

Bayesian Analysis of A Random Coefficient Logit Model Using Aggregate Data

Peter Rossi
University of Chicago, GSB
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Joint with Renna Jiang and Puneet Manchanda

Introduction

- Often only aggregate sales/share data are available
- Berry, Levinsohn & Pakes (1995)
 - Introduce an aggregate logit model of products' market shares
 - Integrate over individual choice probabilities to obtain market shares
 - Need an aggregate error term to avoid a deterministic system after aggregation
 - Can handle endogeneity problems
 - Generalized Method of Moments (GMM)
 - Key step: invert shares to obtain mean utilities
- We conduct Bayesian inference for this model

GMM Pros and Cons

- GMM's advantages
 - Fewer distributional assumptions
 - Easier to implement
- GMM's disadvantages
 - Inefficiency
 - Inference for functions of model parameters(e.g., price elasticity, price-cost margin) of parameter estimates is difficult or computationally intensive
 - Numerical problems | multiple local maxima etc ...

Bayes Pros and Cons

- Bayes advantages
 - Could be more efficient
 - Stochastic search can handle irregular criterion functions
 - Finite sample inference w/o resort to asymptotic approximations for all functions of model parms
- Bayes disadvantages
 - Requires one additional distributional assumption (so what?)
 - Derivation of the likelihood
 - Reliable MCMC algorithm

Other Bayesian approaches

- Chen and Yang (2003)
 - No aggregate demand shocks
- Musalem, Bradlow & Raju (2006)
 - Apply to situations where there is a fixed, known, and relatively small set of consumers over which aggregate demand is formed.
- Romeo (2007)
 - Use GMM criterion as the basis of a pseudo-likelihood

Model

- Consumer i , product j ($0, 1, \dots, J$), time t

$$U_{ijt} = X_{jt} \theta_i + \eta_{jt} + \varepsilon_{ijt}$$

where X_{jt} is observed product attribute

θ_i is consumer specific coefficient

η_{jt} is unobservable (to researchers) aggregate demand shock

ε_{ijt} is iid Type I (maximum) Extreme Value (0,1)

- Assumptions
 - Consumers maximize current period utility
 - $\theta_i \sim MVN(\bar{\theta}, \Sigma)$
 - $\eta_{jt} \sim N(0, \tau^2)$

Market Share

- Utility: $U_{ijt} = X_{jt}\theta_i + \eta_{jt} + \varepsilon_{ijt}$

- Individual choice probability

$$s_{ijt} = \frac{\exp(X_{jt}\theta_i + \eta_{jt})}{1 + \sum_{k=1}^J \exp(X_{kt}\theta_i + \eta_{kt})}$$

- Market share

$$\begin{aligned} s_{jt} &= \int s_{ijt} d\Phi(\theta_i | \bar{\theta}, \Sigma) \\ &= h(\eta_t | \bar{\theta}, \Sigma, X_t) \end{aligned}$$

- s_t inherits randomness from $\eta_t = (\eta_{1t}, \dots, \eta_{Jt})'$

Likelihood

- Can derive density of s_t from density of η_t using Change-of-Variable Theorem

- Density of η_t is

$$\phi\left(h^{-1}\left(s_t \mid \bar{\theta}, \Sigma, X_t\right) \mid \tau\right)$$

Normal pdf, $\eta_{jt} \sim N(0, \tau^2)$

- Therefore, density of s_t is

$$\begin{aligned}\pi\left(s_t \mid \bar{\theta}, \Sigma, \tau, X_t\right) &= \phi\left(h^{-1}\left(s_t \mid \bar{\theta}, \Sigma, X_t\right) \mid \tau\right) J_{(\eta_t \rightarrow s_t)} \\ &= \phi\left(h^{-1}\left(s_t \mid \bar{\theta}, \Sigma, X_t\right) \mid \tau\right) \left(J_{(s_t \rightarrow \eta_t)}\right)^{-1}\end{aligned}$$

Computing h^{-1}

- h is defined by a system of J non-linear equations

$$s_{1t} = \int \frac{\exp(X_{1t}\theta_i + \eta_{1t})}{1 + \sum_{k=1}^J \exp(X_{kt}\theta_i + \eta_{kt})} d\Phi(\theta_i | \bar{\theta}, \Sigma)$$

⋮

$$s_{Jt} = \int \frac{\exp(X_{Jt}\theta_i + \eta_{Jt})}{1 + \sum_{k=1}^J \exp(X_{kt}\theta_i + \eta_{kt})} d\Phi(\theta_i | \bar{\theta}, \Sigma)$$

- BLP propose an iterative procedure which they prove to be a contraction mapping:

$$\eta_{jt}^{new} = \eta_{jt}^{old} + \ln(s_{jt}) - \ln\left(h\left(\eta_{1t}^{old}, \dots, \eta_{Jt}^{old} \mid \bar{\theta}, \Sigma\right)\right)$$

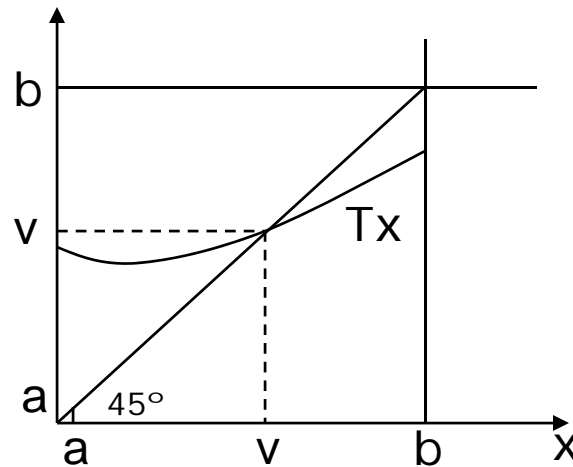
iterate until η_{jt}^{new} and η_{jt}^{old} are close "enough"

Review of contraction mapping

DEFINITION Let $T : S \rightarrow S$ be a function mapping S into itself. T is a **contraction mapping** if $\|Tx - Ty\| < \|x - y\|$, for all $x, y \in S$.

CONTRACTION MAPPING THEOREM

If T is a contraction mapping, then T has **exactly one fixed point** v in S such that $Tv = v$.



In our model, $T : R^J \rightarrow R^J$. BLP prove that $\|T\eta^{\text{new}} - T\eta^{\text{old}}\| < \|\eta^{\text{new}} - \eta^{\text{old}}\|$,
i.e., we are guaranteed for a fixed point $T\eta = \eta$.

Jacobian does not depend on $\bar{\theta}$ or τ^2

- Jacobian: $J_{(s_t \rightarrow \eta_t)} = \left\| \nabla_{\eta_t} s_t \right\|$

where

$$\frac{\partial s_{jt}}{\partial \eta_{kt}} = \begin{cases} \int -s_{ijt} s_{ikt} d\Phi(\theta_i | \bar{\theta}, \Sigma) & \text{if } k \neq j \\ \int s_{ijt} (1 - s_{ijt}) d\Phi(\theta_i | \bar{\theta}, \Sigma) & \text{if } k = j \end{cases}$$

- Re-write utility: $U_{ijt} = X_{jt} \theta_i + \eta_{jt} + \varepsilon_{ijt}$
 $= \underbrace{(X_{jt} \bar{\theta} + \eta_{jt})}_{\mu_{jt}} + X_{jt} v_i + \varepsilon_{ijt}, \quad v_i \sim N(\mathbf{0}, \Sigma)$

- Elements in Jacobian:
$$\int f(s_{igt}) d\Phi(\theta_i | \bar{\theta}, \Sigma) = \int f \left(\frac{\exp(\mu_{jt} + X_{jt} v_i)}{1 + \sum_{k=1}^J \exp(\mu_{kt} + X_{kt} v_i)} \right) d\Phi(v_i | \mathbf{0}, \Sigma) = f(\mu_t, \Sigma | X_t)$$

- But given shares (and covariates), μ_t is determined by Σ
- Thus, given shares (and covariates), Jacobian is only a function of Σ

MCMC: overview

- Recall parameters are: $\Sigma, \bar{\theta}, \tau^2$
- Re-parameterize Σ in terms of r (why?) :

$$\Sigma = U'U, \quad U = \begin{bmatrix} \exp(r_{11}) & r_{12} & \cdots & r_{1K} \\ 0 & \exp(r_{22}) & \ddots & \vdots \\ \vdots & \ddots & \ddots & r_{K-1,K} \\ 0 & \cdots & 0 & \exp(r_{KK}) \end{bmatrix}$$

- Priors

$$\bar{\theta} \sim MVN(\bar{\theta}_0, V_{\bar{\theta}})$$

$$r_{jj} \sim N(0, \sigma_j^2), \quad r_{jk} \sim N(0, \sigma_{off}^2), \quad j \neq k$$

$$\tau^2 \sim \nu_0 s_0^2 / \chi_{\nu_0}^2$$

- Posterior

$$\propto \underbrace{\prod_t \pi(s_t | \bar{\theta}, r, \tau, X_t)}_{\text{likelihood}} \times \text{Prior}(\bar{\theta}, r, \tau)$$

More on the re-parameterization

$$\Sigma = U'U = \begin{bmatrix} \exp(2r_{11}) & r_{12} \exp(r_{11}) & r_{13} \exp(r_{11}) & r_{14} \exp(r_{11}) \\ & r_{12}^2 + \exp(2r_{22}) & r_{12}r_{13} + r_{23} \exp(r_{22}) & r_{12}r_{14} + r_{24} \exp(r_{22}) \\ \text{symmetric} & & r_{13}^2 + r_{23}^2 + \exp(2r_{33}) & r_{13}r_{14} + r_{23}r_{24} + r_{34} \exp(r_{33}) \\ & & & r_{14}^2 + r_{24}^2 + r_{34}^2 + \exp(2r_{44}) \end{bmatrix}$$

- Priors: $r_{jj} \sim N(0, \sigma_j^2)$, $r_{jk} \sim N(0, \sigma_{off}^2)$, $j \neq k$
- Implied prior variance and mean on diagonals of Σ are ($k=1,2,3,4$):

$$\text{var}[\Sigma_{kk}] = 2(k-1)\sigma_{off}^4 + \exp(8\sigma_k^2) - \exp(4\sigma_k^2)$$

$$E[\Sigma_{kk}] = (k-1)\sigma_{off}^2 + \exp(2\sigma_k^2)$$
- Implied prior variance and mean on off-diagonals of Σ are ($k=1,2,3$):

$$\text{var}[\Sigma_{k,(k+1)}] = (k-1)\sigma_{off}^4 + \sigma_{off}^2 \exp(2\sigma_k^2)$$

$$E[\Sigma_{k,(k+1)}] = 0$$
- Goal: let diagonals of Σ have same prior variance
- So, set $\sigma_1^2 = 0.50666596$, $\sigma_{off}^2 = 1.00001292477193$, $\sigma_2^2 = 0.5019265$, $\sigma_3^2 = 0.4969934$, $\sigma_4^2 = 0.4918498$

$$\Rightarrow \text{var}[\Sigma_{11}] = \text{var}[\Sigma_{22}] = \text{var}[\Sigma_{33}] = \text{var}[\Sigma_{44}] = 50$$

$$\text{var}[\Sigma_{12}] = 2.7548, \text{var}[\Sigma_{23}] = 3.7288, \text{var}[\Sigma_{34}] = 4.7021$$

$$E[\Sigma_{11}] = 2.7548, E[\Sigma_{22}] = 3.7288, E[\Sigma_{33}] = 4.7020, E[\Sigma_{44}] = 5.6744$$

$$E[\Sigma_{12}] = E[\Sigma_{23}] = E[\Sigma_{34}] = 0$$

MCMC algorithm

Two sets of conditional draws

$$r \mid \bar{\theta}, \tau^2, \{s_t, X_t\}_{t=1}^T, \sigma_{r_diag}^2, \sigma_{r_off}^2$$
$$\bar{\theta}, \tau^2 \mid r, \{s_t, X_t\}_{t=1}^T, \bar{\theta}_0, V_{\bar{\theta}}, \nu_0, s_0^2$$

1. Random-Walk Metropolis Chain to propose for r

$$r^{new} = r^{old} + MVN(\mathbf{0}, \sigma^2 D_r)$$

2. Gibbs sampler for $\bar{\theta}$ and τ^2
 - A univariate Bayes regression:

$$\mu_{jt} = X_{jt} \bar{\theta} + \eta_{jt}, \quad \eta_{jt} \sim N(0, \tau^2)$$

MCMC part of the code...

```
# ----- (1) Gibbs Sampler for thetabar and taosq -----
output=rniregG(y=mu,X=X,XpX=XpX,Xpy=crossprod(X,mu),sigmasq=taosq,
              A=Athetabar,betabar=thetabar0,nu=nu0,ssq=s0sq)
thetabar=output$betadraw
taosq=output$sigmasqdraw

# ----- (2) Metropolis for r -----
# Random-Walk Chain
rN=r+mvrnorm(1,rep(0,(K*(K+1)/2)),varn_r)

ON=Loglhd(rN,mu,thetabar,taosq)
prior_old=sum(-r[1:K]^2/2/sigmasqR_DIAG)+sum(-r[(K+1):(K*(K+1)/2)]^2/2/sigmasqR_off)
prior_new=sum(-rN[1:K]^2/2/sigmasqR_DIAG)+sum(-rN[(K+1):(K*(K+1)/2)]^2/2/sigmasqR_off)

# Evaluate old r (mu) at new (thetabar,taosq)
eta=mu-X%*%thetabar
llhd_old=sum(log(dnorm(eta,sd=sqrt(taosq))))+OO$sumlogjacob

ratio=exp(ON$llhd+prior_new-llhd_old-prior_old)
alphaS=min(1,ratio) # S stands for Sigma
if (runif(1)<=alphaS) {
  r=rN; OO=ON; ns=ns+1; mu=OO$mu
}
```

Brute-Force log-likelihood code...

```

Loglhd_slow = function(thetabar,r,taosq,mu){
  # Purpose: Evaluate log likelihood. Sigma is re-parameterized as r.

  # (1). Transform r to L, where Sigma=LL'
  L=diag(exp(r[1:K]))
  L[lower.tri(L)]=r[(K+1):(K*(K+1)/2)]

  # (2). At given L, do inversion to get mu. Then compute eta
  temp=invert_slow(L,mu,v,crit,T,H,J,lnactS,indTHJ,indJTH)
  mu = temp$mu; prob = temp$prob; niter = temp$niter
  eta=mu-X%*%thetabar

  # (3). Jacobian
  # Form J diagonal elements at each time t
  diagonal=rowMeans(prob*(1-prob)) # TJ by 1 vector

  # Form the off diagonal elements
  dd=-prob%*%t(prob)/H # TJ by TJ
  cc=aaa*dd+diag(diagonal)#TJ by TJ matrix: block diagonal

  for (t in 1:T){
    cct=cct+diag(diagonal[t,t]) # (t)th block of cc
    logjacob[t]=-log(abs(det(cct)))
  }
  # (4). Form Log Likelihood
  sumlogjacob=sum(logjacob)
  llhd=sum(log(dnorm(eta,sd=sqrt(taosq))))+sumlogjacob

  list(llhd=llhd,mu=mu,niter=niter,sumlogjacob=sumlogjacob)
}

```

$$\int s_{hjt} (1 - s_{hjt}) d\Phi(\theta_h)$$

$$\int -s_{hjt} s_{hkt} d\Phi(\theta_h)$$

$$cct = \nabla_{\eta_t} s_t$$

Slow inversion code

```

invert_slow =
function(L,mu,v,crit,T,H,J,lnactS,indTHJ,indJTH){

# Purpose: Invert observed shares S at give L to get mean utility mu's.

niter=0 # number of iterations taken for the inversion
munew=mu # starting value
muold=munew/2
upart=X%*%L%*%v
while (max(abs((muold-munew)/munew))>crit){

    muold=munew
    num=exp(upart+ muold) # JT by H numerator
    den1=matrix(double(T*H),nrow=T)
    for (t in 1:T){
        den1[t,]=1+colSums(num[((t-1)*J+1):(t*J),]) #T by H
    }
    den=matrix(rep(den1,each=J),ncol=H) #replc each t J times,JT by H
    prob=num/den # JT by H
    sh=t(matrix(rowMeans(prob), nrow=J)) # T by J predicted share
    munew=t(matrix(muold,nrow=J))+log(S)-log(sh) # T by J
    munew=as.vector(t(munew)) # length JT vector
    niter=niter+1
}
List(mu=munew,prob=prob,niter=niter)
}

```

$$1 + \sum_{k=1}^J \exp(X_{kt} \theta_h + \eta_{kt})$$



profile it (on 1000 iterations)...

```
$by.total
      total.time total.pct self.time self.pct
rRLogit_slow    663.68   100.0    0.08    0.0
Loglhd_slow     660.68    99.5    7.96    1.2
invert_slow     513.98    77.4   46.46    7.0
colSums         262.24    39.5  127.60   19.2
exp             98.34    14.8   98.34   14.8
is.data.frame   81.74    12.3    3.58    0.5
inherits        78.16    11.8   69.46   10.5
matrix          74.30    11.2   22.68    3.4
as.vector       52.18     7.9   43.52    6.6
diag            49.26     7.4    8.74    1.3
array           40.30     6.1   40.26    6.1
%*%            30.04     4.5   30.04    4.5
+              29.24     4.4   29.24    4.4
prod           28.52     4.3   26.04    3.9
log            25.24     3.8    7.30    1.1
/             24.14     3.6   24.14    3.6
det            17.48     2.6    2.02    0.3
*             14.14     2.1   14.14    2.1
determinant    11.62     1.8    1.98    0.3
t             11.36     1.7    1.68    0.3
determinant.matrix 9.64     1.5    2.74    0.4
rowMeans       7.70     1.2    6.42    1.0
:              7.56     1.1    7.56    1.1
-              6.50     1.0    6.50    1.0
<              5.62     0.8    5.62    0.8
>              4.56     0.7    4.56    0.7
$              3.34     0.5    3.34    0.5
storage.mode<- 2.96     0.4    0.84    0.1
.Call          2.30     0.3    2.30    0.3
storage.mode   2.04     0.3    1.14    0.2
length         1.78     0.3    1.78    0.3
```

Make it faster: Vectorize it!

```
invert =  
function(L,mu,v,crit,T,H,J,lnactS,indTHJ,indJTH){  
  
# Purpose: Invert observed shares S at give L to get mean utility mu's.  
  
niter=0 # number of iterations taken  
munew=mu # starting value  
muold=munew/2  
upart=X%%L%%v  
while (max(abs((muold-munew)/munew))>crit){  
  
    muold=munew  
    num=exp(upart+ muold) # num is JT x H  
    dim(num)=NULL # convert num to JTH vector  
    num=num[indTHJ] # convert num to THJ vector  
    dim(num)=c(T*H,J) # convert num to TH * J matrix  
    den=1+rowSums(num) # TH vector  
    prob=num/den # TH * J matrix  
    dim(prob)=NULL # convert prob to THJ vector  
    prob=prob[indJTH] # convert prob to JTH vector  
    dim(prob)=c(J*T,H) # convert prob to JT * H matrix  
    sh=rowMeans(prob) # JT vector  
    munew=muold+lnactS-log(sh) # JT vector  
    niter=niter+1  
}  
list(mu=munew,prob=prob,niter=niter)  
}
```

No loop!

Matrix
divided
by a
vector

More on the re-indexing function

```
JTH_THJ=function(J,H,T){
#
# function to convert and index a vector ordered j by t by h (i.e. j
# varies faster than t than h) into a vector ordered t by h by j
#
ind=double(J*H*T)
cnt=1
for (j in 1:J){
  for (h in 1:H) {
    for (t in 1:T) {
      ind[cnt]=(t-1)*J+(h-1)*(T*J)+j
      cnt=cnt+1
    }
  }
}
return(ind)
}
```

- Similarly, THJ_JTH is a function that converts and indexes a vector ordered t by h by j (i.e. t varies faster than h than j), into a vector ordered j by t by h.
- Pre-compute the two indices:
`indTHJ=JTH_THJ(J,H,T)`
`indJTH=THJ_JTH(J,H,T)`

Eliminate Determinants

- Work on algebra to eliminate "det":

$$cct = U'U$$

$$|cct| = |U'| |U| = \left(\prod_{j=1}^J \text{diag}(U) \right)^2$$

$$\begin{aligned} \log(|cct|^{-1}) &= -\log \left(\prod_{j=1}^J \text{diag}(U) \right)^2 \\ &= -2 \sum_{j=1}^J \log(\text{diag}(U)) \end{aligned}$$

- New code:

```
for (t in 1:T){
  cct=cc[((t-1)*J+1):(t*J),((t-1)*J+1):(t*J)] # (t)th block of cc
  logjacob[t]=-2*sum(log(diag(chol(cct))))
  # old code:
  # logjacob[t]=-log(abs(det(cct)))
}
```

profile it again (on 1000 iterations)...

Savings
of
57.5%!

\$by.total	total.time	total.pct	self.time	self.pct
rRCLogit	282.08	100.0	0.04	0.0
Loglhd	279.32	99.0	14.36	5.1
invert	170.66	60.5	47.84	17.0
exp	91.66	32.5	91.66	32.5
/	27.22	9.6	27.22	9.6
diag	23.56	8.4	10.76	3.8
log	20.88	7.4	3.96	1.4
crossprod	19.98	7.1	19.30	6.8
sum	19.38	6.9	0.92	0.3
*	12.74	4.5	12.74	4.5
chol	11.12	3.9	2.50	0.9
-	10.46	3.7	10.46	3.7
rowSums	8.88	3.1	8.42	3.0
diag<-	7.12	2.5	7.02	2.5
rowMeans	6.04	2.1	4.82	1.7
as.matrix	5.26	1.9	4.58	1.6
+	4.72	1.7	4.72	1.7
.Call	1.74	0.6	1.74	0.6
nrow	0.96	0.3	0.86	0.3
is.data.frame	0.96	0.3	0.04	0.0
inherits	0.94	0.3	0.24	0.1
t	0.76	0.3	0.14	0.0
as.matrix.default	0.66	0.2	0.58	0.2
t.default	0.62	0.2	0.62	0.2
mvrnorm	0.62	0.2	0.04	0.0
%*%	0.58	0.2	0.58	0.2
min	0.56	0.2	0.52	0.2
:	0.52	0.2	0.52	0.2
JTH_THJ	0.52	0.2	0.44	0.2
THJ_JTH	0.48	0.2	0.42	0.1
runiregG	0.46	0.2	0.02	0.0

GMM

- Berry, Levinsohn & Pakes (1995):

$$\mu_{jt} = X_{jt} \bar{\theta} + \eta_{jt}$$

- Theoretical moments :

$$E[Z_t' \eta_t] = 0$$

- Sample analog:

$$\hat{m}_T(\bar{\theta}, \Sigma) = \frac{1}{T} \sum_{t=1}^T Z_t' (\hat{\mu}_t(\Sigma) - X_t \bar{\theta})$$

- GMM objective:

$$\min_{\bar{\theta}, \Sigma} \hat{m}_T(\bar{\theta}, \Sigma)' A^{-1} \hat{m}_T(\bar{\theta}, \Sigma)$$

- GMM search can be limited to only Σ , because $\bar{\theta}$ can be concentrated out as follows,

$$\hat{\bar{\theta}}(\Sigma) = (X'ZA^{-1}Z'X)^{-1} X'ZA^{-1}Z'\hat{\mu}(\Sigma)$$

Implementing GMM

- How to form Z ?
 - Total # of parameters for GMM: $\dim(\theta) + \dim(r)$.
 - Form Z by expanding exogenous variables in X or other instruments into polynomials, exponentials, logarithms, and interact with brand intercepts
- How to form A ? Two-step GMM
 - Step 1: Let $A = \frac{1}{T} \sum_{t=1}^T Z_t' Z_t$

Minimize the GMM objective \Rightarrow obtain the residuals $\hat{\eta}_{jt}^{(1)}$, (1) stands for Step 1.
 - Step 2: Construct a new $A = \frac{1}{T^2} \sum_{t=1}^T Z_t' \hat{\eta}_t^{(1)} \hat{\eta}_t^{(1)'} Z_t$

Minimize the GMM objective again.

After convergence, re-start the optimization routine from the converged estimates to ensure that the optimization converged.
- Take the final converged results $\Rightarrow \hat{\Sigma}_{GMM}, \hat{\theta}_{GMM}, \hat{\eta}_{jt}^{(2)}$, and $\hat{\tau}_{GMM} = sd(\hat{\eta}^{(2)})$

A sampling experiment

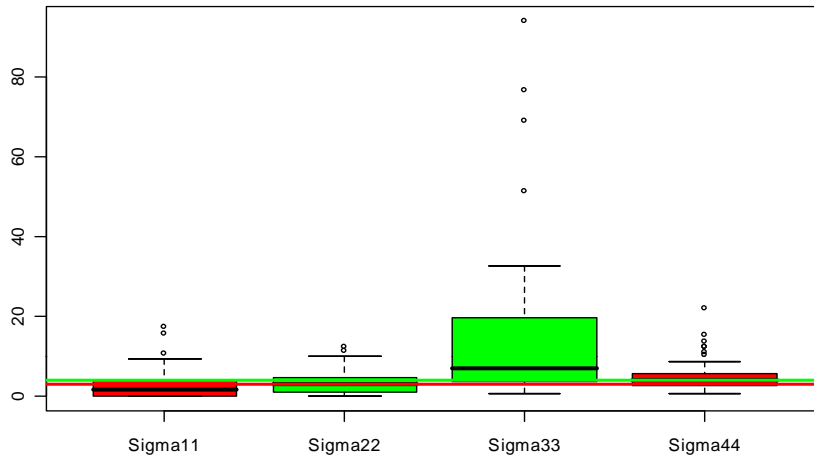
- $J=3$ brands, one outside good
- X contains 3 brand intercepts and a uniform[0,1] distributed attribute
- $T=300$ time periods
- Base parameters

$$\bar{\theta}_{base} = (-2, -3, -4, -5)$$
$$\Sigma_{base} = \begin{bmatrix} 3 & 2 & 1.5 & 1 \\ & 4 & -1 & 1.5 \\ & & 4 & -0.5 \\ & & & 3 \end{bmatrix}$$
$$\tau_{base}^2 = 1$$

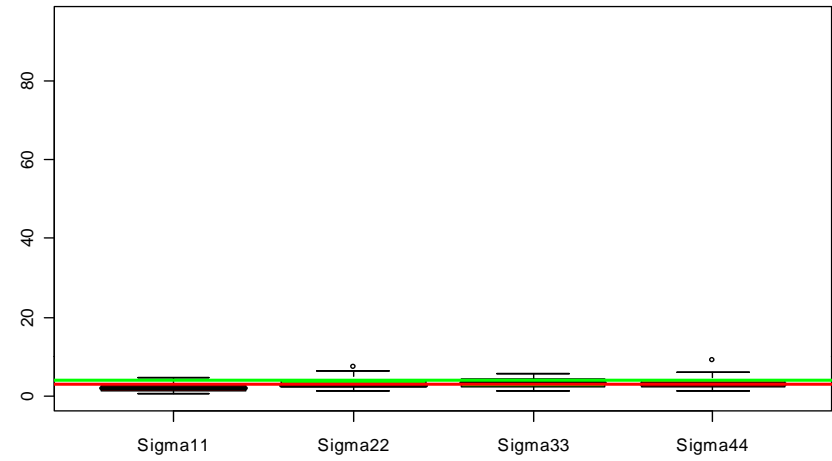
- Generate 50 datasets
 - E.g, in one dataset, implied average shares (stdev) over time for each brand:
8.0% (8.5%), 5.1% (4.8%), 1.7% (2.6%)

Results: diagonals of Σ

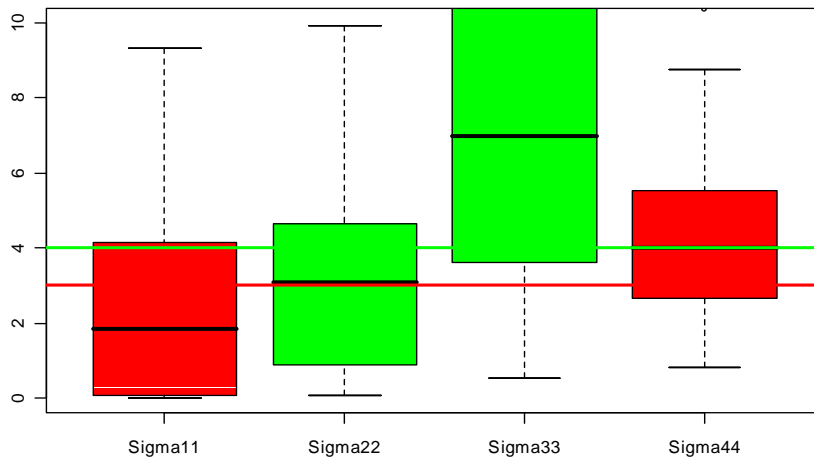
GMM point estimates across 50 reps within main cell



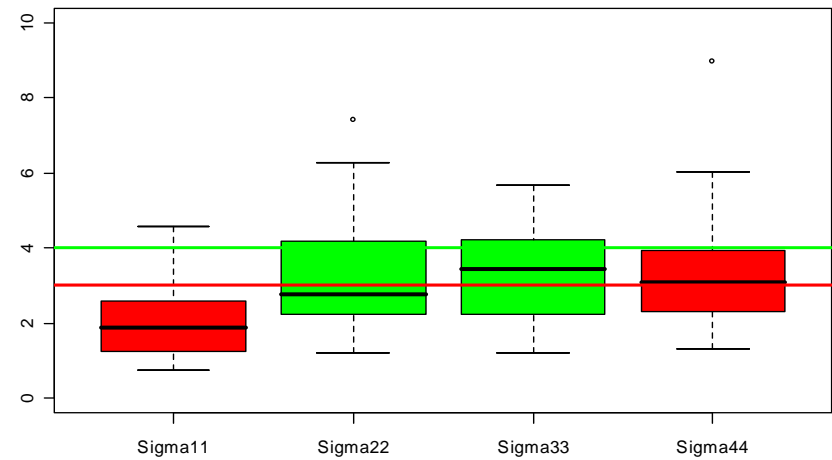
Bayesian posterior mean across 50 reps within main cell



GMM point estimates across 50 reps within main cell: zoomed in

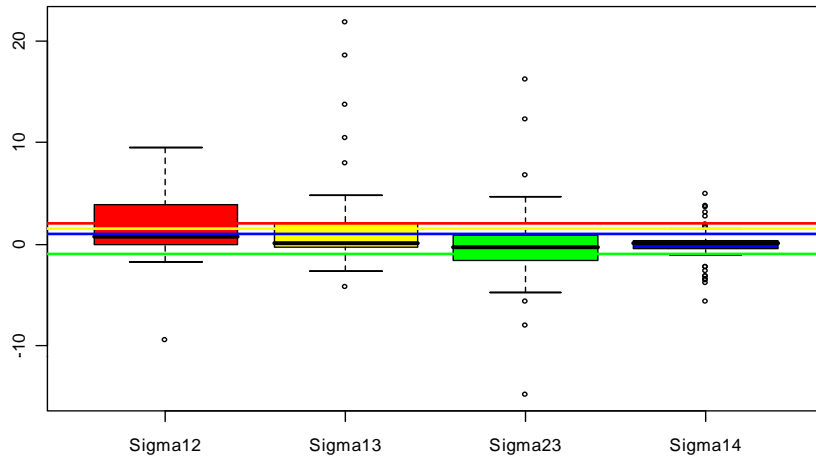


Bayesian posterior mean across 50 reps within main cell: zoomed in

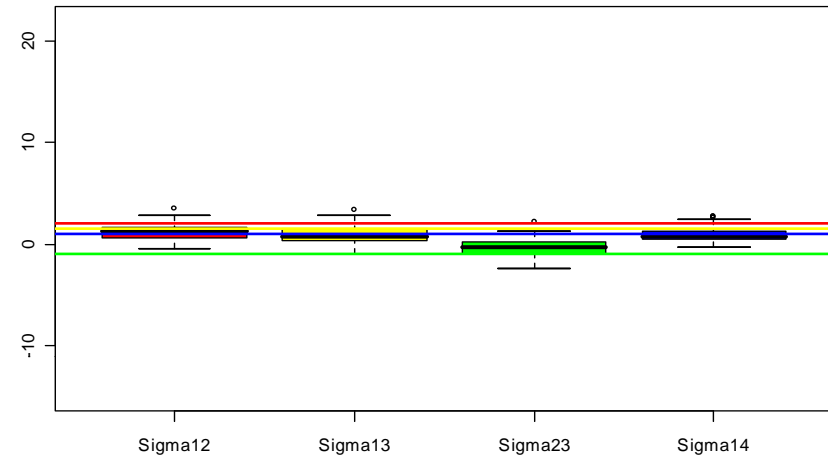


Results: off-diagonals of Σ

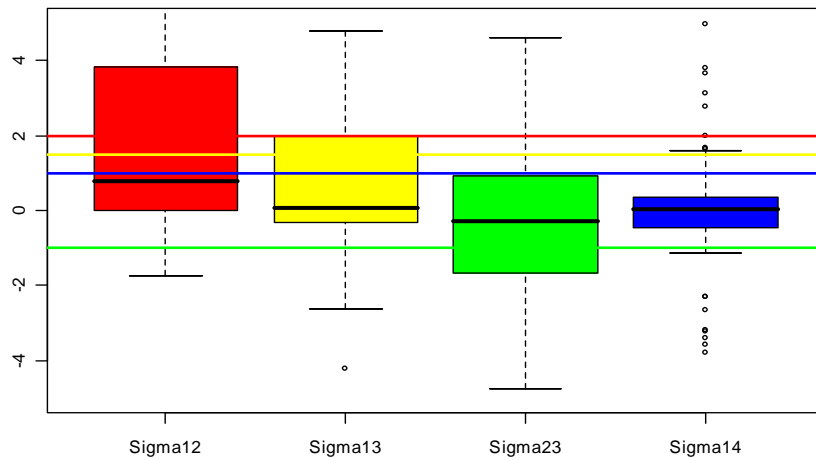
GMM point estimates across 50 reps within main cell



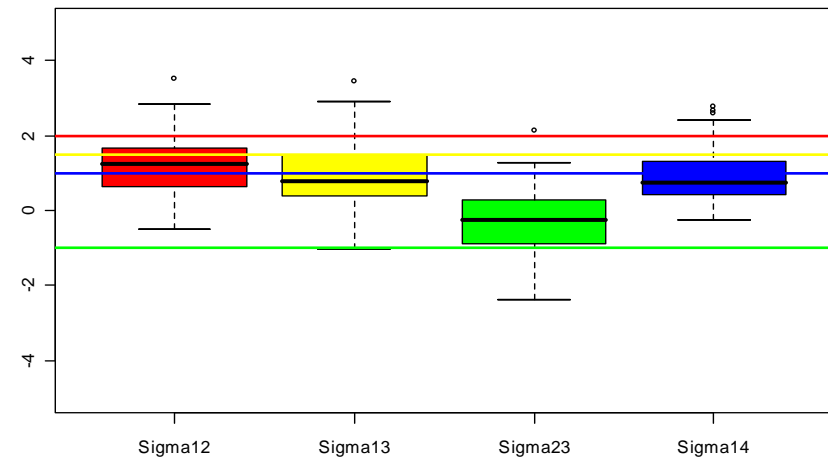
Bayesian posterior mean across 50 reps within main cell



GMM point estimates across 50 reps within main cell: zoomed in

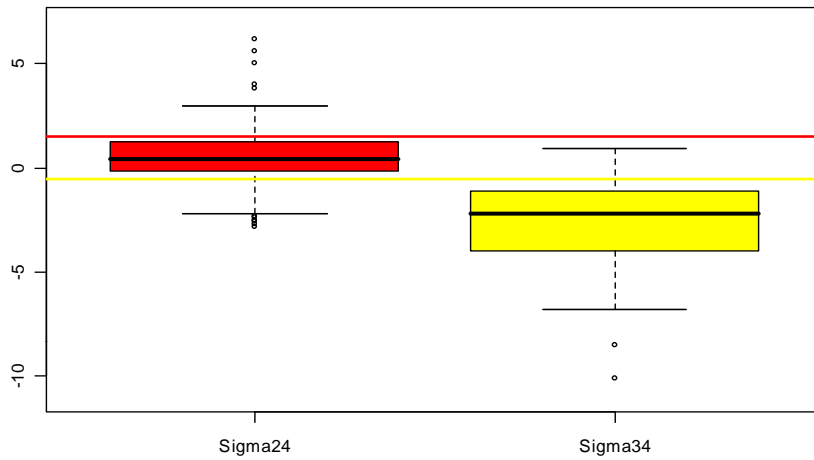


Bayesian posterior mean across 50 reps within main cell

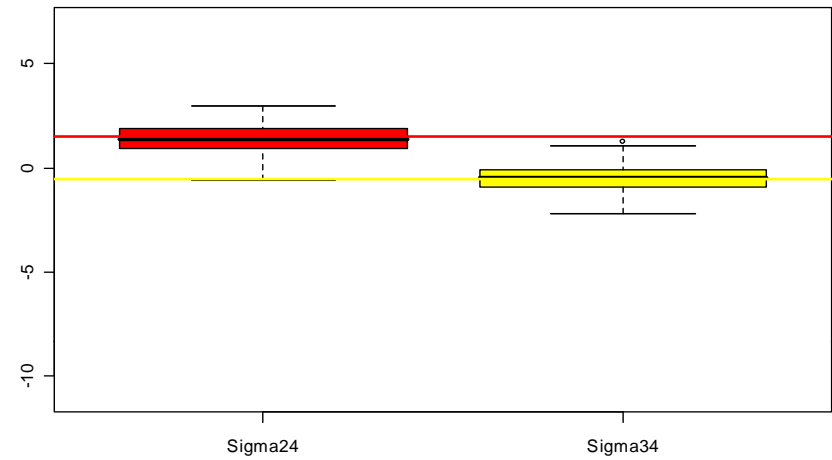


Results: off-diagonals of Σ

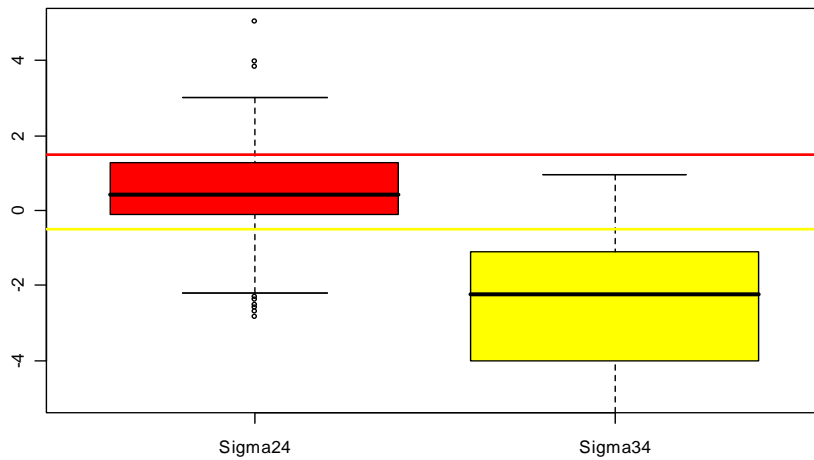
GMM point estimates across 50 reps within main cell



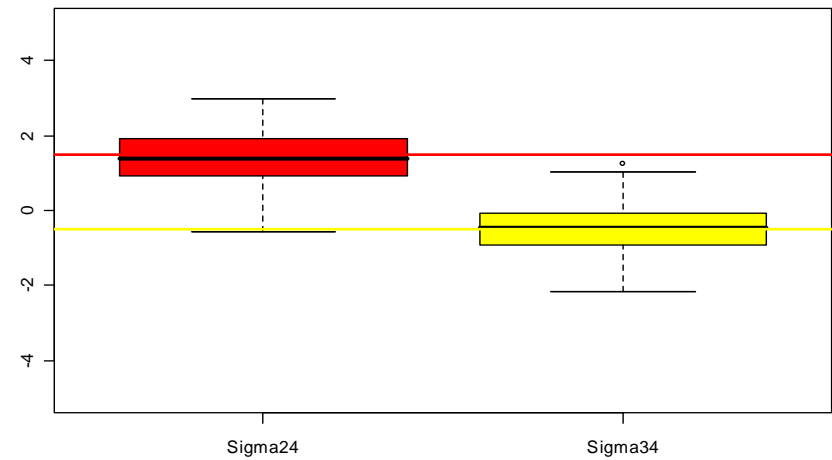
Bayesian posterior mean across 50 reps within main cell



GMM point estimates across 50 reps within main cell: zoomed in

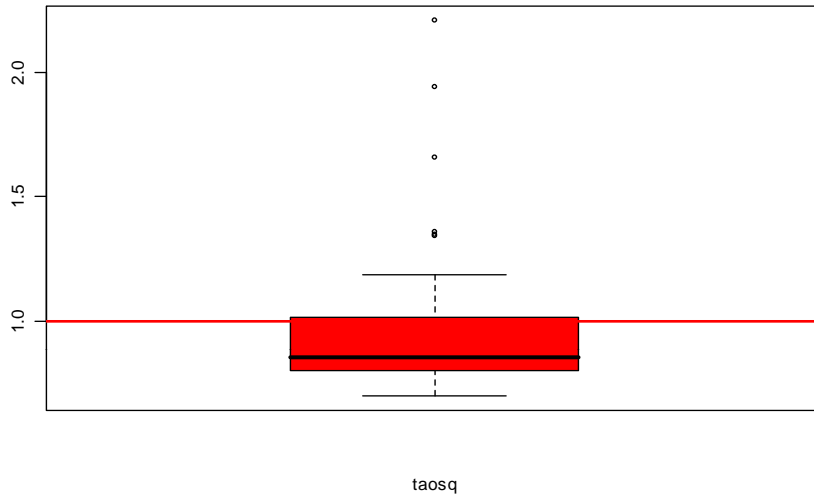


Bayesian posterior mean across 50 reps within main cell

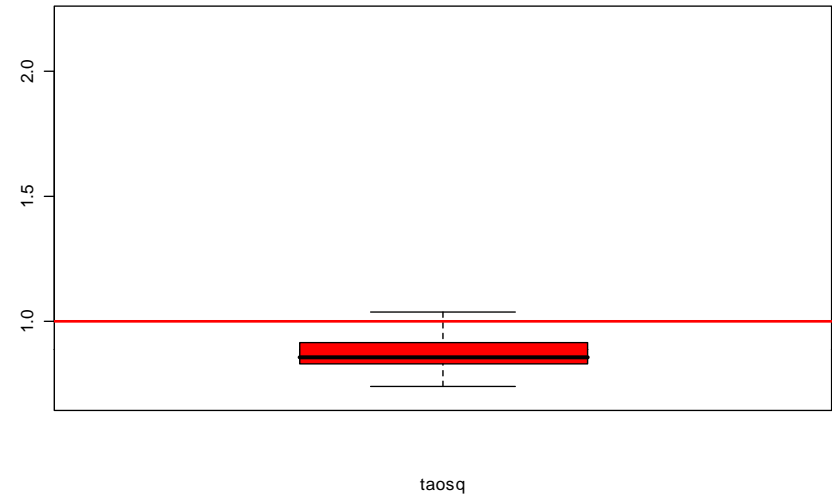


Results: $\tau^2, \bar{\theta}$

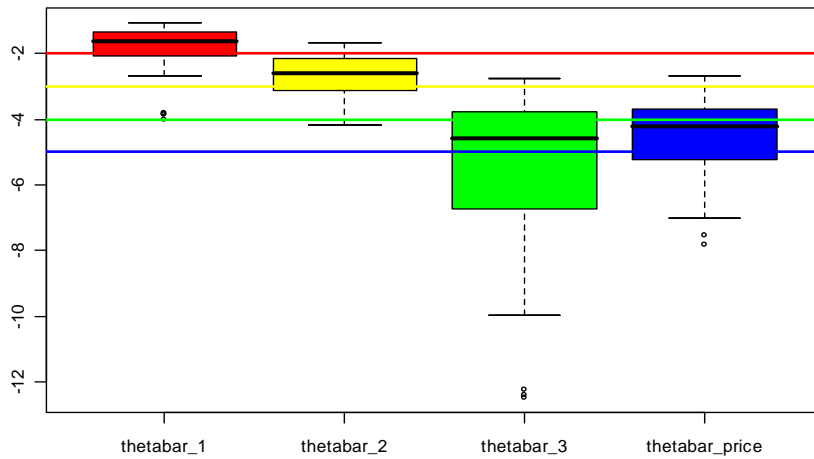
GMM point estimates across 50 reps within main cell



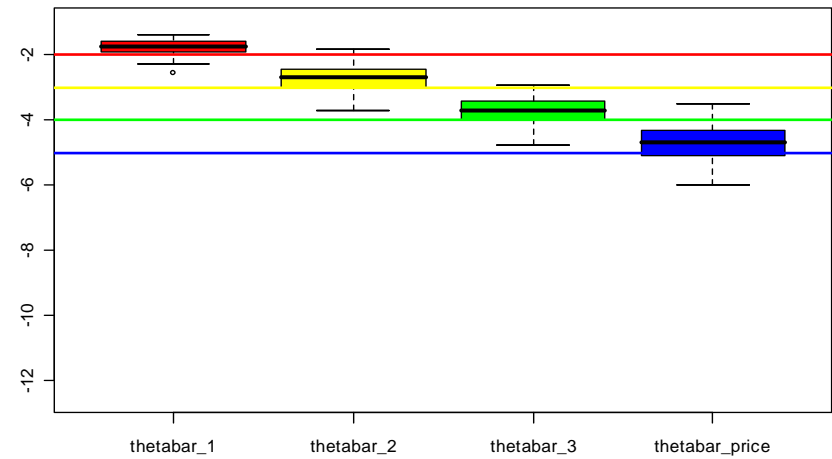
Bayesian posterior mean across 50 reps within main cell



GMM point estimates across 50 reps within main cell



Bayesian posterior mean across 50 reps within main cell



MSE and bias

	MSE		Bias	
	Bayes	GMM	Bayes	GMM
τ^2	0.02	0.09	-0.13	-0.03
$\bar{\theta}_1$	0.11	0.54	0.22	0.13
$\bar{\theta}_2$	0.26	0.54	0.25	0.29
$\bar{\theta}_3$	0.25	8.51	0.27	-1.51
$\bar{\theta}_{price}$	0.41	1.71	0.28	0.47
Σ_{11}	1.94	14.89	-1.04	0.13
Σ_{22}	2.63	9.52	-0.70	-0.46
Σ_{33}	1.95	498.86	-0.62	10.68
Σ_{44}	2.23	21.73	0.45	2.19
Σ_{12}	1.24	10.02	-0.75	-0.29
Σ_{13}	1.15	23.49	-0.56	0.35
Σ_{23}	1.41	19.77	0.67	0.86
Σ_{14}	0.54	5.13	-0.06	-1.09
Σ_{24}	0.63	5.19	-0.08	-0.91
Σ_{34}	0.52	10.05	0.02	-2.22

GMM: 0.7 hr/rep, Bayes: 2.2 hr/rep (Pentium 4, CPU 3 GHz, 1GB RAM)

GMM Estimation: numerical stability

The default setting in GMM estimation

- Optimizer: "optim" in R
- Search algorithm: gradient-based "BFGS" does not work (can't improve objective value, keeps stopping, etc). Use simplex-based "Nelder-Mead." All samples "converged" in the sense of meeting conv criterion
- Convergence criterion for simplex search: reltol=1e-6
 - Relative convergence tolerance. Percentage change in objective function < reltol
- Starting value for Sigma: identity matrix
- Convergence criterion for the inversion step: crit=1e-5
- # of heterogeneity draws to evaluate share integral: H=50 (seed=456)

Sensitivity checks on one outlier of GMM estimate

- Start Sigma from true value
- Tighten reltol=1e-10
- Tighten crit=1e-10

Examine one GMM outlier (Rep 29)

- Default

$$\hat{\Sigma} = \begin{bmatrix} 15.1 & 15.5 & 8.2 & 3.0 \\ & 16.3 & 7.6 & 4.7 \\ & & 18.4 & -4.6 \\ & & & 13.1 \end{bmatrix} \quad \text{objval} = .298$$

$$\hat{\theta} = [-3.5, -4.6, -6.7, -9.6], \quad \hat{\tau}^2 = 2.2$$

- Start from true

$$\hat{\Sigma} = \begin{bmatrix} 15.7 & 17.2 & 1.7 & -2.6 \\ & 19.6 & 2.0 & -3.2 \\ & & 21.4 & -5.7 \\ & & & 20.5 \end{bmatrix} \quad \text{objval} = .179$$

$$\hat{\theta} = [-3.6, -5.2, -7.5, -7.9], \quad \hat{\tau}^2 = 1.7$$

- Tighten reltol to 1e-10:

$$\hat{\Sigma} = \begin{bmatrix} 11.8 & 11.0 & 15.4 & 1.4 \\ & 10.5 & 13.4 & 3.7 \\ & & 28.5 & -6.9 \\ & & & 21.6 \end{bmatrix} \quad \text{objval} = .127$$

$$\hat{\theta} = [-3.2, -3.8, -8.0, -8.2], \quad \hat{\tau}^2 = 1.7$$

- Tighten crit=1e-10:

$$\hat{\Sigma} = \begin{bmatrix} 9.6 & 9.9 & 9.7 & 3.7 \\ & 10.6 & 9.1 & 4.6 \\ & & 15.7 & -0.008 \\ & & & 10.8 \end{bmatrix} \quad \text{objval} = .348$$

$$\hat{\theta} = [-2.8, -3.9, -6.1, -9.1], \quad \hat{\tau}^2 = 1.6$$

Address mis-specification concerns

- So far, we first generate iid η_{jt} from $N(0, 1)$, then conduct inferences assuming $\eta_{jt} \sim N(0, \tau^2)$
- Now let's investigate the performance of our estimator in situations where the model is mis-specified.
- Specifically, we generate η_{jt} from as follows, then fit the model assuming $\eta_{jt} \sim N(0, \tau^2)$
 - Conditional Heteroskedasticity
 - AR(1): 0.9
 - Asymmetric Beta distribution
 - Symmetric Beta distribution

Mis-specification: Heteroskedasticity

- Instead of homoskedastic shocks, generate shocks from

$$\eta_{jt} \sim N(0, V_{jt} \equiv f(X_{jt}))$$

- To keep things comparable, we require $E[V_{jt}] = 1$
- So,

$$1 = E[V_{jt}] = E[\exp(a + bP_{jt})] = \frac{\exp(a)}{b}(\exp(b) - 1)$$

where $P_{jt} \sim Unif[0,1]$

- Set $b=1$, then $a = -\log(\exp(1) - 1) \approx -0.5413$

Mis-specification: AR(1)

- Instead of iid, generate shocks for product j ($j=1,2,3$) according to

$$\eta_{j,t+1} = \rho\eta_{j,t} + u_{j,t+1} \quad u_{j,t} \stackrel{iid}{\sim} N(0, \sigma_u^2)$$

- We require $\text{var}[\eta_{j,t}] = 1$

- So,

$$1 = \text{var}[\eta_{j,t}] = \frac{\sigma_u^2}{1 - \rho^2}$$

- Set $\rho = 0.9$, we get $\sigma_u \approx 0.4359$

Mis-specification: Asymmetric Beta

- Instead of Normal shocks, generate

$$\eta_{jt} \sim d \cdot \text{Beta}[\alpha, \beta] - c$$

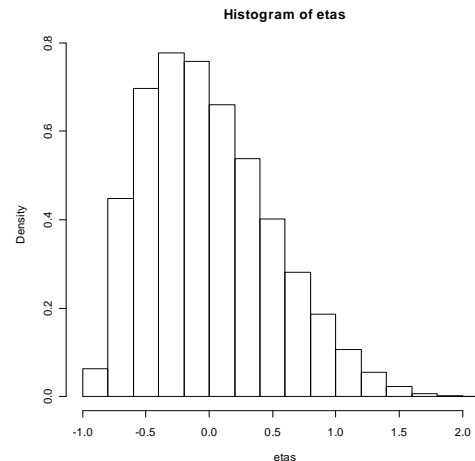
- Set $\alpha = 2, \beta = 5$ to get a logNormal-shaped asymmetric Beta distribution

- We require that η be centered at zero and $sd[\eta_{jt}] = 0.5$

- So, $0.5 = sd[\eta_{jt}] = d \sqrt{\frac{\alpha\beta}{(\alpha+\beta)^2(\alpha+\beta+1)}} = d \sqrt{\frac{2 \times 5}{(2+5)^2(2+5+1)}} \Rightarrow d \approx 3.1305$

- Finally, calculate the de-mean factor c to keep η centered at zero:

$$c = d \cdot \frac{\alpha}{\alpha + \beta} \approx 0.8944$$



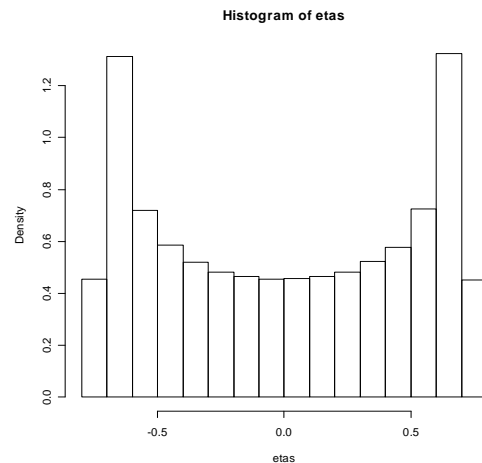
Mis-specification: Symmetric Beta

- Instead of Normal shocks, generate

$$\eta_{jt} \sim d \cdot \text{Beta}[\alpha, \beta] - c$$

- Set $\alpha = 0.5, \beta = 0.5$ to get a U-shaped symmetric Beta distribution
- We require that η centered at zero and $sd[\eta_{jt}] = 0.5$
- Follow the same procedure as asymmetric Beta, we get

$$d \approx 1.4142, c \approx 0.7071$$



MSE and bias in mis-specified cells

		MSE		Bias	
		Bayes	GMM	Bayes	GMM
Σ_{11}	Main	1.94	14.89	-1.04	0.13
	Hetro	11.07	25.81	1.85	0.23
	AR1	3.91	35.43	-0.38	0.32
	AsyBeta	2.17	66.28	-1.22	1.20
	SymBeta	2.14	8.49	-1.19	-1.05
Σ_{22}	Main	2.63	9.52	-0.70	-0.46
	Hetro	15.73	26.65	2.78	-0.33
	AR1	5.3	181.16	0.13	0.83
	AsyBeta	4.00	87.09	-1.71	1.96
	SymBeta	2.34	38.38	-0.92	0.46
Σ_{33}	Main	1.95	498.86	-0.62	10.68
	Hetro	40.2	566.35	3.6	12.35
	AR1	8.08	1927.91	0.50	15.95
	AsyBeta	3.47	601.03	-1.33	8.83
	SymBeta	3.04	163.88	-0.42	6.04
Σ_{44}	Main	2.23	21.73	0.45	2.19
	Hetro	5.12	23.11	1.16	2.33
	AR1	5.41	24.05	1.44	2.40
	AsyBeta	0.71	64.72	-0.16	2.73
	SymBeta	2.42	21.58	0.50	1.91
Σ_{12}	Main	1.24	10.02	-0.75	-0.29
	Hetro	1.75	22.0	0.46	-0.25
	AR1	3.14	13.49	-0.29	-0.35
	AsyBeta	0.97	23.92	-0.79	-0.84
	SymBeta	1.22	9.84	-0.80	-0.87

MSE and bias in mis-specified cells

		MSE		Bias	
		Bayes	GMM	Bayes	GMM
Σ_{13}	Main	1.15	23.49	-0.56	0.35
	Hetro	3.85	53.12	0.40	1.00
	AR1	2.09	49.7	-0.63	0.50
	AsyBeta	0.50	10.28	-0.54	-0.77
	SymBeta	0.77	8.08	-0.37	-0.82
Σ_{23}	Main	1.41	19.77	0.67	0.86
	Hetro	3.85	49.15	0.45	2.13
	AR1	3.86	173.38	0.83	4.04
	AsyBeta	1.50	15.91	0.99	1.15
	SymBeta	1.43	17.75	0.61	1.95
Σ_{14}	Main	0.54	5.13	-0.06	-1.09
	Hetro	4.05	17.47	-1.34	-1.70
	AR1	1.13	8.85	0.16	-0.83
	AsyBeta	0.41	12.42	-0.16	-0.53
	SymBeta	0.78	4.95	0.19	-0.07
Σ_{24}	Main	0.63	5.19	-0.08	-0.91
	Hetro	3.48	17.44	-1.02	-1.72
	AR1	1.42	9.47	0.08	-1.13
	AsyBeta	0.45	16.02	-0.30	-0.53
	SymBeta	0.73	8.15	0.22	0.10
Σ_{34}	Main	0.52	10.05	0.02	-2.22
	Hetro	3.01	31.25	-1.08	-3.02
	AR1	1.16	28.4	0.07	-1.66
	AsyBeta	0.3	27.72	-0.03	-0.44
	SymBeta	0.74	10.43	0.09	-0.25

MSE and bias in mis-specified cells

		MSE		Bias	
		Bayes	GMM	Bayes	GMM
τ^2	Main	0.02	0.09	-0.13	-0.03
	Hetro	0.009	0.134	-0.021	-0.011
	AR1	0.049	0.227	-0.172	-0.058
	AsyBeta	0.002	0.007	-0.044	-0.002
	SymBeta	0.001	0.006	-0.03	-0.01
$\bar{\theta}_1$	Main	0.11	0.54	0.22	0.13
	Hetro	0.53	0.43	-0.37	0.18
	AR1	0.22	0.55	0.24	0.16
	AsyBeta	0.12	0.5	0.23	0.02
	SymBeta	0.17	0.29	0.31	0.33
$\bar{\theta}_2$	Main	0.26	0.54	0.25	0.29
	Hetro	0.87	1.04	-0.52	0.25
	AR1	0.39	1.7	0.22	0.33
	AsyBeta	0.29	2.04	0.45	-0.14
	SymBeta	0.25	1.52	0.33	0.10
$\bar{\theta}_3$	Main	0.25	8.51	0.27	-1.51
	Hetro	2.00	12.08	-0.93	-2.02
	AR1	0.84	10.92	0.14	-1.46
	AsyBeta	0.41	9.39	0.50	-1.11
	SymBeta	0.38	5.01	0.32	-1.04
$\bar{\theta}_{price}$	Main	0.41	1.71	0.28	0.47
	Hetro	0.85	2.16	0.62	0.67
	AR1	0.59	2.39	-0.10	0.33
	AsyBeta	0.51	2.27	0.6	0.37
	SymBeta	0.34	2.48	0.23	0.29

Inclusion of instrumental variables

- Denote $X_{jt} = \{W_{jt}, P_{jt}\}_t$, where price P_{jt} is endogenous:

$$P_{jt} = Z_{jt} \delta + \xi_{jt} \quad \begin{pmatrix} \xi_{jt} \\ \eta_{jt} \end{pmatrix} \sim N \left(\begin{pmatrix} 0 \\ 0 \end{pmatrix}, \Omega \equiv \begin{pmatrix} \Omega_{11} & \Omega_{12} \\ \Omega_{12} & \Omega_{22} \end{pmatrix} \right)$$

- Joint density of price and share at time t is

$$\begin{aligned} \pi(P_t, s_t | \bar{\theta}, r, \delta, \Omega) &= \pi(\xi_t, \eta_t | \bar{\theta}, r, \delta, \Omega) J_{(\xi_t, \eta_t \rightarrow P_t, s_t)} \\ &= \pi(\xi_t, \eta_t | \bar{\theta}, r, \delta, \Omega) \left(J_{(P_t, s_t \rightarrow \xi_t, \eta_t)} \right)^{-1} \end{aligned}$$

where the Jacobian:

$$J_{(P_t, s_t \rightarrow \xi_t, \eta_t)} = \begin{vmatrix} \nabla_{\xi_t} P_t & \nabla_{\eta_t} P_t \\ \nabla_{\xi_t} s_t & \nabla_{\eta_t} s_t \end{vmatrix} = \begin{vmatrix} I & \mathbf{0} \\ \nabla_{\xi_t} s_t & \nabla_{\eta_t} s_t \end{vmatrix} = \left\| \nabla_{\eta_t} s_t \right\| = J_{(s_t \rightarrow \eta_t)}$$

- Two sets of conditionals

$$\begin{aligned} &\bar{\theta}, \delta, \Omega | r, \{s_t, X_t, Z_t\}_{t=1}^T, \bar{\theta}_0, V_{\bar{\theta}}, \bar{\delta}, V_{\delta}, \nu_0, V_{\Omega} \\ &r | \bar{\theta}, \delta, \Omega, \{s_t, X_t, Z_t\}_{t=1}^T, \sigma_{r_diag}^2, \sigma_{r_off}^2 \end{aligned}$$

MSE and bias in IV cell

	MSE		Bias	
	Bayes	GMM	Bayes	GMM
$\bar{\theta}_1$	0.50	9.89	0.49	-0.93
$\bar{\theta}_2$	0.44	13.46	0.51	-1.28
$\bar{\theta}_3$	0.41	34.11	0.41	-2.16
$\bar{\theta}_{price}$	0.28	10	0.33	-0.02
Σ_{11}	3.82	315.49	-1.59	6.86
Σ_{22}	3.11	383.2	-1.51	8.74
Σ_{33}	3.68	6301.31	-1.30	19.09
Σ_{44}	0.75	104.68	-0.06	4.02
Σ_{12}	2.33	117.63	-1.24	1.59
Σ_{13}	1.64	82.45	-1.00	1.20
Σ_{23}	1.92	139.48	0.78	2.65
Σ_{14}	0.36	38.25	-0.25	-1.42
Σ_{24}	0.56	24.03	-0.32	-1.05
Σ_{34}	0.20	24.87	0.10	-1.89
δ_1	0.002	0.002	0.003	0.001
δ_2	0.002	0.002	0.002	0.001
δ_3	0.002	0.002	-0.002	-0.003
δ_4	0.004	0.004	-0.005	-0.002
Ω_{11}	0.0002	0.0002	0.0012	-0.0003
Ω_{12}	0.002	0.003	-0.02	-0.01
Ω_{22}	0.03	1.28	-0.16	0.39
$Corr_{\Omega}$	0.003	0.008	-0.002	-0.062

Summary of robustness checks

Bayes outperforms GMM in all following sampling experiments (cells):

- Basic cells : $\eta_{jt} \sim N(0, \tau^2)$
 - $T=300, \tau=1$ (main)
 - $T=300, \tau=2$
 - $T=300, \tau=0.5$
 - $T=1000, \tau=1$
- Cells with mis-specified η :
 - Heteroskedasticity: $\eta_{jt} \sim N(0, V_{jt} \equiv f(X_{jt}))$
 - AR(1): 0.9
 - Asymmetric Beta[2,5]: log-normal shaped
 - Symmetric Beta[0.5,0.5]: U-shaped
- Instrumental Variable cell:
 - $\text{corr}(\text{price shock}, \text{demand shock})=0.46,$
instruments explain 32% of total price variation

Conclusions

- The impression that aggregate share models are hard to estimate is partly because of the method used, not the intrinsic property of the model
- Bayesian inference is dramatically more efficient than GMM
- Distribution of aggregate shocks
 - There is concern that Bayes makes more distributional assumptions than GMM. We found that it doesn't matter much
 - Straightforward to extend to a Mixture of Normals. Probably requires more data to identify the shape of this distribution