Performance of the BIC_q : Simulation Experiment

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Abstract

A more detailed presentation is given for the simulation experiment reported in Xu and McLeod (2010, Table 2) comparing the performance of the BIC_q with the AIC, BIC and BIC_γ for linear model selection. The scripts displayed in this document may be manually extracted using an editor from the file SimExperimentBICq.Rnw which is located in the subdirectory inst/doc. Alternatively if the R package Sweave is installed the command: R CMD Stangle SimExperimentBICq.Rnw may be used.

Keywords: AIC, extended BIC.

1. Introduction

The performance of the AIC, BIC, BIC_{γ} and BIC_q were compared by simulation. We used the regression, $y_i = \beta_1 x_{i,1} + \beta_2 x_{i,2} + \beta_3 x_{i,3} + \beta_4 x_{i,4} + \beta_5 x_{i,5} + e_i$, $i = 1, \ldots, 40$, where e_i are independent and identically distributed as N(0,1), $x_{i,1} = 1$ and $x_{i,2}, \ldots, x_{i,5}$ are specified in Shao (1993, Table 1).

To compare the models we may consider the model error, $\|X\beta - X(S)\hat{\beta}(S)\|^2$, as well as counts of the number of times a correct model is chosen, a model which underfits is chosen or a model which overfits is chosen.

For the BIC_qwe consider q = 0.15, 0.25, 0.5, and 0.75 as well as $q = \hat{q}_1 \in (q_{1,k_1}, q_{2,k_1})$ and $q = \hat{q}_2 \in (q_{1,k_2}, q_{2,k_2})$. The interval (q_{1,k_1}, q_{2,k_2}) is obtained using the method given in Xu and McLeod (2010) with $\alpha = 0.99$.

2. Main Simulation Function

The basic workhorse for the simulation is our function BICqSimulation shown below.

```
R> `BICqSimulation` <- 
+ function(X, beta, NumSim, g=c(0.5,1), q=c(0.15,0.25,0.75)) 
+ { 
+ ### y=[1,X]*beta+sigma*N(0,1) 
+ #Arguments: X - n*p design matrix for regression 
+ # beta - vector of p+1 coefficients (including intercept) 
+ # NumSim - number of simulations 
+ # g - adjustable parameters for EBIC 
+ # q - tuning parameters for BICq
```

```
+ #
+ stopifnot(ncol(X)==length(beta)-1)
+ n<- nrow(X)
+ K < - ncol(X)
+ b0<- beta[1]
+ b<- beta[-1]
+ s0<- (b!=0)
+ X1<- as.matrix(X[,b!=0])
+ b1<- b[b!=0]
+ y0<- b0+X1%*%b1
+ k0 < - length(b1)
+ out<- 0
+ meMean<- 0
+ meSum2<- 0
+ for (i in 1:NumSim){
      y < -y0 + rnorm(n)
      Xy \leftarrow data.frame(X, y=y)
+ #AIC
      m<- bestglm(Xy,IC="AIC")</pre>
+
      Subs<- m$Subsets[,-1]
      sAIC<- as.matrix(Subs[which.min(Subs[,ncol(Subs)]),][1:K])</pre>
      fit<- m$BestModel
      meAIC<- sum((y0-fit$fitted.values)^2)</pre>
      sAICst \leftarrow as.matrix(Subs[m$Bestq[1,3]+1,][1:K])
      sBICst<- as.matrix(Subs[m$Bestq[2,3]+1,][1:K])
      fit<- lm(y~., data=data.frame(X[,sAICst],y=y))</pre>
      meAICst<- sum((y0-fit$fitted.values)^2)</pre>
      fit<- lm(y~., data=data.frame(X[,sBICst],y=y))</pre>
      meBICst<- sum((y0-fit$fitted.values)^2)</pre>
+ #BIC
      m<- bestglm(Xy,IC="BIC")</pre>
+
      Subs<- m$Subsets[,-1]
      sBIC<- as.matrix(Subs[which.min(Subs[,ncol(Subs)]),][1:K])
      fit<- m$BestModel</pre>
      meBIC<- sum((y0-fit$fitted.values)^2)</pre>
+ #EBICs
      sBICg<- matrix(rep(NA,K*length(g)),ncol=K)</pre>
      meBICg<- rep(NA,length(g))</pre>
      for (j in 1:length(g)){
      m<- bestglm(Xy,IC="BICg",t=g[j])</pre>
      Subs<- m$Subsets[,-1]
      sBICg[j,]<- as.matrix(Subs[which.min(Subs[,ncol(Subs)]),][1:K])</pre>
      fit<- m$BestModel
      meBICg[j]<- sum((y0-fit$fitted.values)^2)</pre>
+ #BICq's
      sBICq<- matrix(rep(NA,K*length(q)),ncol=K)</pre>
```

```
meBICq<- rep(NA,length(q))</pre>
      for (j in 1:length(q)){
      m<- bestglm(Xy,IC="BICq",t=q[j])</pre>
      Subs<- m$Subsets[,-1]
      sBICq[j,]<- as.matrix(Subs[which.min(Subs[,ncol(Subs)]),][1:K])</pre>
      fit<- m$BestModel
      meBICq[j]<- sum((y0-fit$fitted.values)^2)</pre>
      SS<- rbind(sAIC,sBIC,sBICg,sBICq,sAICst,sBICst)
      rownames(SS)<- c("AIC", "BIC", paste("BICg(g=",g,")", sep=""),
                  paste("BICq(q=",q,")",sep=""), "BICq1","BICq2")
      meSS<- c(meAIC,meBIC,meBICg,meBICq,meAICst,meBICst)</pre>
      Delta<- meSS-meMean
      meMean<- meMean+Delta/i
      meSum2<- meSum2+Delta*(meSS-meMean)</pre>
      SS_s0<- SS-matrix(rep(s0,nrow(SS)),nrow=nrow(SS),byrow=TRUE)
      SS_s0<- rowSums(SS_s0)
      kSS<- rowSums(SS)
      overfit <- kSS>k0
      underfit <- kSS<k0
      correct <- as.numeric(SS_s0==0)</pre>
      out <- out+cbind(overfit=(kSS>k0), underfit=(kSS<k0), correct=(SS_s0==0))</pre>
+ overfit <- out[, "overfit"]/NumSim
+ underfit <- out[,"underfit"]/NumSim
+ correct <- out[,"correct"]/NumSim
+ cbind(overfit=overfit, underfit=underfit, correct=correct, me=meMean, se.me= sqrt(meSum2
+ }
```

3. R Script for Simulations

Here we do only 10 simulations to get some timings. It took about 4.8 seconds on our computer, so for 10^5 simulations, it would take about 13.3 hours or about 3.3 hours per model.

The results are in the R list, OutTable, and show for each model the model error as well as counts for number correct, number overfit and number underfit.

```
R> library(bestglm)
Loaded lars 0.9-8
R> data(Shao)
R> NumSim <- 10^1
R> SEED<-123123321</pre>
```

4. Using Rmpi

The main simulations were run on a Mac computer with 8 CPU's using the package **Rmpi** (Yu 2002, 2009). Since 10^5 simulations were needed, we simply divided the job up into four parts, giving each CPU one model to simulate. In this case, essentially all that is needed is to replace apply by mpi.apply in the above R script.

Running the script below required about 3.4 hours. Making use of the multi-core CPU by using the **Rmpi** package, the total time to run all four models is about the same as running just one model on a single CPU. The results are identical to what is obtained if the R script in Section 3 was run on a single CPU NumSim increased to 100,000. This was verified by running the script in Section 3 overnight and setting NumSim to 100000.

4.1. R Script for Simulation Using Rmpi

```
mpi.spawn.Rslaves(nslaves=4)
mpi.bcast.Robj2slave(SEED)
mpi.bcast.Robj2slave(X)
mpi.bcast.Robj2slave(BETA)
mpi.bcast.Robj2slave(NumSim)
mpi.bcast.Robj2slave(BICqSimulation)
mpi.bcast.Robj2slave(GetTable)
mpi.bcast.cmd(library(bestglm))
#note: argument name is 'fun' not 'FUN'
OutTable<-mpi.apply(1:4, fun=GetTable)</pre>
End <- proc.time()[3]</pre>
EndDate<-date()</pre>
Total <- End-Start
#output results
for (i in 1:4)
    write.table(OutTable[[i]], file=paste("tb",i,".dat",sep=""))
TotalTime<-paste("started:",StartDate,"\nended:",EndDate,"\nTotal elapsed time in seconds"
write(TotalTime, file="TotalTime.txt")
#display files at console
dir()
file.show("TotalTime.txt")
#close and quit
mpi.close.Rslaves()
mpi.quit()
```

The script saves the output results in the files tb1.dat, tb2.dat, tb3.dat and tb4.dat. These files may be processed afterwards to produce the tables shown below.

5. Tables

In the case of proportions, the maximum standard error is when p=0.5 and corresponds to $(0.25\times 10^{-5})^{-\frac{1}{2}}\approx 0.0016$.

	overfit	underfit	correct	me	se.me
AIC	0.44	0.00	0.56	3.83	0.01
BIC	0.19	0.00	0.81	3.00	0.01
BICg(g=0.5)	0.16	0.00	0.84	2.91	0.01
BICg(g=1)	0.13	0.00	0.87	2.87	0.01
BICq(q=0.15)	0.03	0.00	0.97	2.25	0.01
BICq(q=0.25)	0.06	0.00	0.94	2.42	0.01
BICq(q=0.75)	0.56	0.00	0.44	4.15	0.01
BICq1	0.04	0.00	0.96	2.31	0.01
$\mathrm{BICq2}$	0.04	0.00	0.96	2.32	0.01

Table 1: 10^5 Simulations with $\beta = (2, 0, 0, 4, 0)$. The percentage of overfit, underfit and correct models as well as the model error, ME, are shown. The maximum sd of the percentages is 0.0016. For ME the sd was about 0.01 in all cases.

	overfit	underfit	correct	me	se.me
AIC	0.32	0.00	0.68	4.23	0.01
BIC	0.13	0.00	0.87	3.68	0.01
BICg(g=0.5)	0.17	0.00	0.83	3.84	0.01
BICg(g=1)	0.23	0.00	0.77	4.08	0.01
BICq(q=0.15)	0.02	0.00	0.98	3.18	0.01
BICq(q=0.25)	0.04	0.00	0.96	3.30	0.01
BICq(q=0.75)	0.43	0.00	0.57	4.44	0.01
BICq1	0.03	0.00	0.97	3.22	0.01
$\mathrm{BICq2}$	0.03	0.00	0.97	3.22	0.01

Table 2: 10^5 Simulations with $\beta = (2, 0, 0, 4, 8)$. The percentage of overfit, underfit and correct models as well as the model error, ME, are shown. The maximum sd of the percentages is 0.0016. For ME the sd was about 0.01 in all cases.

	overfit	underfit	correct	me	se.me
AIC	0.19	0.00	0.81	4.62	0.01
BIC	0.07	0.00	0.93	4.36	0.01
BICg(g=0.5)	0.16	0.00	0.84	4.57	0.01
BICg(g=1)	0.37	0.00	0.63	4.85	0.01
BICq(q=0.15)	0.01	0.00	0.99	4.15	0.01
BICq(q=0.25)	0.02	0.00	0.98	4.18	0.01
BICq(q=0.75)	0.26	0.00	0.74	4.73	0.01
BICq1	0.02	0.00	0.98	4.16	0.01
BICq2	0.02	0.00	0.98	4.16	0.01

Table 3: 10^5 Simulations with $\beta = (2, 9, 0, 4, 8)$. The percentage of overfit, underfit and correct models as well as the model error, ME, are shown. The maximum sd of the percentages is 0.0016. For ME the sd was about 0.01 in all cases.

	overfit	underfit	correct	me	se.me
AIC	0.00	0.00	1.00	5.00	0.01
BIC	0.00	0.00	1.00	5.01	0.01
BICg(g=0.5)	0.00	0.00	1.00	5.01	0.01
BICg(g=1)	0.00	0.00	1.00	5.00	0.01
BICq(q=0.15)	0.00	0.01	0.99	5.18	0.01
BICq(q=0.25)	0.00	0.00	1.00	5.08	0.01
BICq(q=0.75)	0.00	0.00	1.00	5.00	0.01
BICq1	0.00	0.01	0.99	5.13	0.01
$\mathrm{BICq2}$	0.00	0.01	0.99	5.13	0.01

Table 4: 10^5 Simulations with $\beta = (2, 9, 6, 4, 8)$. The percentage of overfit, underfit and correct models as well as the model error, ME, are shown. The maximum sd of the percentages is 0.0016. For ME the sd was about 0.01 in all cases.

6. Comments on the Simulation Results

The results for the mean error, ME, are identical to those reported in Xu and McLeod (2010, Table 2). It is interesting to note that in the only case where the BICq1 or BICq2 did not do better than all the other model selection criteria was with the full model $\beta = (2, 9, 6, 4, 8)$ and in this case the difference was very small and was apparently due to a very slight tendency to underfit. With the other models, it was interesting to see that most of the criteria tended to overfit the model. Underfitting was very rare. This agrees with asymptotic theory. Even in the case of the AIC, the results of Shibata (1980) indicated that asymptotically the probability of underfitting was zero.

Our simulation results are also available in the files tb1.dat, tb2.dat, tb3.dat and tb4.dat included with this vignette.

References

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