Bayesian Variable Selection & Bayesian Model Averaging

Hoff Chapter 9, Liang et al 2008, Hoeting et al (1999), Clyde & George (2004)

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Prior & Posterior Recap

$$\begin{split} \mathbf{Y} \mid \alpha, \beta_{\gamma}, \phi, \gamma &\sim \mathsf{N}(1\alpha + \mathsf{X}_{\gamma}\beta_{\gamma}, \mathsf{I}_{n}/\phi) \\ p(\alpha, \phi) &\propto 1/\phi \quad \beta_{\gamma} \mid \gamma, \phi \sim \mathsf{N}(0, \frac{g}{\phi}(\mathsf{X}_{\gamma}^{T}\mathsf{X}_{\gamma})^{-1}) \quad \gamma \sim p(\gamma) \\ \alpha \mid \phi, y \sim \mathsf{N}\left(\bar{y}, \frac{1}{n\phi}\right) \\ \beta_{\gamma} \mid \gamma, \phi, g, y \sim \mathsf{N}\left(\frac{g}{1+g}\hat{\beta}_{\gamma}, \frac{g}{1+g}\frac{1}{\phi}\left[\mathsf{X}_{\gamma}^{T}\mathsf{X}_{\gamma}\right]^{-1}\right) \\ \phi \mid \gamma, y \sim \mathsf{Gamma}\left(\frac{n-1}{2}, \frac{\mathsf{TotalSS} - \frac{g}{1+g}\mathsf{RegSS}}{2}\right) \\ p(\gamma \mid \mathsf{Y}) &= \frac{BF(\gamma : \gamma 0)p(\gamma)/p(\gamma 0)}{\sum_{\gamma' \in \Gamma} BF(\gamma' : \gamma 0)p(\gamma')/p(\gamma 0)} \\ BF(\gamma : \gamma 0) &= (1+g)^{(n-1-p\gamma)/2}(1+g(1-R_{\gamma}^{2}))^{-(n-1)/2} \end{split}$$

Choice of g: Bartlett's Paradox

The Bayes factor for comparing γ to the null model:

$$BF(\gamma:\gamma 0) = (1+g)^{(n-1-p_{\gamma})/2} (1+g(1-R_{\gamma}^2))^{-(n-1)/2}$$

- For fixed sample size n and R^2_{γ} , consider taking values of g that go to infinity
- Increasing vagueness in prior
- ▶ What happens to BF as $g \to \infty$?
- Why is this a paradox?

Information Paradox

The Bayes factor for comparing γ to the null model:

$$BF(\gamma:\gamma_0) = (1+g)^{(n-1-p_{\gamma})/2} (1+g(1-R_{\gamma}^2))^{-(n-1)/2}$$

- Let g be a fixed constant and take n fixed.
- $\blacktriangleright \text{ Let } F = \frac{R_{\gamma}^2/p_{\gamma}}{(1-R_{\gamma}^2)/(n-1-p_{\gamma})}$
- As $R^2_{\gamma} \to 1$, $F \to \infty$ LR test would reject γ_0 where F is the usual F statistic for comparing model γ to γ_0
- ▶ BF converges to a fixed constant $(1+g)^{n-1-p_{\gamma}/2}$ (does not go to infinity

"Information Inconsistency" see Liang et al JASA 2008

Mixtures of g priors & Information consistency

Need
$$BF o \infty$$
 if $R_{\gamma}^2 o 1$

Put a prior on
$$g$$

$$C \int (1+\pi)^{(n-1-p_{\alpha})/2} (1+\pi)^{(n-1-p_{\alpha})/2} dx$$

to diverge

paradox!

$$BF(\gamma:\gamma_0)=rac{C\int(1+g)}{C}$$

 $BF(\gamma:\gamma_0) = \frac{C\int (1+g)^{(n-1-p_{\gamma})/2} (1+g(1-R_{\gamma}^2))^{-(n-1)/2} \pi(g) dg}{C}$

 $\mathsf{E}_{\sigma}[(1+g)^{(n-1-p_{\gamma})/2}]$

 $p(g) = \frac{a-2}{2}(1+g)^{-a/2}$

ightharpoonup Consider minimal model $p_{\gamma}=1$ and n=3 (can estimate intercept, one coefficient, and σ^2 , then a > 3 integral exists \triangleright For 2 < a < 3 integral diverges and resolves the information

lacktriangle interchange limit and integration as $R^2 o 1$ want

▶ Need $BF \to \infty$ if $R^2_{\sim} \to 1$

hyper-g prior (Liang et al JASA 2008)

or $g/(1+g) \sim Beta(1, (a-2)/2)$

• prior expectation converges if $a > n + 1 - p_{\gamma}$

Mixtures of g priors & Information consistency

Need $BF \to \infty$ if $R^2 \to 1 \Leftrightarrow \mathsf{E}_g[(1+g)^{(n-1-p_\gamma)/2}]$ diverges (proof in Liang et al)

▶ hyper-g prior (Liang et al JASA 2008)

$$p(g) = \frac{a-2}{2}(1+g)^{-a/2}$$

or
$$g/(1+g) \sim Beta(1,(a-2)/2)$$
 need $2 < a \leq 3$

- ▶ Jeffreys prior on g corresponds to a = 2 (improper)
- ► Hyper-g/n $(g/n)(1+g/n) \sim (Beta(1,(a-2)/2))$
- ► Zellner-Siow Cauchy prior $1/g \sim G(1/2, n/2)$
- robust prior (Bayarri et al Annals of Statistics 2012
- ► Intrinsic prior (Womack et al JASA 2015)

All have prior tails for β that behave like a Cauchy distribution and (the latter 4) marginal likelihoods that can be computed using special hypergeometric functions (${}_2F_1$, Appell F_1)

USair Data

```
> library(BAS)
> data(usair, package="HH")
> poll.bma = bas.lm(log(SO2) ~ temp + log(mfgfirms) +
                                log(popn) + wind +
+
+
                                precip + raindays,
                    data=usair,
+
+
                    prior="JZS", #Jeffrey-Zellner-Siow
                    alpha=nrow(usair), # n
                    n.models=2^6.
+
+
                    modelprior = uniform(),
                    method="deterministic")
+
```

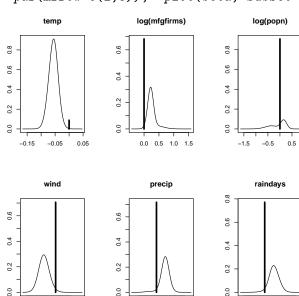
Summary

> summary(poll.bma)

```
P(B != 0 | Y) model 1 model 2 model 3
                                                          model
                 1.00000000 1.00000 1.0000000 1.0000000 1.000000
Intercept
                 0.91158530 1.00000 1.0000000 1.0000000 1.000000
temp
log(mfgfirms)
                 0.31718916 0.00000 0.0000000 0.0000000 1.000000
log(popn)
                 0.09223957 0.00000 0.0000000 0.0000000 0.000000
                 0.29394451 0.00000 0.0000000 0.0000000 1.000000
wind
                 0.28384942 0.00000 1.0000000 0.0000000 1.000000
precip
                 0.22903262 0.00000 0.0000000 1.0000000 0.000000
raindays
BF
                         NA 1.00000 0.3286643 0.2697945 0.265587
PostProbs
                         NA 0.29410 0.0967000 0.0794000 0.078100
R.2
                         NA 0.29860 0.3775000 0.3714000 0.542700
dim
                         NA 2.00000 3.0000000 3.0000000 5.000000
                         NA 3.14406 2.0313422 1.8339656 1.818248
logmarg
```

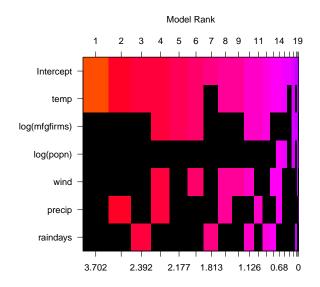
Plots

- > beta = coef(poll.bma)
- > par(mfrow=c(2,3)); plot(beta, subset=2:7,ask=F)



Posterior Distribution with Uniform Prior on Model Space

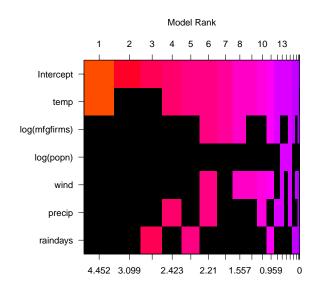
> image(poll.bma, rotate=FALSE)



Posterior Distribution with BB(1,1) Prior on Model Space

BB(1,1) Prior on Model Space

> image(poll.bb.bma, rotate=FALSE)



Summary

- ightharpoonup Choice of prior on β_{γ}
- \triangleright g-priors or mixtures of g (sensitivity)
- priors on the models (sensitivity)
- posterior summaries select a model or "average" over all models

Diabetes Example from Hoff p = 64

> source("yX.diabetes.train.txt")

> source("yX.diabetes.test.txt")

> colnames(diabetes.test)[1] = "y"

> diabetes.train = as.data.frame(diabetes.train)

> diabetes.test = as.data.frame(diabetes.test)

> set.seed(8675309)

> str(diabetes.train)

```
'data.frame':
                    342 obs. of 65 variables:
$ у
                -0.0147 -1.0005 -0.1444 0.6987 -0.2222 ...
         : num
$ age
         : num
                0.7996 -0.0395 1.7913 -1.8703 0.113 ...
$ sex
         : num
                1.064 -0.937 1.064 -0.937 -0.937 ...
                1.296 -1.081 0.933 -0.243 -0.764 ...
$ bmi
         : num
                0.459 - 0.553 - 0.119 - 0.77 0.459 \dots
$ map
         : num
$ tc
                -0.9287 -0.1774 -0.9576 0.256 0.0826 ...
         : num
$ 1d1
         : num
                -0.731 -0.402 -0.718 0.525 0.328 ...
$ hdl
                -0.911 1.563 -0.679 -0.757 0.171 ...
         : num
$ tch
                -0.0544 -0.8294 -0.0544 0.7205 -0.0544 ...
         : num
$ ltg
         : num
                0.4181 -1.4349 0.0601 0.4765 -0.6718 ...
```

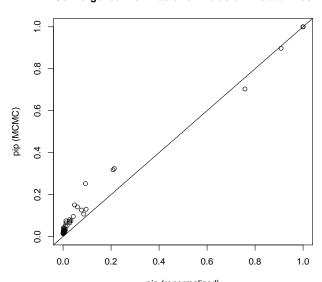
MCMC with BAS

```
> library(BAS)
> diabetes.bas = bas.lm(y ~ ., data=diabetes.train,
                        prior = "JZS",
+
                        method="MCMC",
+
                        n.models = 10000.
+
                        MCMC.iterations=150000.
+
                        thin = 10.
                        initprobs="eplogp",
+
                        force.heredity=FALSE)
+
> system.time(bas.lm(y ~ ., data=diabetes.train,
                     prior = "JZS",
+
                     method="MCMC", n.models = 10000,
                     MCMC.iterations=150000,
                     thin = 10, initprobs="eplogp",
+
                     force.heredity=FALSE))
  user system elapsed
  6.846 0.278 7.128
```

Diagnostics

> diagnostics(diabetes.bas, type="pip")

Convergence Plot: Posterior Inclusion Probabilities

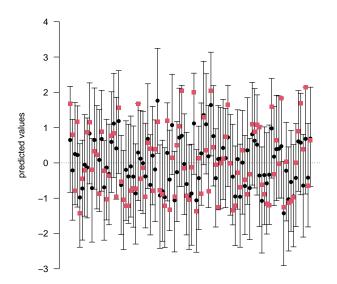


Prediction

```
> pred.bas = predict(diabetes.bas,
+
                     newdata=diabetes.test,
                     estimator="BMA",
+
                     se=TRUE)
+
> mean((pred.bas$fit- diabetes.test$y)^2)
[1] 0.4552798
> ci.bas = confint(pred.bas);
> coverage = mean(diabetes.test$y > ci.bas[,1] & diabetes.t
> coverage
Γ1 1
```

95% prediction intervals

> plot(ci.bas); points(diabetes.test\$y, col=2, pch=15)
NULL



Selection and Prediction

- ► BMA optimal for squared error loss Bayes
- ► HPM: Highest Posterior Probability model (not optimal for prediction) but for selection
- MPM: Median Probabilty model (select model where PIP > 0.5) (optimal under certain conditions; nested models)
- BPM: Best Probability Model Model closest to BMA under loss (usually includes more predictors than HPM or MPM)

Selection

```
> pred.bas = predict(diabetes.bas,
                     newdata=diabetes.test,
+
                      estimator="BPM",
+
+
                     se=TRUE)
> #MSE
> mean((pred.bas$fit- diabetes.test$y)^2)
[1] 0.4740667
> #Coverage
> ci.bas = confint(pred.bas)
> mean(diabetes.test$y > ci.bas[,1] &
       diabetes.test$y < ci.bas[,2])
[1] 0.98
```

Alternatives to MCMC

- "Stochastic Search" (no guarantee samples represent posterior)
- Variational, EM, etc to find modal model
- in BMA all variables are included, but coefficients are shrunk to 0; alternative is to use shrinkage methods without point mass at zero
- If p > n, can use a generalized inverse, but requires care for prior on $\gamma!$

Model averaging versus Model Selection - what are objectives?

Effect Estimation

- Coefficients in each model are adjusted for other variables in the model
- ➤ OLS: leave out a predictor with a non-zero coefficient then estimates are biased!
- Model Selection in the presence of high correlation, may leave out "redundant" variables;
- improved MSE for prediction (Bias-variance tradeoff)
- in BMA all variables are included, but coefficients are shrunk to 0
- Care needed for "causal" questions and confounder adjustment! With confounding, should not use plain BMA.
 Need to change prior to include potential confounders (advanced topic)