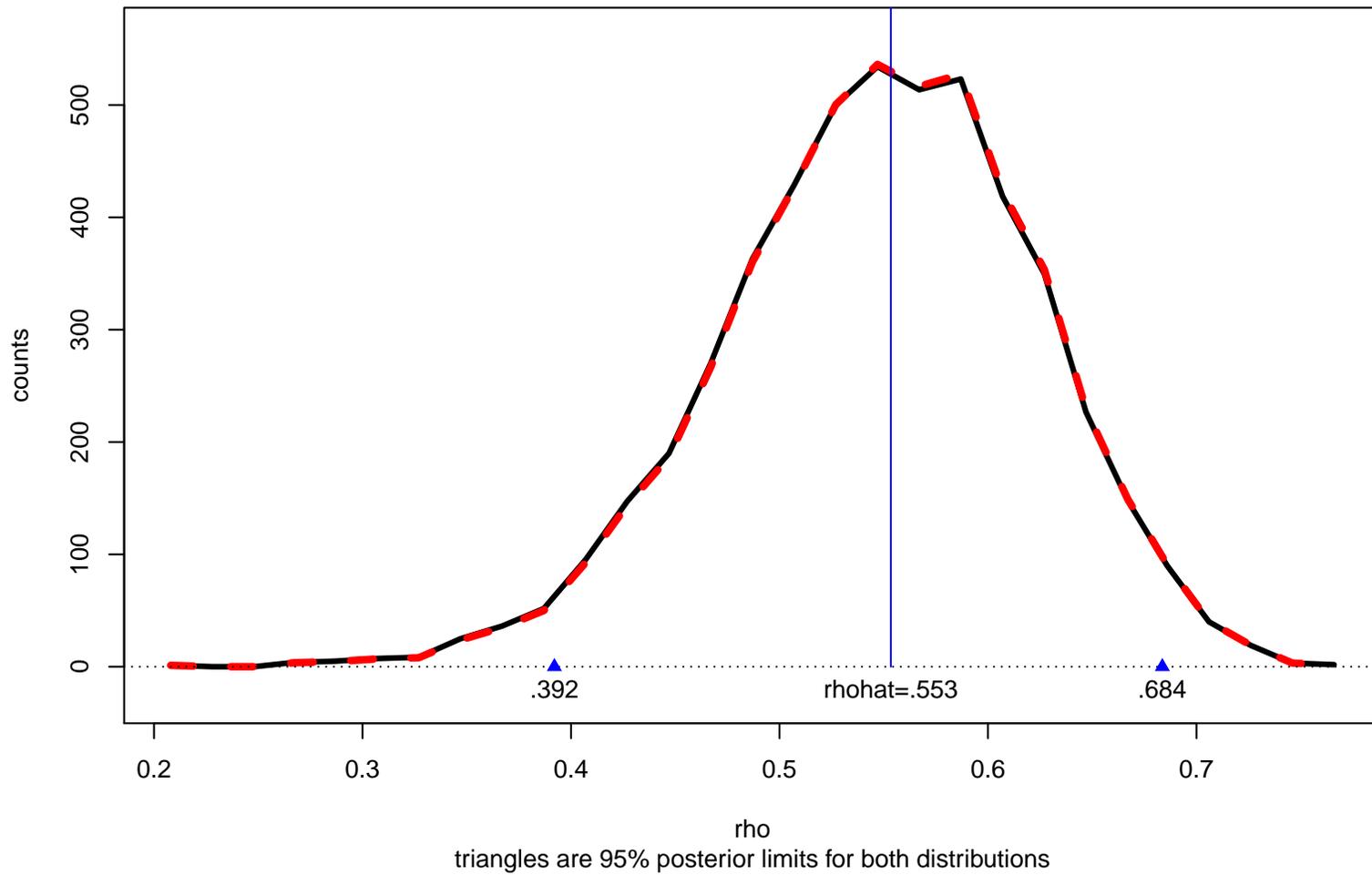
Jeffreys Prior for ρ

- $\pi(\rho) = 1/(1 - \rho^2)$ “uninformative for ρ ” [makes posterior intervals have nearly correct frequentist coverage]
- **BCa method** frequentist weights w_i that reweight raw bootstrap counts to have nearly correct coverages:

$$\hat{E}_{\text{BCa}}\{t\} = \sum t_i w_i / w_i$$

- In this case Bayes \doteq frequentist (not always so!)

Corrected BCa bootstrap distribution (black) and Jeffreys Bayes posterior density (red) for $\rho|\hat{\rho}$



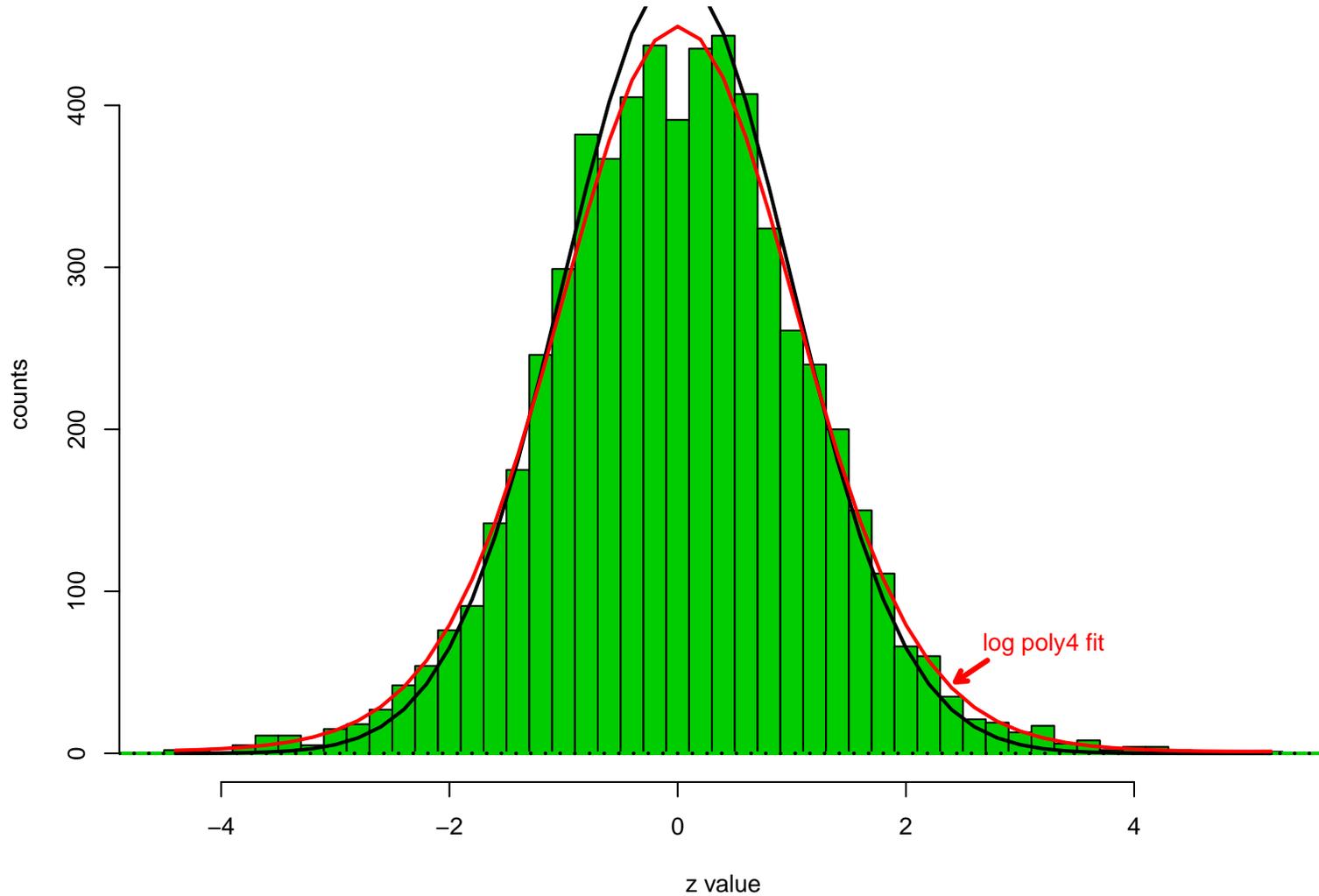
Comparison of Intervals

	.025	.975	
Fisher	.389	.684	(confidence intervals)
Jeffreys	.392	.684	(credible intervals)
BCa	.393	.682	(both?)

Prostate Cancer Study (Singh et al, 2002)

- *Microarray study*
102 men: 52 prostate cancer, 50 healthy controls
- 6033 genes z_i test statistic for H_{0i} : “no difference”
$$H_{0i} : z_i \sim \mathcal{N}(0, 1)$$
- *Goal* Identify genes involved in prostate cancer

Prostate cancer z-values for 6033 genes; 52 patients vs
50 healthy controls. $N(0,1)$ black; LogPoly4 red



False Discovery Rate: $\text{Fdr}(z_0) = \Pr\{H_{0i} | z_i \geq z_0\}$

- Let $S_0(z_0) = \Pr\{\mathcal{N}(0,1) \geq z_0\}$ and $S(z) = \Pr\{z \geq z_0\}$
- $\text{Fdr}(z_0) \doteq S_0(z_0)/S(z_0)$
- Empirical Bayes $\widehat{\text{Fdr}}(z_0) = S_0(z_0)/\hat{S}_{\text{nonpar}}(z_0)$ where
 $\hat{S}_{\text{nonpar}}(z_0) = \#\{z_i \geq z_0\}/n$ (Benjamini–Hochberg 1995)
- Reject H_{0i} if $\widehat{\text{Fdr}}(z_i)$ small (≤ 0.1 or 0.2)

Poisson Model

- Let y_1, y_2, \dots, y_K be heights of histogram bars (i.e., bin counts), $\mu_1, \mu_2, \dots, \mu_K$ their unknown expectations.
- **Poisson model** $y_k \stackrel{\text{ind}}{\sim} \text{Poi}(\mu_k)$
- $\log(\mu)$ polynomial function of z
- **GLM** gives parametric cdf estimate $\hat{S}_{\text{par}}(z_0)$

$$\widehat{\text{Fdr}}_{\text{par}}(z_0) = S_0(z_0) / \hat{S}_{\text{par}}(z_0) = \hat{\theta}$$

- *Example* polynomial degree 4: $\hat{\theta} = \widehat{\text{Fdr}}_{\text{par}}(3.0) = 0.199 \pm ?$

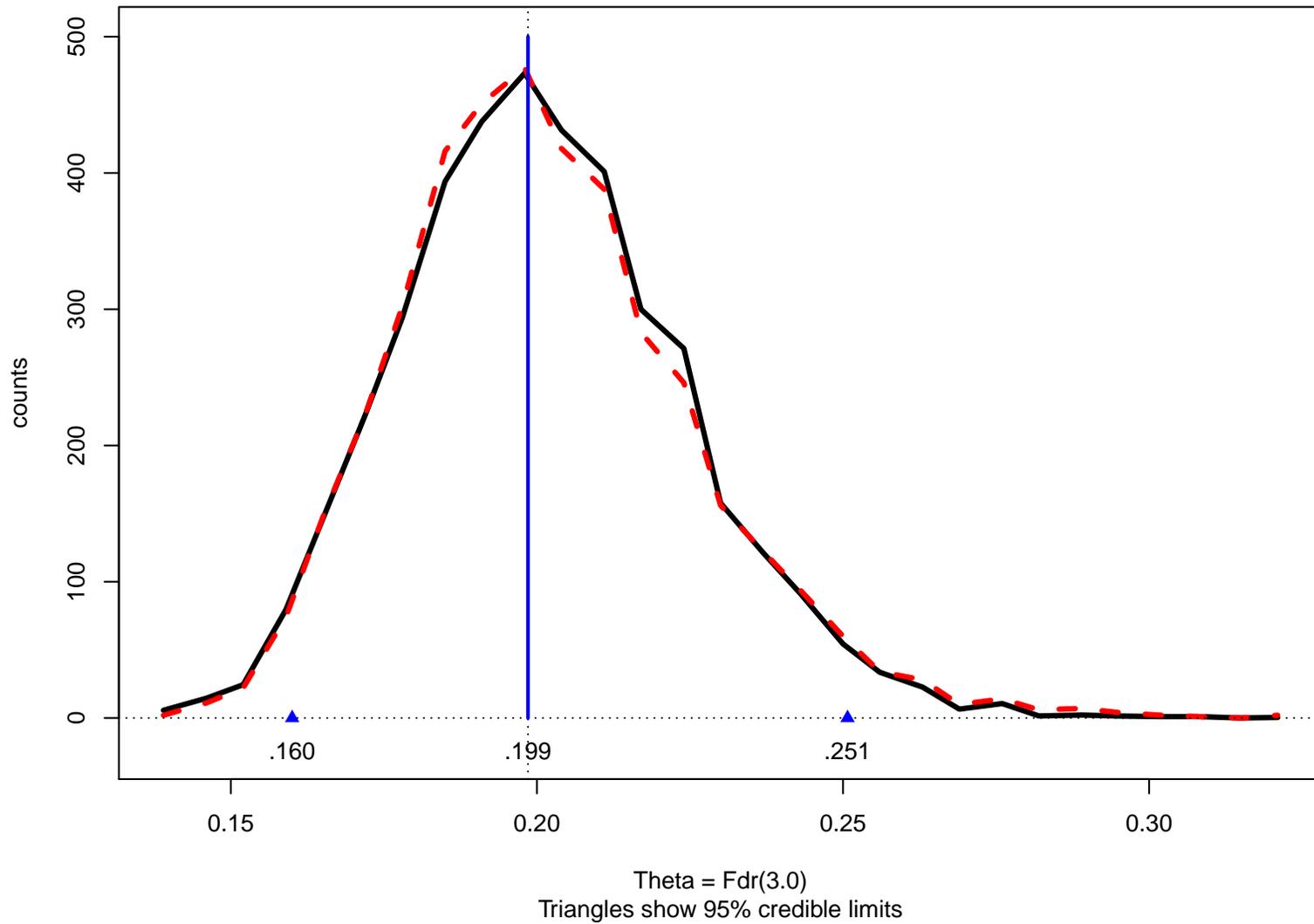
Parametric Bootstrap Replications

- Use estimates $\hat{\mu}_k$ from Poisson GLM
- Bootstrap sample $y_k^* \stackrel{\text{ind}}{\sim} \text{Poi}(\hat{\mu}_k)$ gives $\hat{S}_{\text{par}}^*(z_0)$ and

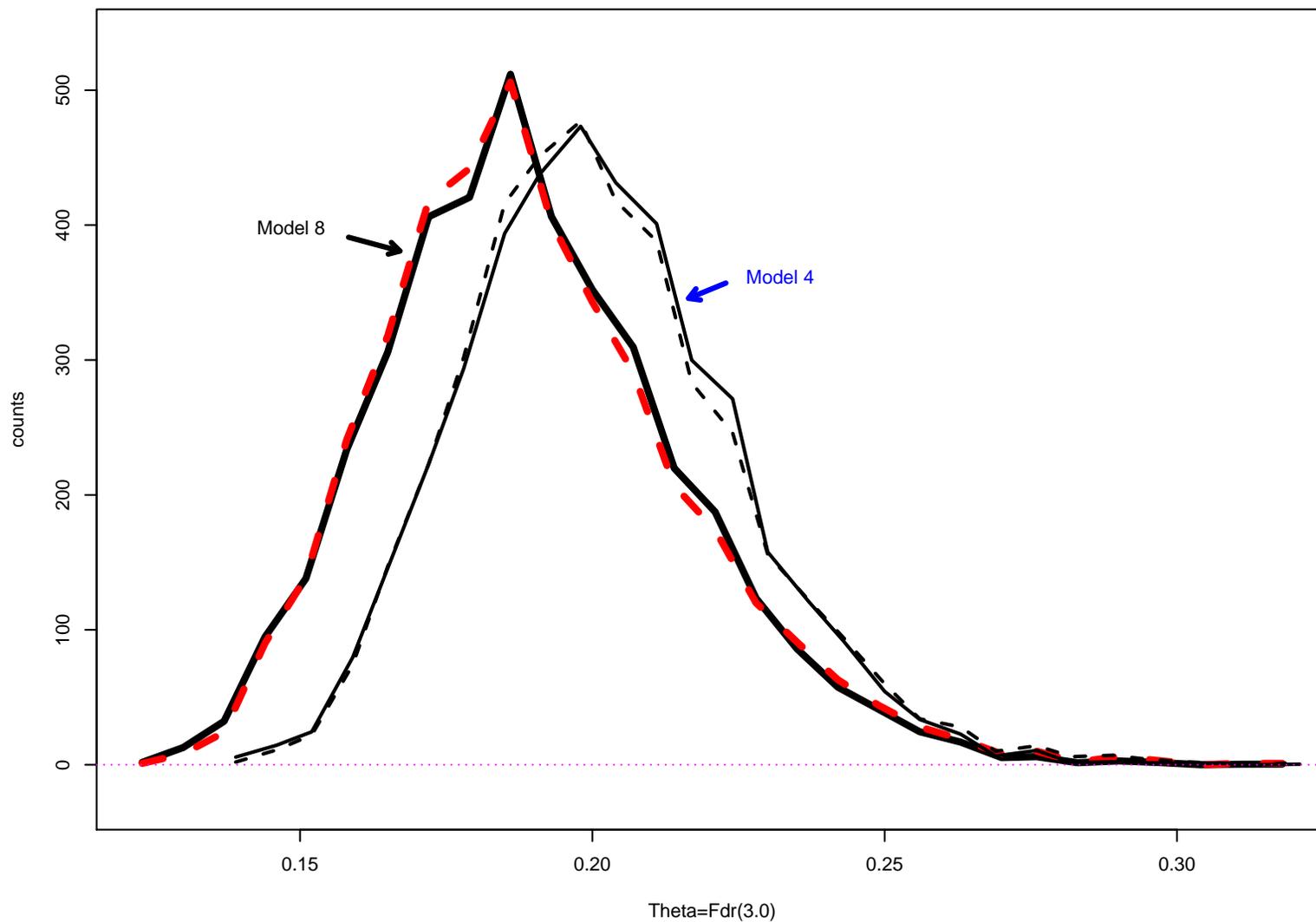
$$\hat{\theta}^* = S_0(z_0) / \hat{S}_{\text{par}}^*(z_0)$$

- $B = 4000$ bootstrap replications gave boot standard error **0.024** for $z_0 = 3$

4000 parametric bootstrap replications for Fdr(3.0), prostate data;
red curve is Bayes posterior counts (flat prior)



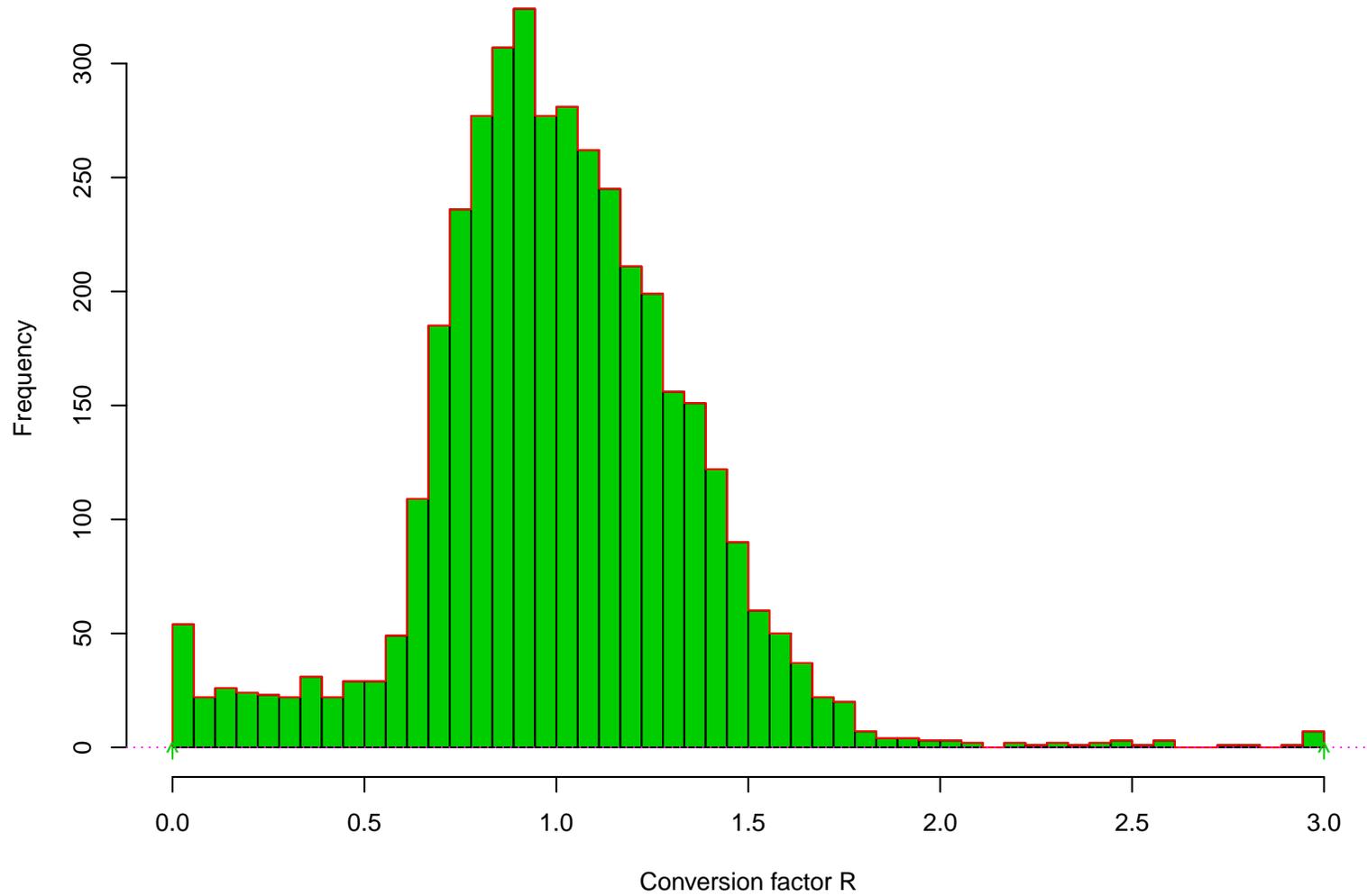
Now Bootstrap/Bayes using 8th degree Poisson model (heavy);
compared with 4th degree (light)



Model Selection: $AIC = Deviance + 2 \cdot df$

Model	deviance	AIC	boot counts	Bayes counts
M2	138.6	142.6	0	0.0
M3	137.1	143.1	0	0.0
M4	65.3	73.3*	1265	1294.8
M5	64.3	74.3	416	421.0
M6	63.8	75.8	215	214.6
M7	63.8	77.8	54	54.7
M8	59.6	75.6	2050	2033.4

Conversion factor R for the prostate data; b=4000 bootreps;
R has mean 1 and standard deviation .365



The Conversion Factor R

- Converts frequentist boot density function $f_{\hat{\beta}}(\hat{\beta}^*)$ to likelihood $f_{\beta}(\hat{\beta})$

- $\hat{t} = \sum_1^B t_i R_i / \sum_1^B R_i$ and $\bar{t} = \sum_1^B t_i / B$, $\overline{\text{sd}}$ = boot st dev of t :

$$\frac{\hat{t} - \bar{t}}{\overline{\text{sd}}} = \overline{\text{cor}}(t, R) \cdot \overline{\text{CV}}(R)$$

- *Prostate example*

$$\overline{\text{CV}} = 0.365 \quad \overline{\text{cor}} = -0.01 \quad \text{so } \frac{\hat{t} - \bar{t}}{\overline{\text{sd}}} = -0.004$$

- *Correlation example*

$$\overline{\text{CV}} = 0.172 \quad \overline{\text{cor}} = -0.999 \quad \text{so " " } = -0.17$$

One-Parameter Exponential Families

- θ the expectation parameter
(e.g., p in binomial, λ in Poisson)

- Repeated sampling

$$\hat{\theta} \sim (\theta, \sigma_{\theta} / \sqrt{n}, \gamma_{\theta} / \sqrt{n}) \quad (\text{mean, sd, skewness})$$

- $Z \equiv \sqrt{n} \frac{\theta - \hat{\theta}}{\sigma_{\hat{\theta}}} \longrightarrow \mathcal{N}(0, 1)$

$$\log R(\theta) \doteq \frac{\gamma_{\hat{\theta}}}{2\sqrt{n}} Z \cdot (1 + Z^2/3)$$

- Rate Bayes \longrightarrow frequentist depends on skewness $\gamma_{\hat{\theta}}$

Some Conclusions

- Parametric bootstrap connects Bayes and frequentist analyses, especially Jeffreys-type uninformative Bayes.
[bootstrap: only one probability distribution]
- Convenient way to compute $\pi(\theta|\mathbf{x})$.
[competition to MCMC]
- Provides easy estimates of simulation error.
- “Behavior” important for Jeffreys-Bayes methods.