

AUGMENTED INVERSE PROBABILITY WEIGHTING

The propensity score for binary exposure, denoted $e(X)$,

$$e(x) = f_{Z|X}^{\circ}(1|x) = \Pr_{Z|X}^{\circ}[Z = 1|X = x]$$

can be used to construct generalized versions of the inverse probability weighted (IPW) estimators based on augmentation via the conditional mean model $\mu(x, z)$

$$\mathbb{E}_{Y|X,Z}^{\circ}[Y|X = x, Z = z] = \mu(x, z)$$

We deduce that augmented IPW (AIPW) estimators are

$$\begin{aligned}\tilde{\mu}_{\text{AIPW}}(0) &= \frac{1}{n} \sum_{i=1}^n \frac{(1 - Z_i)(Y_i - \mu(X_i, 0))}{1 - e(X_i)} + \frac{1}{n} \sum_{i=1}^n \mu(X_i, 0) \\ \tilde{\mu}_{\text{AIPW}}(1) &= \frac{1}{n} \sum_{i=1}^n \frac{Z_i(Y_i - \mu(X_i, 1))}{e(X_i)} + \frac{1}{n} \sum_{i=1}^n \mu(X_i, 1)\end{aligned}$$

or, in standardized weight form

$$\begin{aligned}\hat{\mu}_{\text{AIPW}}(0) &= \sum_{i=1}^n W_{i0}(Y_i - \mu(X_i, 0)) + \frac{1}{n} \sum_{i=1}^n \mu(X_i, 0) \\ \hat{\mu}_{\text{AIPW}}(1) &= \sum_{i=1}^n W_{i1}(Y_i - \mu(X_i, 1)) + \frac{1}{n} \sum_{i=1}^n \mu(X_i, 1)\end{aligned}$$

with

$$W_{i0} = \frac{\frac{(1 - Z_i)}{(1 - e(X_i))}}{\sum_{j=1}^n \frac{(1 - Z_j)}{(1 - e(X_j))}} \quad W_{i1} = \frac{\frac{Z_i}{e(X_i)}}{\sum_{j=1}^n \frac{Z_j}{e(X_j)}}$$

These estimators are generalizations of IPW estimators, and are more robust to misspecification: the estimators are consistent provided **at least one** of the two models $e(x)$ or $\mu(x, z)$ is correctly specified. That is, estimators that use instead the models

$$g(x) \quad \text{and} \quad m(x, z)$$

for the propensity score and conditional mean model respectively are still consistent provided either $g(x) \equiv e(x)$ or $m(x, z) \equiv \mu(x, z)$. If **both** are correctly specified, the estimators are the optimal IPW estimators.

Simulation study: In the following simulation, we have $k = 10$ predictors, but only the first three are confounders. We compare the results for the following models:

Model 0: Outcome regression under correct specification of $\mu(x, z)$

Model 1: IPW with correct specification of $e(x)$

Model 2: AIPW with correct specification of both $e(x)$ and $\mu(x, z)$

Model 3: AIPW with correct specification of $e(x)$ only

Model 4: AIPW with correct specification of $\mu(x, z)$ only

Model 5: AIPW with both models mis-specified

We compute both the original (denoted a) and standardized weight (denoted b) estimators. We perform 2000 replicate analyses with $n = 1000$. In the first analysis, we assume that the propensity score is known.

```

#Monte Carlo study
library(mvnfast)
set.seed(23987)
n<-1000
k<-10
Mu<-sample(-5:5,size=k,rep=T)/2
Sigma<-0.1*0.8^abs(outer(1:k,1:k,'-'))
al<-c(-4,rep(1,6),rep(-2,4))
expit<-function(x){return(1/(1+exp(-x)))};logit<-function(x){return(log(x)-log(1-x))}
be<-rep(0,k+2)
be[1:5]<-c(2,1,-2.0,2.2,3.6)
sig<-2
nreps<-2000
ests<-matrix(0,nrow=nreps,ncol=11)
for(irep in 1:nreps){

  X<-rmvn(n,mu=Mu,Sigma)
  Xal<-cbind(1,X)
  ps.true<-expit(Xal %*% al)
  Z<-rbinom(n,1,ps.true)
  Xm<-cbind(1,Z,X)
  muval<-as.vector(Xm %*% be)

  Y<-rnorm(n,muval,sig)
  eX<-ps.true
  w<-(1-Z)/(1-eX)+Z/eX
  w0<-(1-Z)/(1-eX)
  W0<-w0/sum(w0)
  w1<-Z/eX
  W1<-w1/sum(w1)

  mu0<- cbind(1,0,X) %*% be
  mu1<- cbind(1,1,X) %*% be

  #Intercept only mis-specified model
  m0<- cbind(1,0,0*X) %*% be
  m1<- cbind(1,1,0*X) %*% be

  ests[irep,1]<-mean(mu1)-mean(mu0)
  ests[irep,2]<-mean(w1*Y)-mean(w0*Y)
  ests[irep,3]<-sum(W1*Y)-sum(W0*Y)
  ests[irep,4]<-mean(w1*(Y-mu1))+mean(mu1)-mean(w0*(Y-mu0))-mean(mu0)
  ests[irep,5]<-sum(W1*(Y-mu1))+mean(mu1)-sum(W0*(Y-mu0))-mean(mu0)
  ests[irep,6]<-mean(w1*(Y-m1))+mean(m1)-mean(w0*(Y-m0))-mean(m0)
  ests[irep,7]<-sum(W1*(Y-m1))+mean(m1)-sum(W0*(Y-m0))-mean(m0)

  #Mis-specified PS model using X1 only.
  #Obtain intercept from data
  gX<-expit(logit(mean(Z))+al[2]*X[,1])
  wg0<-(1-Z)/(1-gX)
  Wg0<-wg0/sum(wg0)
  wg1<-Z/gX
  Wg1<-wg1/sum(wg1)

  ests[irep,8]<-mean(wg1*(Y-mu1))+mean(mu1)-mean(wg0*(Y-mu0))-mean(mu0)
  ests[irep,9]<-sum(Wg1*(Y-mu1))+mean(mu1)-sum(Wg0*(Y-mu0))-mean(mu0)

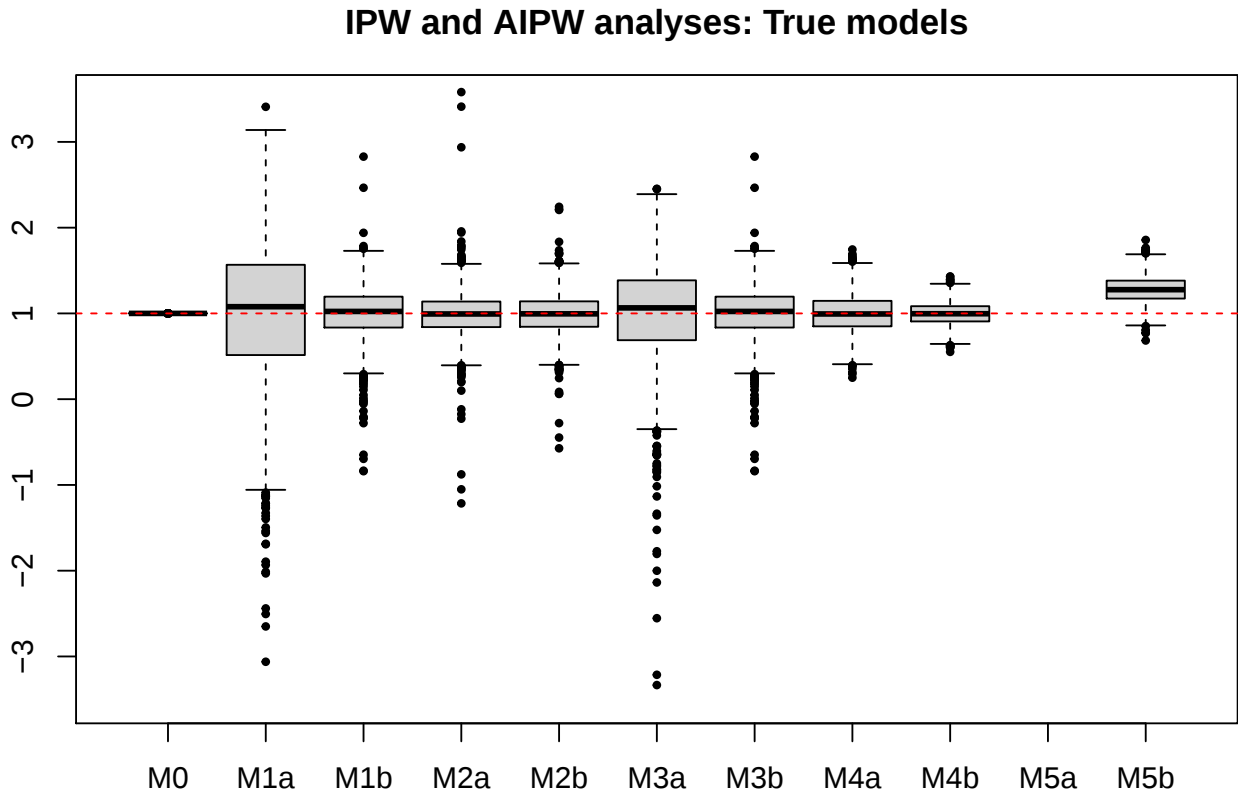
  ests[irep,10]<-mean(wg1*(Y-m1))+mean(m1)-mean(wg0*(Y-m0))-mean(m0)
  ests[irep,11]<-sum(Wg1*(Y-m1))+mean(m1)-sum(Wg0*(Y-m0))-mean(m0)
}

```

```

par(mar=c(4,4,3,0))
nvec<-as.character(c(0,rep(1:5,each=2)))
nvec<-paste('M',nvec,sep='')
nvec[c(2,4,6,8,10)]<-paste(nvec[c(2,4,6,8,10)], 'a',sep='')
nvec[c(3,5,7,9,11)]<-paste(nvec[c(3,5,7,9,11)], 'b',sep='')
boxplot(ests,ylim=range(-3.5,3.5),pch=19,cex=0.5, names=nvec)
title('IPW and AIPW analyses: True models')
abline(h=1,lty=2,col='red')

```



```

#Bias
b<-sqrt(n)*apply(ests-1,2,mean)
#Variance
v<-n*apply(ests,2,var)
#MSE
mse<-b^2+v
res<-rbind(b,v,mse)
colnames(res)<-nvec
rownames(res)<-c('Bias','Var.','MSE')
round(res,2)

```

	M0	M1a	M1b	M2a	M2b	M3a	M3b	M4a	M4b	M5a	M5b
Bias	0	0.15	0.01	-0.30	-0.30	-0.04	0.01	-0.10	-0.14	231.94	8.80
Var.	0	779.64	86.73	70.00	56.71	384.10	86.73	47.20	16.99	84.50	24.18
MSE	0	779.66	86.73	70.09	56.80	384.10	86.73	47.21	17.01	53881.25	101.63

The results are as expected in light of the theory.

In a second analysis, we now **estimate** the propensity score model.

```

#Monte Carlo study
set.seed(23987)
nreps<-2000
ests2<-matrix(0,nrow=nreps,ncol=11)
for(irep in 1:nreps){

  X<-rmvn(n,mu=Mu,Sigma)
  Xal<-cbind(1,X)
  ps.true<-expit(Xal %*% al)
  Z<-rbinom(n,1,ps.true)
  Xm<-cbind(1,Z,X)
  muval<-as.vector(Xm %*% be)

  Y<-rnorm(n,muval,sig)
  ps.fitted<-fitted(glm(Z~X,family=binomial))
  eX<-ps.fitted
  w<-(1-Z)/(1-eX)+Z/eX
  w0<-(1-Z)/(1-eX)
  W0<-w0/sum(w0)
  w1<-Z/eX
  W1<-w1/sum(w1)

  mu0<- cbind(1,0,X) %*% be
  mu1<- cbind(1,1,X) %*% be

  #Intercept only mis-specified model
  m0<- cbind(1,0,0*X) %*% be
  m1<- cbind(1,1,0*X) %*% be

  ests2[irep,1]<-mean(mu1)-mean(mu0)
  ests2[irep,2]<-mean(w1*Y)-mean(w0*Y)
  ests2[irep,3]<-sum(W1*Y)-sum(W0*Y)
  ests2[irep,4]<-mean(w1*(Y-mu1))+mean(mu1)-mean(w0*(Y-mu0))-mean(mu0)
  ests2[irep,5]<-sum(W1*(Y-mu1))+mean(mu1)-sum(W0*(Y-mu0))-mean(mu0)
  ests2[irep,6]<-mean(w1*(Y-m1))+mean(m1)-mean(w0*(Y-m0))-mean(m0)
  ests2[irep,7]<-sum(W1*(Y-m1))+mean(m1)-sum(W0*(Y-m0))-mean(m0)

  #Mis-specified PS model using X1 only.
  gX<-fitted(glm(Z~X[,1]),family=binomial)
  wg0<-(1-Z)/(1-gX)
  Wg0<-wg0/sum(wg0)
  wg1<-Z/gX
  Wg1<-wg1/sum(wg1)

  ests2[irep,8]<-mean(wg1*(Y-mu1))+mean(mu1)-mean(wg0*(Y-mu0))-mean(mu0)
  ests2[irep,9]<-sum(Wg1*(Y-mu1))+mean(mu1)-sum(Wg0*(Y-mu0))-mean(mu0)

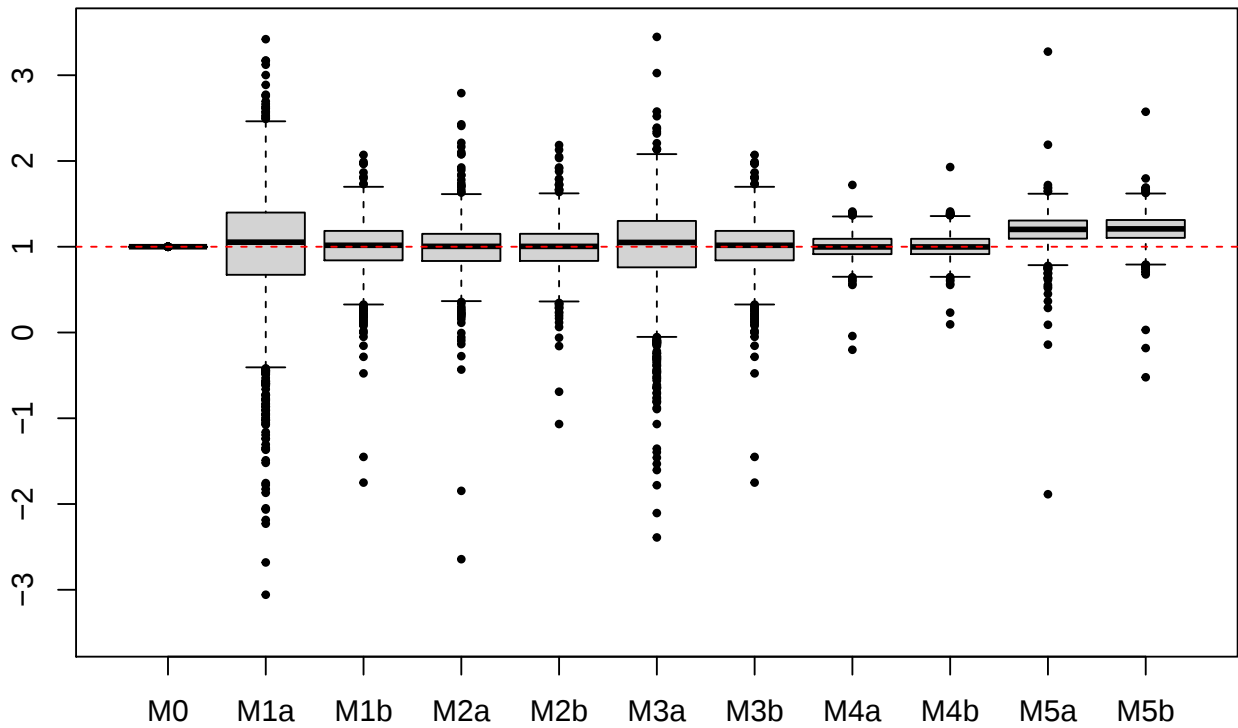
  ests2[irep,10]<-mean(wg1*(Y-m1))+mean(m1)-mean(wg0*(Y-m0))-mean(m0)
  ests2[irep,11]<-sum(Wg1*(Y-m1))+mean(m1)-sum(Wg0*(Y-m0))-mean(m0)

}

par(mar=c(4,4,3,0))
nvec<-as.character(c(0,rep(1:5,each=2)))
nvec<-paste('M',nvec,sep='')
nvec[c(2,4,6,8,10)]<-paste(nvec[c(2,4,6,8,10)], 'a', sep='')
nvec[c(3,5,7,9,11)]<-paste(nvec[c(3,5,7,9,11)], 'b', sep='')
boxplot(ests2,ylim=range(-3.5,3.5),pch=19,cex=0.5,names=nvec)
title('IPW and AIPW analyses: estimated PS model')
abline(h=1,lty=2,col='red')

```

IPW and AIPW analyses: estimated PS model



```
#Bias
b<-sqrt(n)*apply(ests2-1,2,mean)
#Variance
v<-n*apply(ests2,2,var)
#MSE
mse<-b^2+v
res2<-rbind(b,v,mse)
colnames(res2)<-nvec
rownames(res2)<-c('Bias','Var.','MSE')
round(res2,2)
```

	M0	M1a	M1b	M2a	M2b	M3a	M3b	M4a	M4b	M5a	M5b
+ Bias	0	-0.41	0.15	-0.15	-0.14	-0.28	0.15	0.00	0.02	6.19	6.56
+ Var.	0	571.46	84.49	80.33	65.25	303.00	84.49	19.42	18.93	55.76	28.30
+ MSE	0	571.63	84.51	80.35	65.27	303.08	84.51	19.42	18.93	94.01	71.31

The variances are decreased after estimation

In a third analysis, we now estimate the propensity score **and** the conditional mean models.

```

#Monte Carlo study
set.seed(23987)
nreps<-2000
ests3<-matrix(0,nrow=nreps,ncol=11)
for(irep in 1:nreps){

  X<-rmvn(n,mu=Mu,Sigma)
  Xal<-cbind(1,X)
  ps.true<-expit(Xal %*% al)
  Z<-rbinom(n,1,ps.true)
  Xm<-cbind(1,Z,X)
  muval<-as.vector(Xm %*% be)

  Y<-rnorm(n,muval,sig)
  ps.fitted<-fitted(glm(Z~X,family=binomial))
  eX<-ps.fitted
  w<-(1-Z)/(1-eX)+Z/eX
  w0<-(1-Z)/(1-eX)
  W0<-w0/sum(w0)
  w1<-Z/eX
  W1<-w1/sum(w1)

  fit.mu<-lm(Y~Z+X)
  mu0<- predict(fit.mu,newdata=data.frame(Z=0,X=X))
  mu1<- predict(fit.mu,newdata=data.frame(Z=1,X=X))

  #Intercept only mis-specified model
  X1<-X[,1]
  fit.m<-lm(Y~Z+X1)
  m0<- predict(fit.m,newdata=data.frame(Z=0,X1=X[,1]))
  m1<- predict(fit.m,newdata=data.frame(Z=1,X1=X[,1]))

  ests3[irep,1]<-mean(mu1)-mean(mu0)
  ests3[irep,2]<-mean(w1*Y)-mean(w0*Y)
  ests3[irep,3]<-sum(W1*Y)-sum(W0*Y)
  ests3[irep,4]<-mean(w1*(Y-mu1))+mean(mu1)-mean(w0*(Y-mu0))-mean(mu0)
  ests3[irep,5]<-sum(W1*(Y-mu1))+mean(mu1)-sum(W0*(Y-mu0))-mean(mu0)
  ests3[irep,6]<-mean(w1*(Y-m1))+mean(m1)-mean(w0*(Y-m0))-mean(m0)
  ests3[irep,7]<-sum(W1*(Y-m1))+mean(m1)-sum(W0*(Y-m0))-mean(m0)

  #Mis-specified PS model using X1 only.
  gX<-fitted(glm(Z~X[,1]),family=binomial)
  wg0<-(1-Z)/(1-gX)
  Wg0<-wg0/sum(wg0)
  wg1<-Z/gX
  Wg1<-wg1/sum(wg1)

  ests3[irep,8]<-mean(wg1*(Y-mu1))+mean(mu1)-mean(wg0*(Y-mu0))-mean(mu0)
  ests3[irep,9]<-sum(Wg1*(Y-mu1))+mean(mu1)-sum(Wg0*(Y-mu0))-mean(mu0)

  ests3[irep,10]<-mean(wg1*(Y-m1))+mean(m1)-mean(wg0*(Y-m0))-mean(m0)
  ests3[irep,11]<-sum(Wg1*(Y-m1))+mean(m1)-sum(Wg0*(Y-m0))-mean(m0)

}

par(mar=c(4,4,3,0))
nvec<-as.character(c(0,rep(1:5,each=2)))
nvec<-paste('M',nvec,sep='')
nvec[c(2,4,6,8,10)]<-paste(nvec[c(2,4,6,8,10)], 'a',sep='')
nvec[c(3,5,7,9,11)]<-paste(nvec[c(3,5,7,9,11)], 'b',sep='')

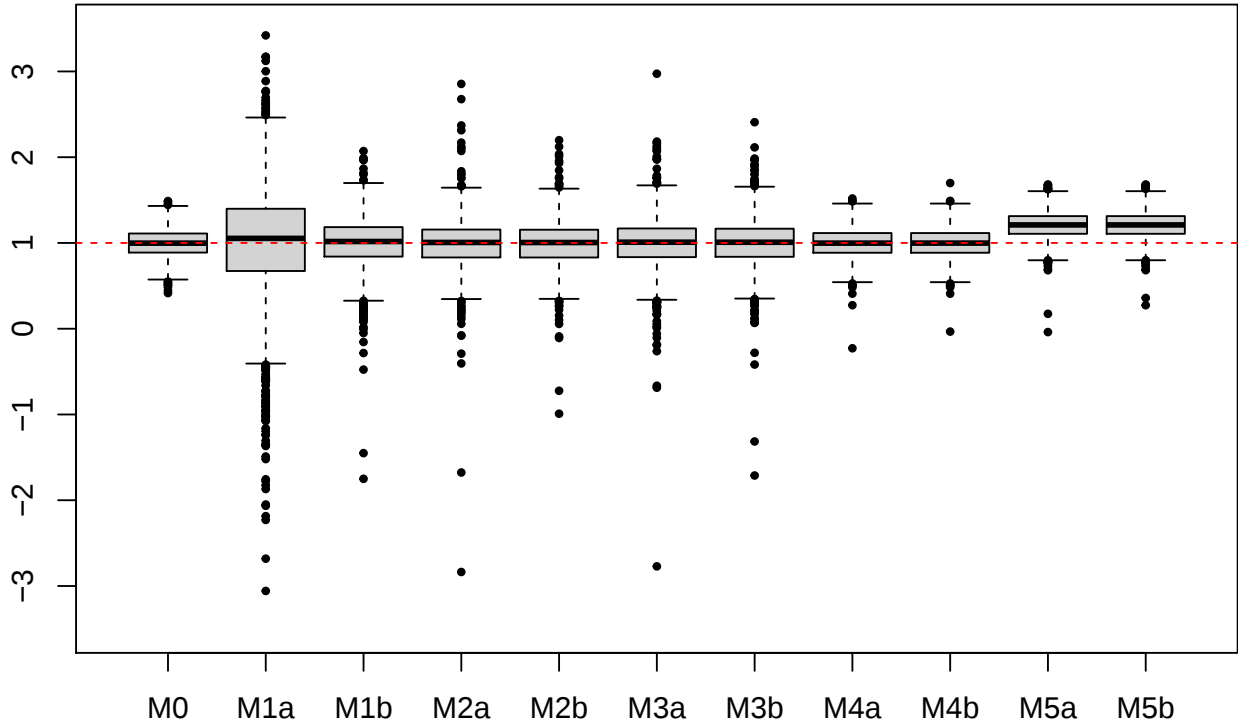
```

```

boxplot(ests3,ylim=range(-3.5,3.5),pch=19,cex=0.5,names=nvec)
title('IPW and AIPW analyses: Both models estimated')
abline(h=1,lty=2,col='red')

```

IPW and AIPW analyses: Both models estimated



```

#Bias
b<-sqrt(n)*apply(ests3-1,2,mean)
#Variance
v<-n*apply(ests3,2,var)
#MSE
mse<-b^2+v
res3<-rbind(b,v,mse)
colnames(res3)<-nvec
rownames(res3)<-c('Bias','Var.','MSE')
round(res3,2)

```

	M0	M1a	M1b	M2a	M2b	M3a	M3b	M4a	M4b	M5a	M5b
+ Bias	-0.01	-0.41	0.15	-0.16	-0.14	-0.13	-0.01	-0.03	-0.02	6.70	6.71
+ Var.	25.98	571.46	84.49	80.93	65.24	104.90	79.83	27.30	27.02	25.39	24.81
+ MSE	25.98	571.63	84.51	80.95	65.26	104.92	79.83	27.30	27.02	70.30	69.82