

Bayesian Inference and the Parametric Bootstrap

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Importance Sampling for Bayes Posterior Distribution

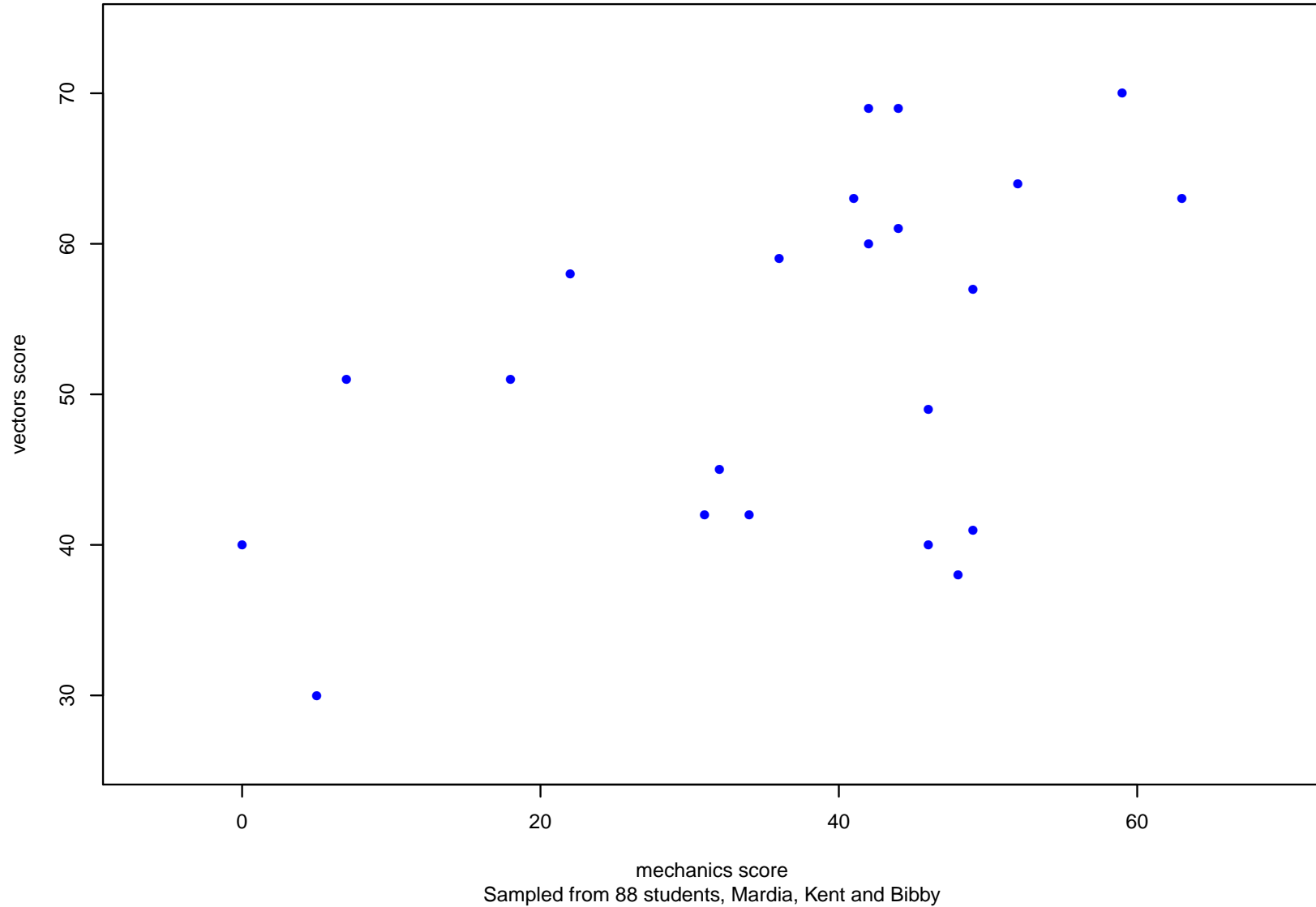
- Newton and Raftery (1994 *JRSS-B*)
“Nonparametric Bootstrap: good choice”
(actually used a smoothed nonparametric bootstrap)
- Today Parametric Bootstrap
 - Good computational properties when applicable
 - Connection between Bayes and frequentist inference

Student Score Data

(Mardia, Kent and Bibby)

- $n = 22$ students' scores on two tests: *mechanics, vectors*
- **Data** $\mathbf{y} = (y_1, y_2, \dots, y_{22})$ with $y_i = (\text{mec}_i, \text{vec}_i)$
- *Parameter of interest* $\theta = \text{correlation}(\text{mec}, \text{vec})$
- *Sample correlation coefficient* $\hat{\theta} = 0.498 \pm ??$

Scores of 22 students on two tests 'mechanics' and 'vectors';
Sample Correlation Coefficient is .498 +-??



R. A. Fisher (1915)

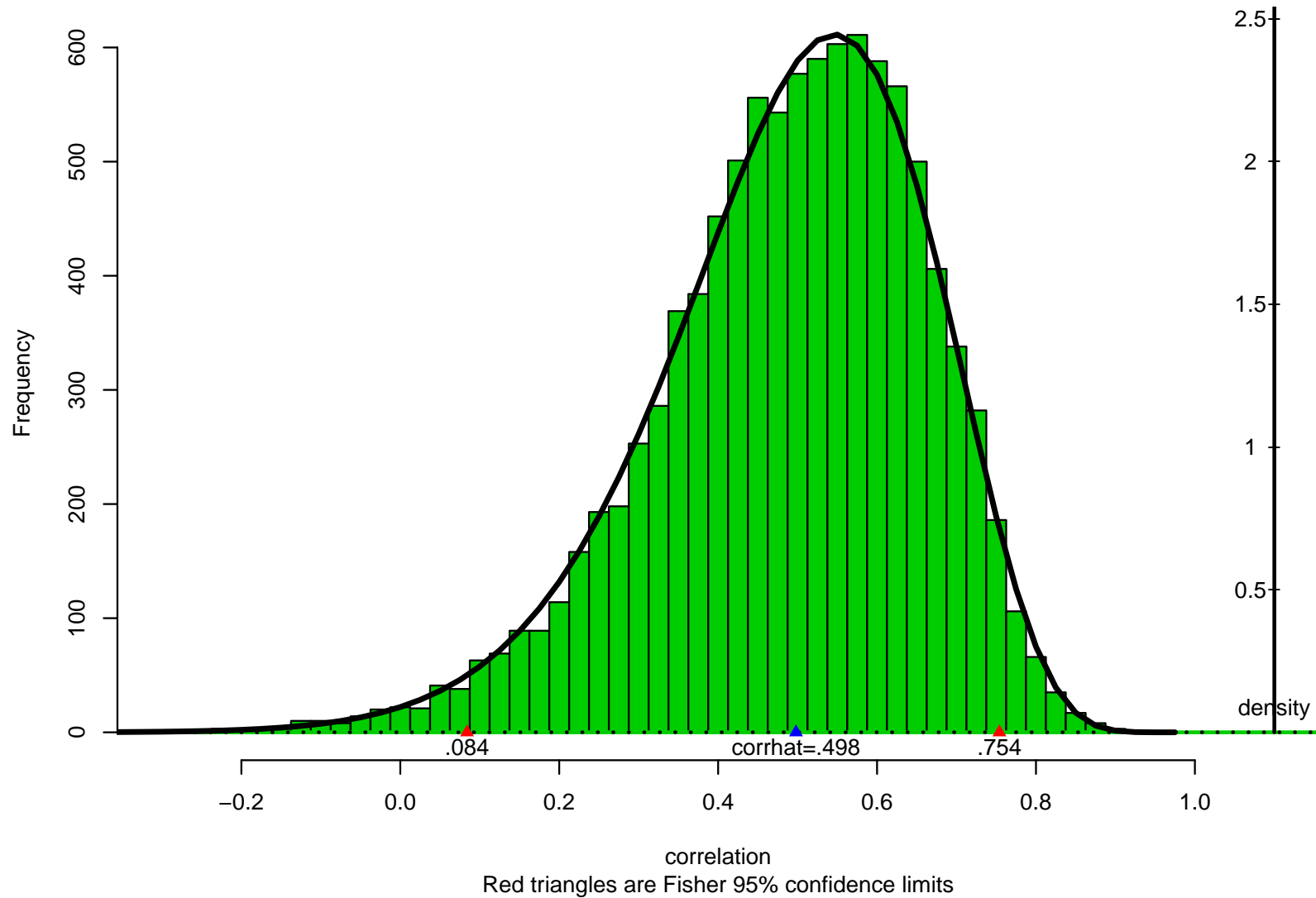
- Density $f_{\theta}(\hat{\theta})$

$$= \frac{n-2}{\pi} (1-\theta^2)^{(n-1)/2} (1-\hat{\theta}^2)^{(n-4)/2} \int_0^{\infty} [\cosh(w) - \theta\hat{\theta}]^{-(n-1)} dw$$

- Bivariate normal $y_i \stackrel{\text{ind}}{\sim} \mathcal{N}_2(\mu, \Sigma)$ $i = 1, 2, \dots, n$
- 95% confidence limits (from z transform)

$$\theta \in (0.084, 0.754)$$

Fisher density for correlation coefficient if $\text{corr}=.498$
And histogram for 10000 parametric bootstrap replications



Parametric Bootstrap Computations

- Assume $y_i \stackrel{\text{ind}}{\sim} \mathcal{N}_2(\mu, \Sigma)$
- Estimate $\hat{\mu}$ and $\hat{\Sigma}$ by MLE
- Bootstrap sample $y_i^* \stackrel{\text{ind}}{\sim} \mathcal{N}_2(\hat{\mu}, \hat{\Sigma})$ for $i = 1, 2, \dots, 22$
- Bootstrap replication $\hat{\theta}^* = \text{corr coeff for } \mathbf{y}^* = (y_1^*, y_2^*, \dots, y_{22}^*)$
- Bootstrap standard deviation

$$\text{sd}(\hat{\theta}) = 0.168 \quad (B = 10,000)$$

- Percentile 95% confidence limits (0.109, 0.761)

Better Bootstrap Confidence Limits (BCa)

- **Idea** Reweighting the B bootstrap replications improves coverage
- Weights involve z_0 (bias correction) and a (acceleration)
- $W_{\text{BCa}}(\theta) = \frac{\varphi(Z_\theta / (1 + aZ_\theta) - z_0)}{(1 + aZ_\theta)^2 \varphi(Z_\theta + z_0)}$ where $Z_\theta = \Phi^{-1} G(\theta) - z_0$
 \uparrow
bootstrap cdf
- Reweighting the $B = 10,000$ bootstraps gave weighted percentiles

(0.075, 0.748)

Fisher: (0.084, 0.754)

Bayes Posterior Intervals

- *Prior* $\pi(\theta)$
- Posterior expectation for $t(\theta)$:

$$E\{t(\theta)|\hat{\theta}\} = \int t(\theta)\pi(\theta)f_{\theta}(\hat{\theta})d\theta \Big/ \int \pi(\theta)f_{\theta}(\hat{\theta})d\theta$$

- **Conversion factor** Ratio of likelihood to bootstrap density:

$$R(\theta) = f_{\theta}(\hat{\theta})/f_{\hat{\theta}}(\theta) \quad (\text{"}\theta\text{"} = \hat{\theta}^*)$$

- **Bootstrap integrals**

$$E\{t(\theta)|\hat{\theta}\} = \frac{\int t(\theta)\pi(\theta)R(\theta)f_{\hat{\theta}}(\theta)d\theta}{\int \pi(\theta)R(\theta)f_{\hat{\theta}}(\theta)d\theta}$$

Bootstrap Estimation of Bayes Expectation $E\{t(\theta)|\hat{\theta}\}$

- Parametric bootstrap replications $f_{\hat{\theta}}(\cdot) \rightarrow \theta_1, \theta_2, \dots, \theta_i, \dots, \theta_B$
- $t_i = t(\theta_i), \quad \pi_i = \pi(\theta_i), \quad R_i = R(\theta_i):$

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

- *Reweighting* Weight $\pi_i R_i$ on θ_i
- Importance sampling estimate:

$$\hat{E}\{t(\theta)|\hat{\theta}\} \rightarrow E\{t(\theta)|\hat{\theta}\} \quad \text{as } B \rightarrow \infty$$

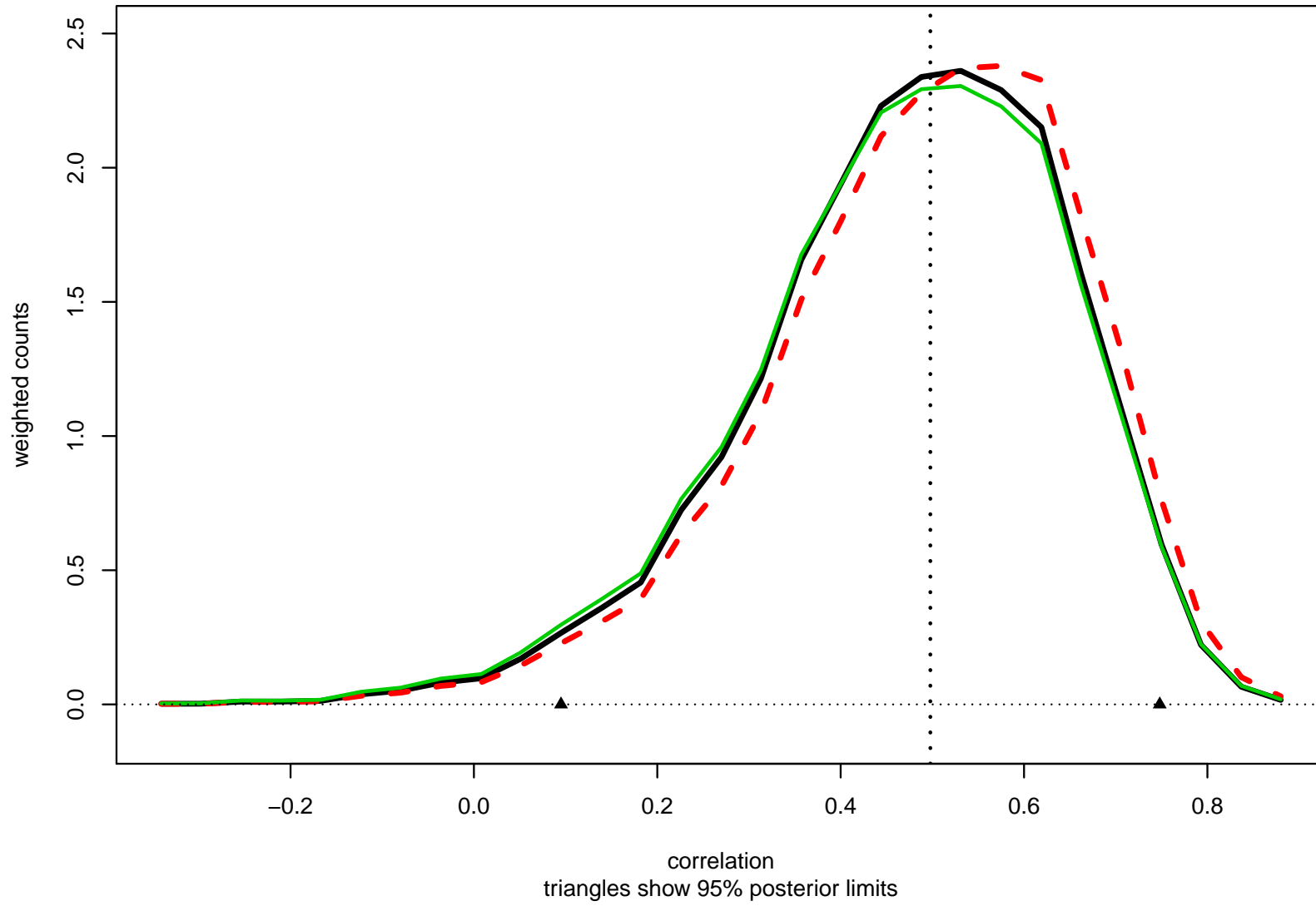
Jeffreys Priors

- **Harold Jeffreys** (1930s):
Theory of “uninformative” prior distributions
(“invariant”, “objective”, “reference”, ...)
- For density family $f_{\theta}(y)$, $\theta \in \mathcal{R}^p$:

$$\pi(\theta) = c|I(\theta)|^{1/2} \quad I(\theta) = \text{Fisher info matrix}$$

- *Normal correlation* $\pi(\theta) = 1/(1 - \theta^2)$

Importance sampling posterior density for correlation (black)
using Jeffreys prior $\pi(\theta)=1/(1-\theta^2)$;
BCa weighted density (green); raw bootstrap density (red)



Multiparameter Exponential Families \mathcal{R}^P

- p -dimensional sufficient statistic $\hat{\beta}$
- p -dimensional parameter vector $\beta = E\{\hat{\beta}\}$
(Poisson: $e^{-\lambda}\lambda^x/x!$ has $\hat{\beta} = x$, $\beta = \lambda$)
- Hoeffding $f_{\beta}(\hat{\beta}) = f_{\hat{\beta}}(\hat{\beta}) e^{-D(\hat{\beta},\beta)/2}$
- Deviance $D(\beta_1, \beta_2) = 2E_{\beta_1} \log \left\{ f_{\beta_1}(\hat{\beta}) / f_{\beta_2}(\hat{\beta}) \right\}$

Conversion Factor in Exponential Families

- Conversion factor $R(\beta) = f_{\beta}(\hat{\beta}) / f_{\hat{\beta}}(\beta)$ (with $\hat{\beta}$ fixed)

$$R(\beta) = v(\beta)e^{\Delta(\beta)}$$

where $\Delta(\beta) = [D(\beta, \hat{\beta}) - D(\hat{\beta}, \beta)] / 2$

- $v(\beta) = f_{\hat{\beta}}(\hat{\beta}) / f_{\beta}(\beta) \doteq 1/\text{Jeffreys prior}$ (Laplace)
- Jeffreys prior $\pi(\beta)R(\beta) \doteq e^{\Delta(\beta)}$

Example: Multivariate Normal

- $y_i \stackrel{\text{ind}}{\sim} \mathcal{N}_d(\mu, \Sigma)$ for $i = 1, 2, \dots, n$

- $\Delta =$

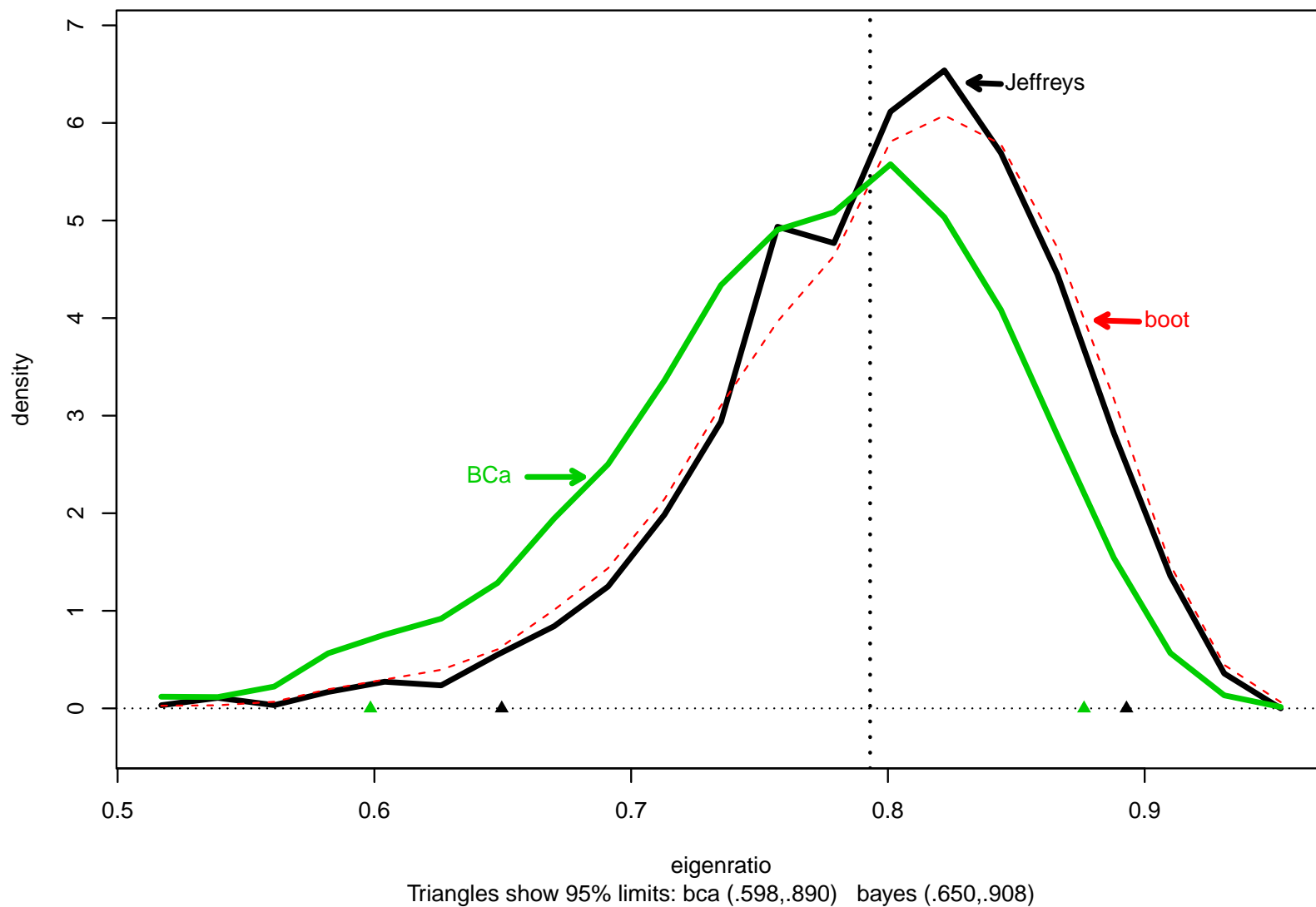
$$n \left\{ (\mu - \hat{\mu})' \frac{\hat{\Sigma}^{-1} - \Sigma^{-1}}{2} (\mu - \hat{\mu}) + \frac{1}{2} \text{tr} \left(\Sigma \hat{\Sigma}^{-1} - \hat{\Sigma} \Sigma^{-1} \right) + \log \frac{|\hat{\Sigma}|}{|\Sigma|} \right\}$$

- “eigenratio” (student score data)

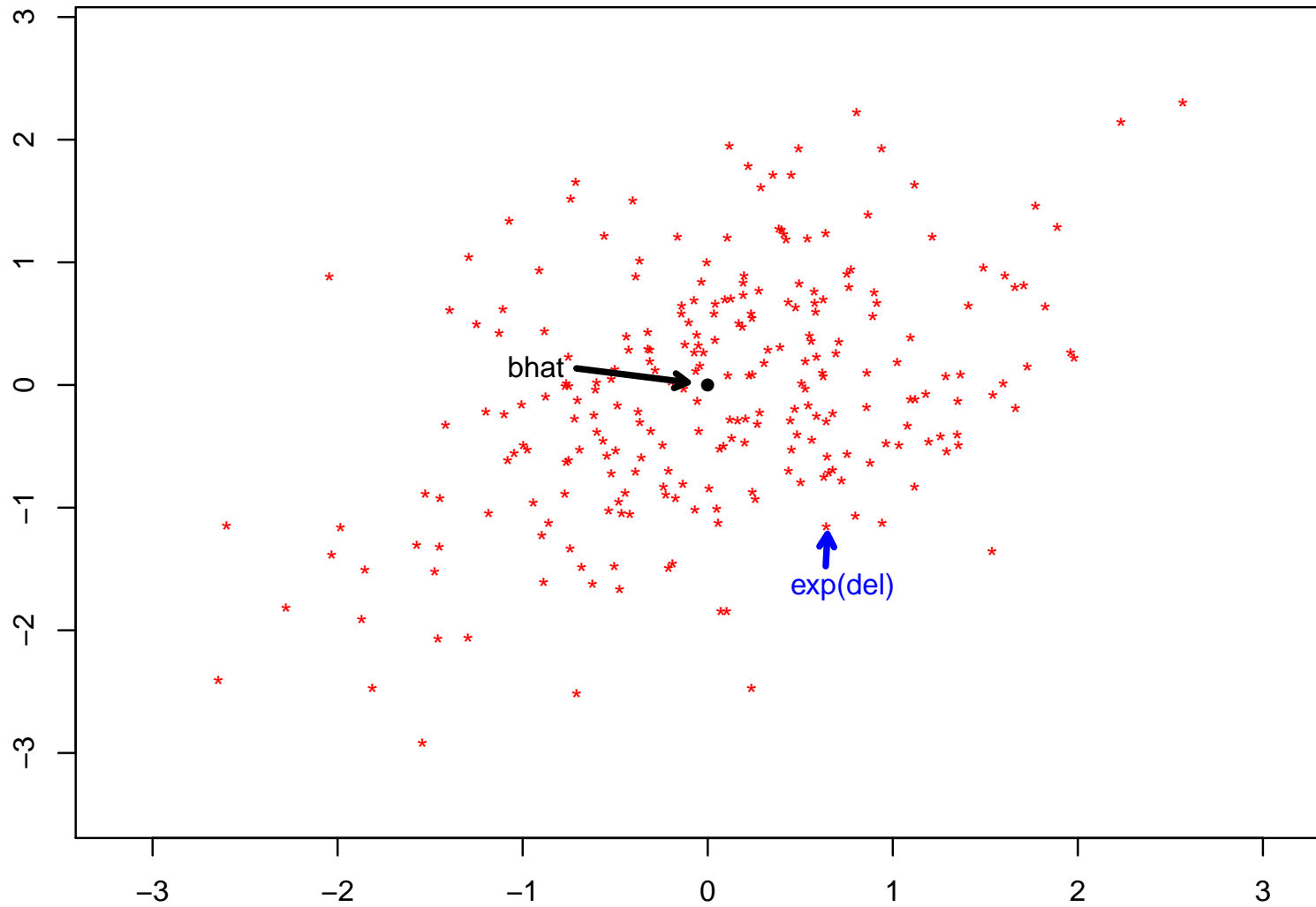
$$\bullet \theta = \frac{\lambda_1}{\lambda_1 + \lambda_2} \quad \bullet \hat{\theta} = \frac{\hat{\lambda}_1}{\hat{\lambda}_1 + \hat{\lambda}_2} = 0.793 \pm ??$$

- **Bootstrap** $y_i^* \stackrel{\text{ind}}{\sim} \mathcal{N}_d(\hat{\mu}, \hat{\Sigma}) \quad i = 1, 2, \dots, n \rightarrow \theta_1, \theta_2, \dots, \theta_B$

Density estimates for student score eigenratio (B=10,000);
Boot (red),Jeffreys (5-dim prior) Black, and BCa Green (z0=-.182, a=0)



Reweighting the parametric bootstrap points



Prostate Cancer Study

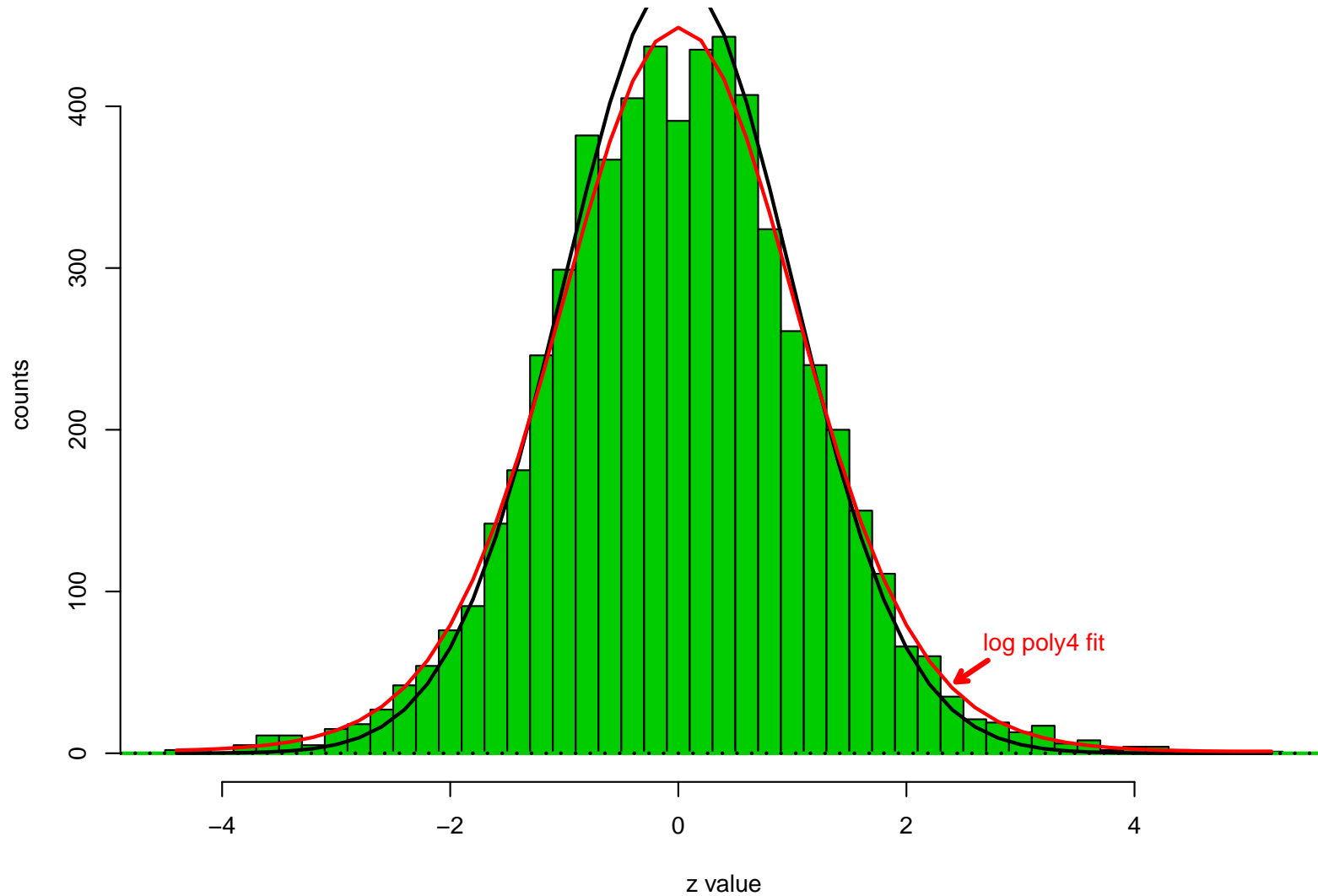
(Singh et al, 2002)

- *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



Poisson Regression Models

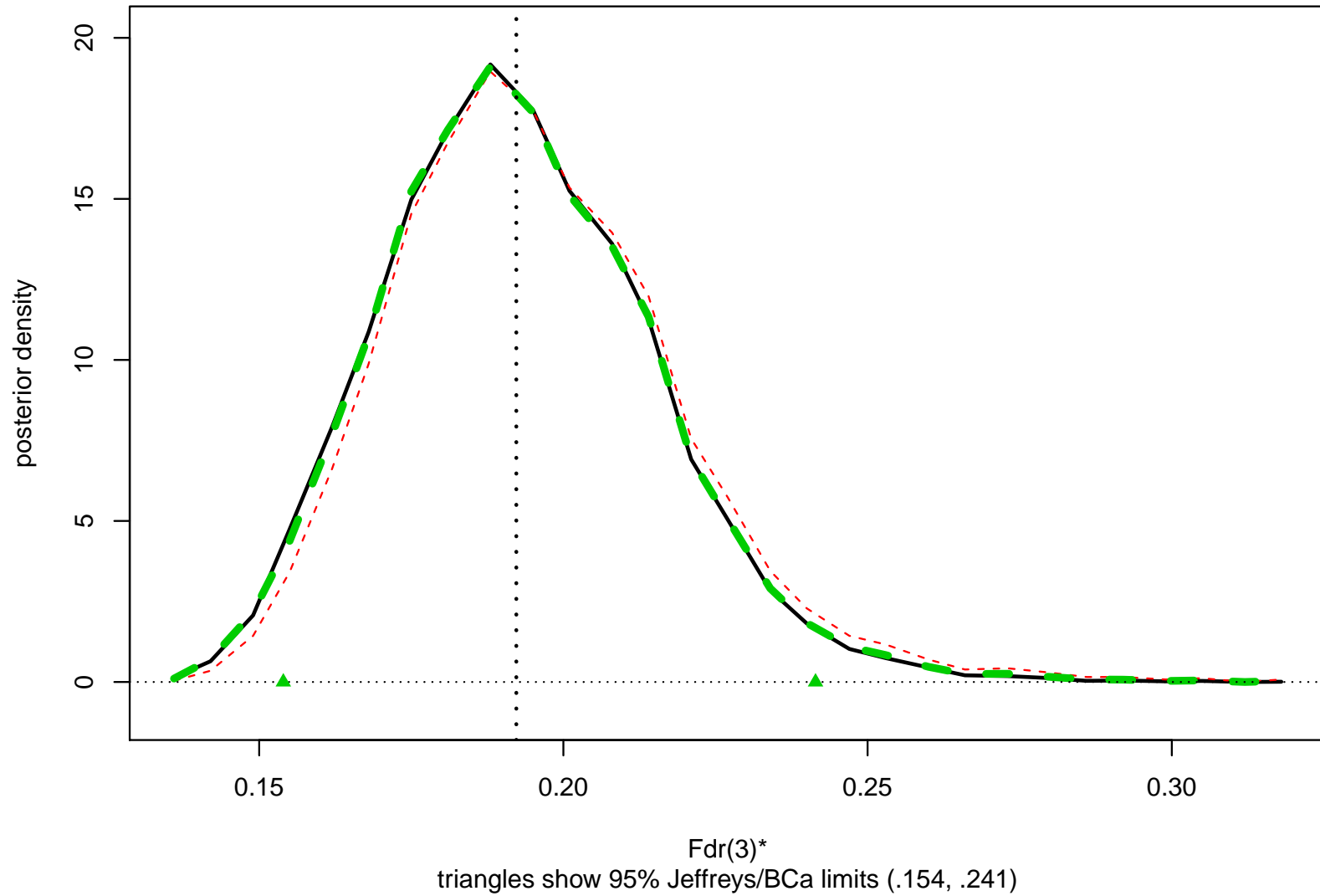
- *Histogram* $y_j = \#\{z_i \in \text{bin}_j\}$
- $x_j = \text{midpoint bin}_j, j = 1, 2, \dots, J$
- *Poisson model* $y_j \stackrel{\text{ind}}{\sim} \text{Poi}(\mu_j)$
with $\log(\mu_j) = \text{poly}(x_j, \text{degree} = m)$
- *Exponential family* degree $p = m + 1$ • Let $\eta_j = \log(\mu_j)$

$$\Delta = (\boldsymbol{\eta} - \hat{\boldsymbol{\eta}})' (\boldsymbol{\mu} + \hat{\boldsymbol{\mu}}) - 2 \left(\sum \mu_j - \sum \hat{\mu}_j \right)$$

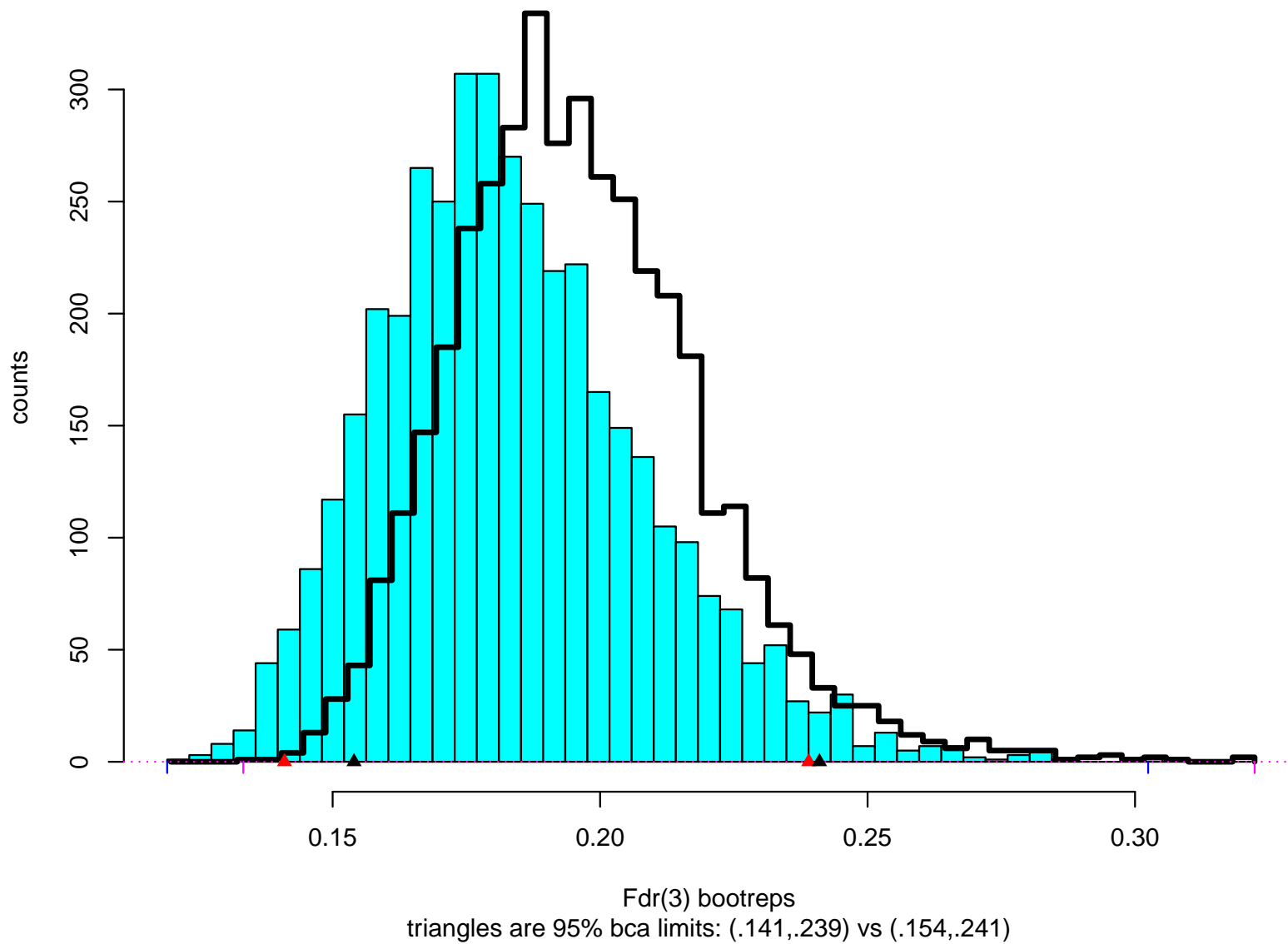
Parametric False Discovery Rate Estimates

- $\text{glm}(y \sim \text{poly}(x, 4), \text{Poisson}) \longrightarrow \{\hat{\mu}_j\}$ (“log poly 4”)
- $\theta = \widehat{\text{Fdr}}(3) = [1 - \Phi(3)] / [1 - \hat{F}(3)] \approx \Pr\{\text{null} | z \geq 3\}$
(\hat{F} is smoothed CDF estimate)
- Model 4: $\widehat{\text{Fdr}}(3) = 0.192 \pm ??$
- Parametric bootstrap $y_j^* \stackrel{\text{ind}}{\sim} \text{Poi}(\hat{\mu}_j) \longrightarrow \{\hat{\mu}_j^*\} \longrightarrow \widehat{\text{Fdr}}(3)^*$

Fdr(3) estimation, Model 4; prostate data from 4000 parboots;
Jeffreys Bayes (black), raw boot (red), bca (green)



Compare Model 4 (line) with Model 8 (solid);
4000 parametric bootstrap replications of Fdr(3)



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

Model	AIC	boot counts	Bayes counts [†]	
M2	142.6	0	0.0	
M3	143.1	0	0.0	
M4	73.3	1266	1458	(36%)
M5	74.3	415	462	(12%)
M6	75.8	215	197	(5%)
M7	77.8	54	67	(2%)
M8	75.6	2050	1816	(45%)

[†] Model 8, Jeffreys prior

Internal Accuracy of Bayes Estimates

- B for computing $\hat{E} \{t(\beta)|\hat{\beta}\} = \sum_1^B t_i \pi_i R_i / \sum_1^B \pi_i R_i$?

- Define $r_i = \pi_i R_i$ and $s_i = t_i \pi_i R_i$
- $\hat{c}_{ss} = \sum_1^B (s_i - \bar{s})^2 / B$, etc.

$$\text{CV} \{ \hat{E} \}^2 = \frac{1}{B} \left(\frac{\hat{c}_{ss}}{\bar{s}^2} - 2 \frac{\hat{c}_{sr}}{\bar{s}\bar{r}} + \frac{\hat{c}_{rr}}{\bar{r}^2} \right)$$

- *Example* $t = \text{Fdr}(3)$

4000 parametric bootreps, Jeffreys Bayes

- Model 4 $\hat{E} = 0.193$ $\widehat{\text{CV}} = \mathbf{0.0019}$

- Model 8 $\hat{E} = 0.179$ $\widehat{\text{CV}} = \mathbf{0.0025}$

Sampling Variability of Bayes Estimates

- *How much* would $\hat{E} \{t(\beta)|\hat{\beta}\}$ vary for new $\hat{\beta}$'s?
(i.e., frequentist properties of Bayes estimates)
- Need to bootstrap the parametric bootstrap calculations for $\hat{E} \{t(\beta)|\hat{\beta}\}$
- **Shortcut** “Bootstrap after bootstrap”

Bootstrap-after-Bootstrap Calculations

- $f_{\hat{\beta}}(\cdot) \longrightarrow \beta_1, \beta_2, \dots, \beta_i, \dots, \beta_B$,
the original bootstrap reps under $\hat{\beta}$
- Want bootreps under some other value $\hat{\gamma}$
- **Idea** Reweight the originals
- **Weight** $Q_{\hat{\gamma}}(\beta) = f_{\beta}(\hat{\gamma}) / f_{\beta}(\hat{\beta})$:

$$\hat{E}\{t|\hat{\gamma}\} = \frac{\sum_1^B t_i \pi_i R_i Q_{\hat{\gamma}}(\beta_i)}{\sum_1^B \pi_i R_i Q_{\hat{\gamma}}(\beta_i)}$$

Bootstrap-after-Bootstrap Standard Errors

- BAB

(1) $f_{\hat{\beta}}(\cdot) \rightarrow \gamma_1, \gamma_2, \dots, \gamma_k$

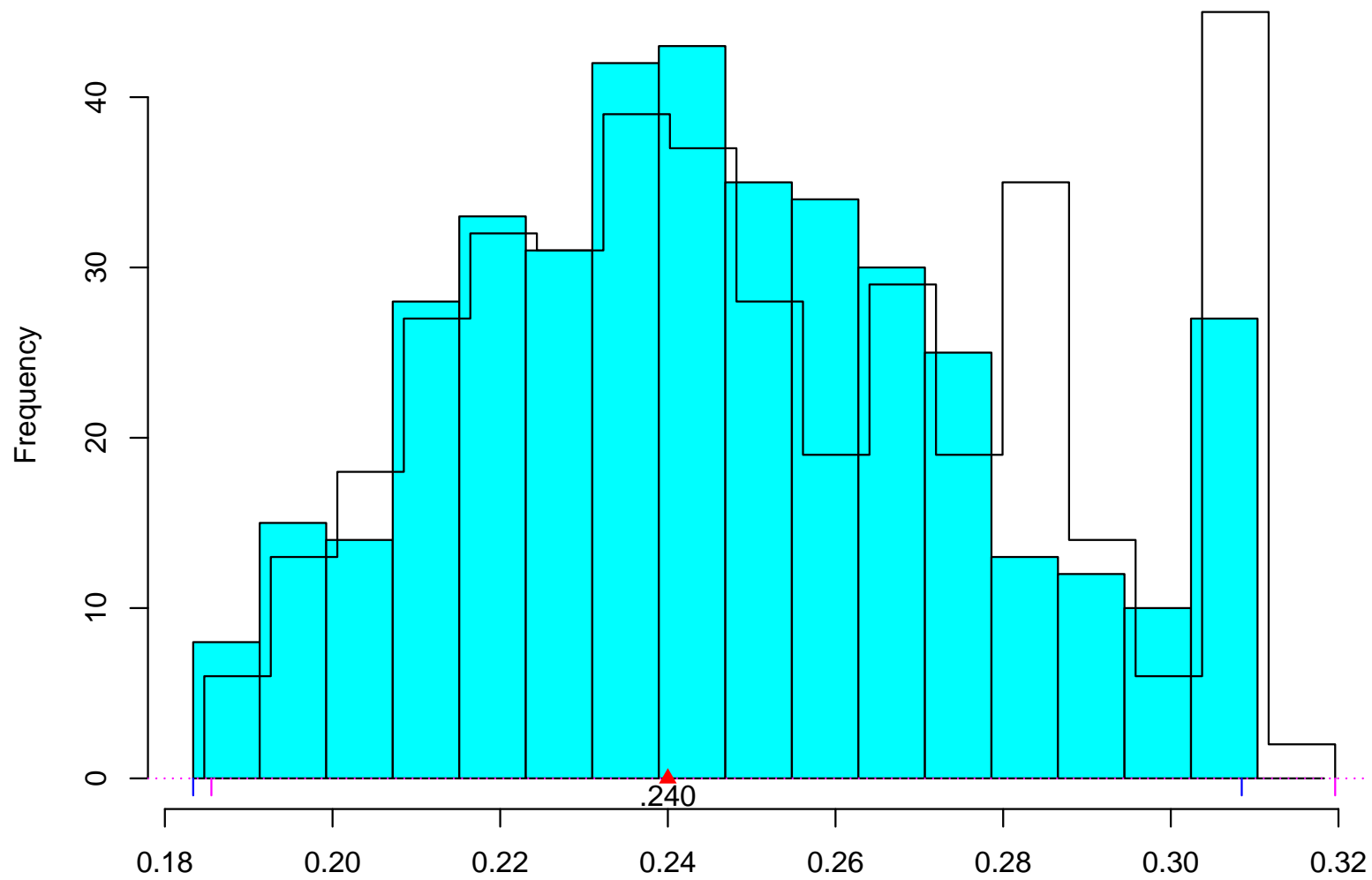
(2) $\hat{E}_k = \hat{E} \{t | \hat{\gamma}_k\}$

(3) $\text{se} \{ \hat{E} \} = \left[\sum (\hat{E}_k - \hat{E}.)^2 / K \right]^{1/2}$

- *Example* Model 4

97.5% credible limit for $\text{Fdr}(3) = 0.241 \pm ??$

400 BAB replications of 97.5% credible upper limit for Fdr(3);
line histogram: same for BCa upper limit



97.5% credible upper limit
BAB Standard Errors: .030 (Bayes) .034 (BCa); corr .98

BAB SDs for Bayes Model Selection Proportions

Model	Bayes post. prob.	BAB SD
M4	36%	$\pm 20\%$
M5	12%	$\pm 14\%$
M6	5%	$\pm 8\%$
M7	2%	$\pm 6\%$
M8	45%	$\pm 27\%$

- *Model averaging?*

A Few Conclusions ...

- Parametric bootstrap closely related to objective Bayes.
(That's why it's a good importance sampling choice.)
- When it applies, parboot approach has both computational and interpretational advantages over MCMC/Gibbs.
- Objective Bayes analyses should be checked frequentistically.

References

- **Bootstrap** DiCiccio & Efron 1996 *Statist. Sci.* 189–228
- **Bayes and Bootstrap** Newton & Raftery 1994 *JRSS-B* 3–48 (see Discussion!)
- **Objective Priors** Berger & Bernardo 1989 *JASA* 200–207; Ghosh 2011 *Statist. Sci.* 187–202
- **Exponential Families** Efron 2010 *Large-Scale Inference*, Appendix 1
- **MCMC and Bootstrap** Efron 2011 *J. Biopharm. Statist.* 1052–1062