

# Gorilla Faceshape (Male vs. Female) Bayesian Procrustes:

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## Load the Landmark Data of Gorilla

```
#install.packages("githubinstall")
#require(githubinstall)
#githubinstall("BPviGM1")

#install.packages("devtools")
#require(devtools)
#install_github("debashischatterjee111/BPviGM1")

require(shapes)

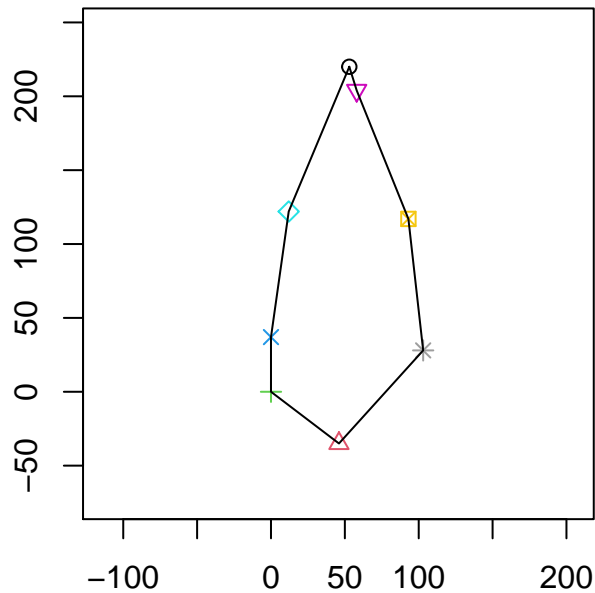
## Loading required package: shapes
require(Morpho)

## Loading required package: Morpho
require(BPviGM1)

## Loading required package: BPviGM1
data(gorf.dat)
data(gorm.dat)

ogf=gf=gorf.dat
ogm=gf=gorm.dat

plotshapes(gm[, , 1], joinline=c(1,6,7,8,2,3,4,5,1), col=1:8, symbol=1:8)
```



```
#polygon(gm[,1,1],gm[,2,1])
```

## Check for proper dimension of 3D array

```
dim(ogm)
```

```
## [1] 8 2 29
```

## Sort according to Landmark order

```
sortcoldat=function(dat,seq)
{dat[,]=dat[seq,,]
  return(dat)
}
```

```
o.gf=ogf=sortcoldat(ogf,c(1,6,7,8,2,3,4,5))
o.gm=ogm=sortcoldat(ogm,c(1,6,7,8,2,3,4,5))
```

## Keep the data in Pre-shape space

```
for(k in 1:dim(ogf)[3])
{
  o.gf[, ,k] <- rotonto(ogf[, ,1],ogf[, ,k],scale=T)$yrot
```

```

}

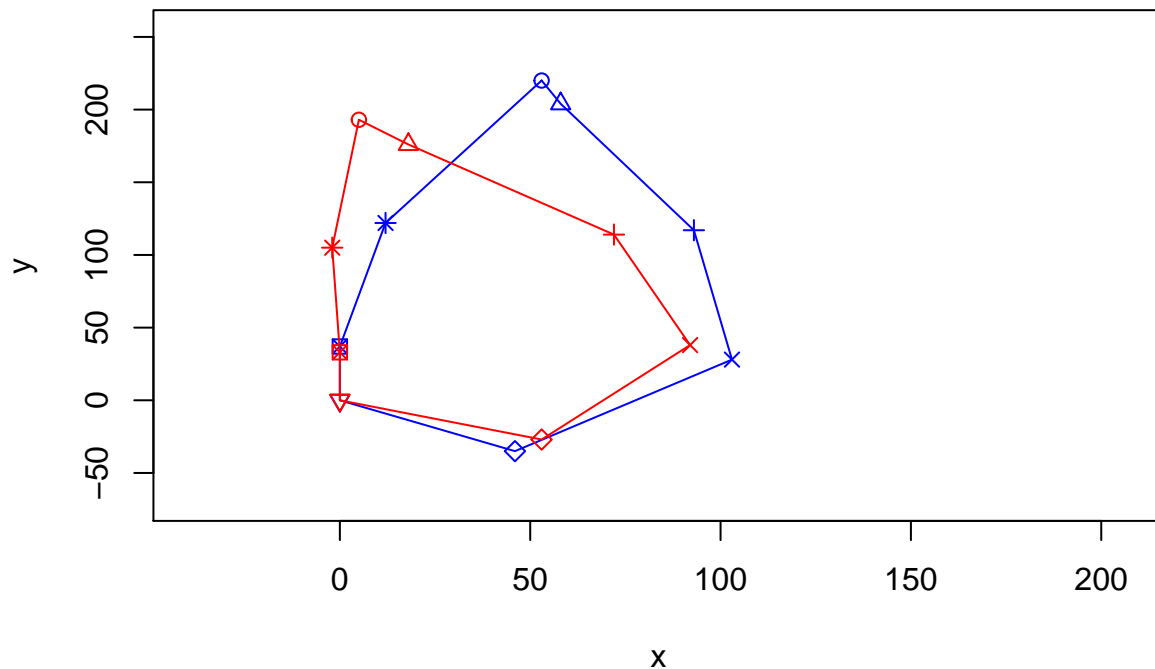
for(k in 1:dim(ogm)[3])
{
  o.gm[, ,k] <- rotonto(ogm[, ,1],ogm[, ,k],scale=T)$yrot
}

plot(c(min(o.gm[,1,])-30, 100+max(o.gm[,1,])), c(min(o.gm[,2,])-30, max(o.gm[,2,])+30), type = "n", main = "Raw Data-plot of Apes: male & female")
#plotshapes(gm[, ,1],joinline=c(1,6,7,8,2,3,4,5,1), col=1:8, symbol=1:8)

polygon(o.gm[,1,1],o.gm[,2,1],density=0, col="skyblue", border="blue")
points(o.gm[,1,1],o.gm[,2,1], col="blue",pch=1:8)
polygon(o.gf[,1,1],o.gf[,2,1],density=0, col="pink", border="red")
points(o.gf[,1,1],o.gf[,2,1], col="red",pch=1:8)

```

## Raw Data-plot of Apes: male & female



Load the novel R package BPviGM1

Start & Run MCMC for Gorilla Male upto 10 objects

```

require(BPviGM1)
apem10000.10=MCMCpostPsample2D(1.5,
rep(0.5,1),o.gm[, ,1:10],10000)

```

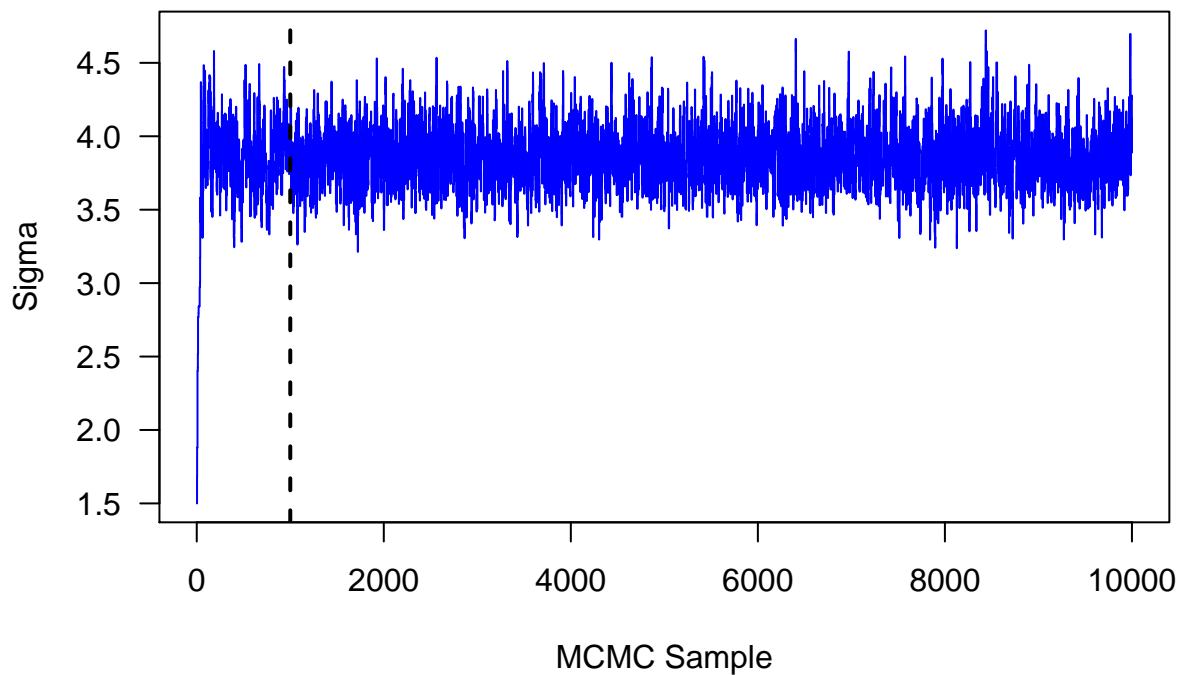
```

##      |                                     | 0%current sample:[1] mcm
## : Done :[1] "***"

```

```
## [1] "Time taken:"
##      user  system elapsed
## 232.47    0.47   243.88
PLOTpostvar2D(apem10000.10, 1000,colu="blue")
```

### (MCMC plot from Posterior of parameters



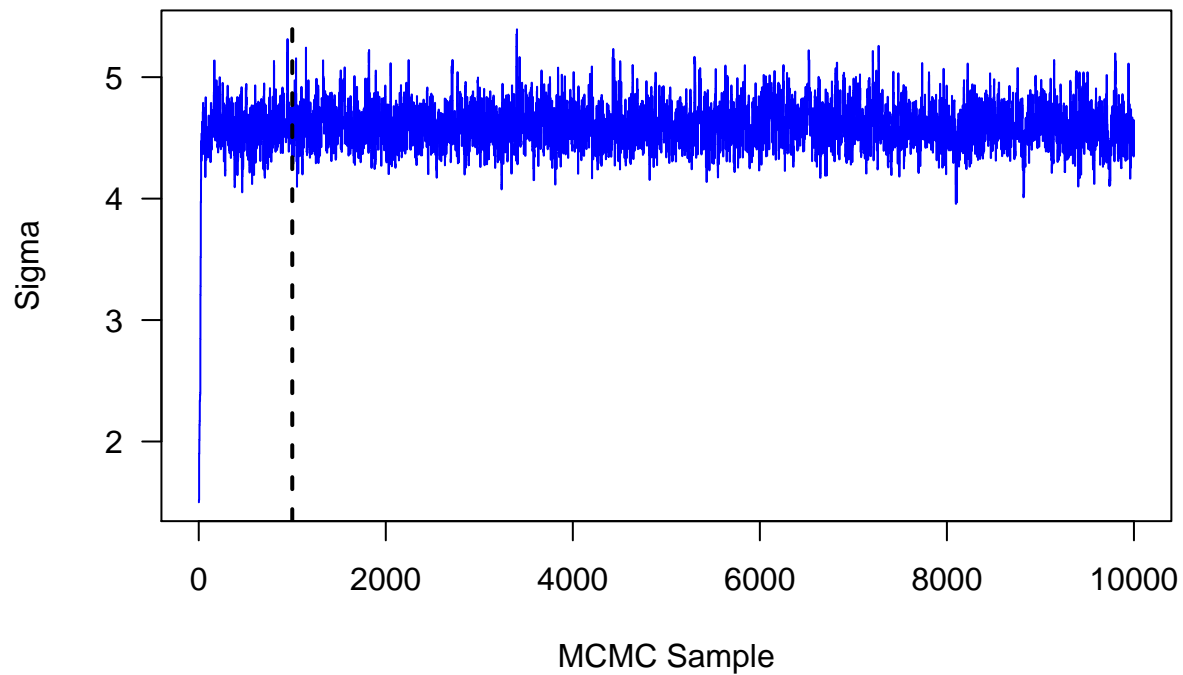
### Start & Run MCMC for Gorilla Male upto 20 objects

```
apem10000.20=MCMCpostPsample2D(1.5,
rep(0.5,1),o.gm[,1:20],10000)
```

```
##      |                                     | 0%current sample:[1] mcm
## : Done :[1] "****"
## [1] "Time taken:"
##      user  system elapsed
## 449.72    0.89   469.03
```

```
PLOTpostvar2D(apem10000.20, 1000,colu="blue")
```

### (MCMC plot from Posterior of parameters



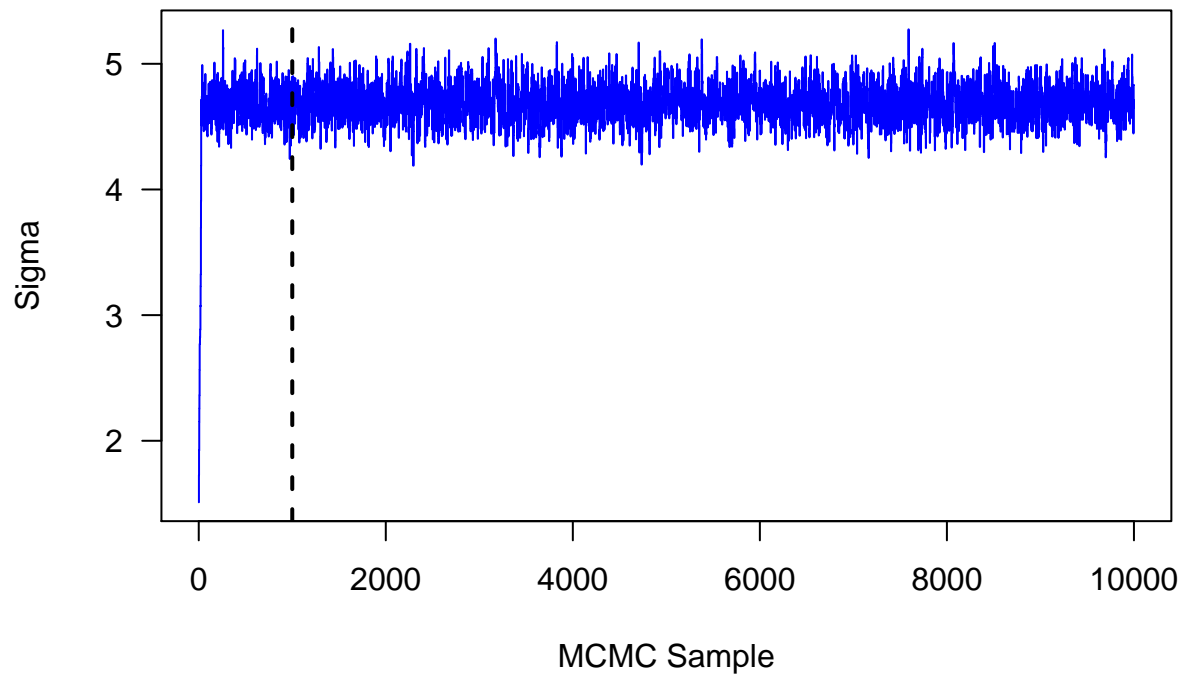
### Start & Run MCMC for Gorilla Male upto 30 objects

```
apem10000.30=MCMCpostPsample2D(1.5,  
rep(0.5,1),o.gm[,],10000)
```

```
##      |                                     |      0%current sample:[1] mcm  
## : Done :[1] "***"  
## [1] "Time taken:"  
##      user  system elapsed  
## 633.05    1.14   651.22
```

```
PLOTpostvar2D(apem10000.30, 1000,colu="blue")
```

### (MCMC plot from Posterior of parameters



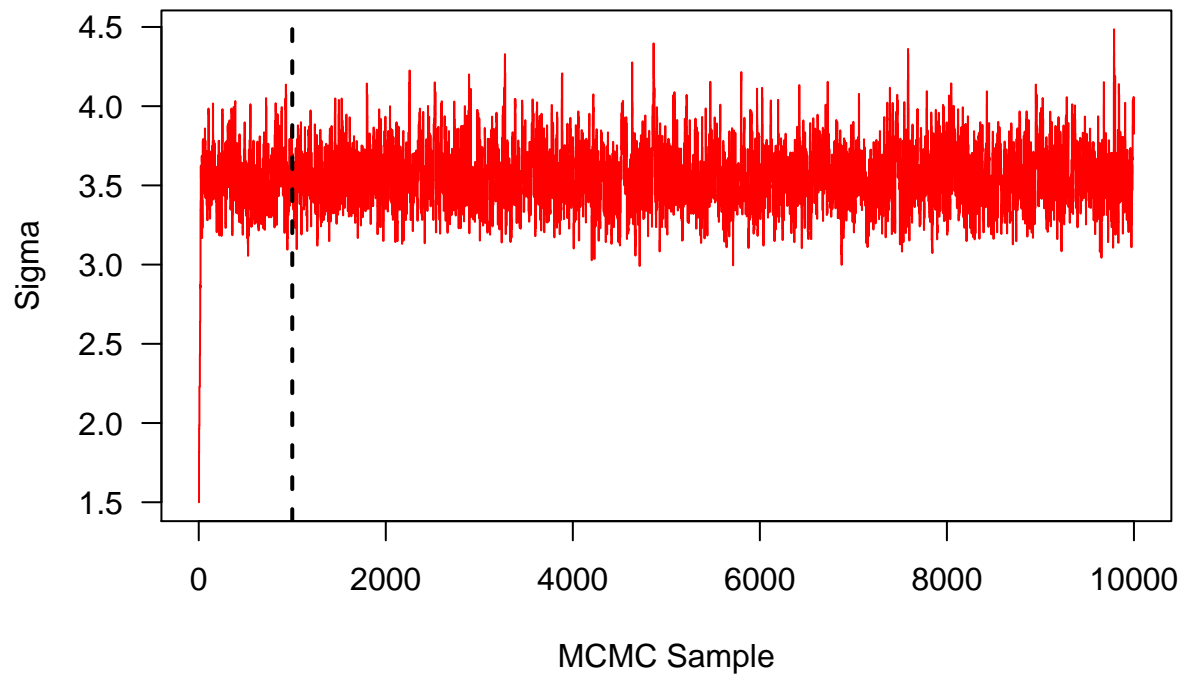
### Start & Run MCMC for Gorilla Female upto 10 objects

```
apec10000.10=MCMCpostPsample2D(1.5,  
rep(0.5,1),o.gf[, ,1:10],10000)
```

```
##      |                                     |      0%current sample:[1] mcm  
## : Done :[1] "***"  
## [1] "Time taken:"  
##      user  system elapsed  
## 221.22    0.44   228.61
```

```
PLOTpostvar2D(apec10000.10, 1000,colu="red")
```

### (MCMC plot from Posterior of parameters)



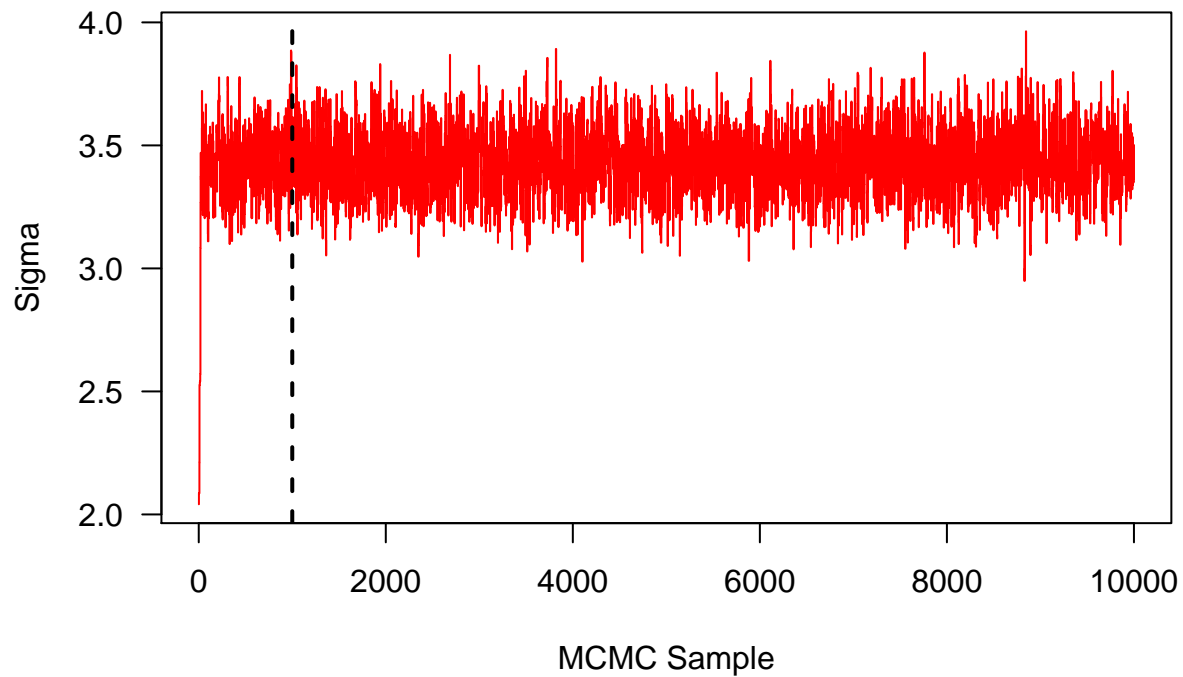
### Start & Run MCMC for Gorilla Female upto 20 objects

```
apef10000.20=MCMCpostPsample2D(1.5,  
rep(0.5,1),o.gf[, ,1:20],10000)
```

```
##      |                                     |      0%current sample:[1] mcm  
## : Done :[1] "***"  
## [1] "Time taken:"  
##      user  system elapsed  
## 447.34    1.00  465.04
```

```
PPL0Tpostvar2D(apef10000.20, 1000,colu="red")
```

### (MCMC plot from Posterior of parameters



### Start & Run MCMC for Gorilla Female upto 30 objects

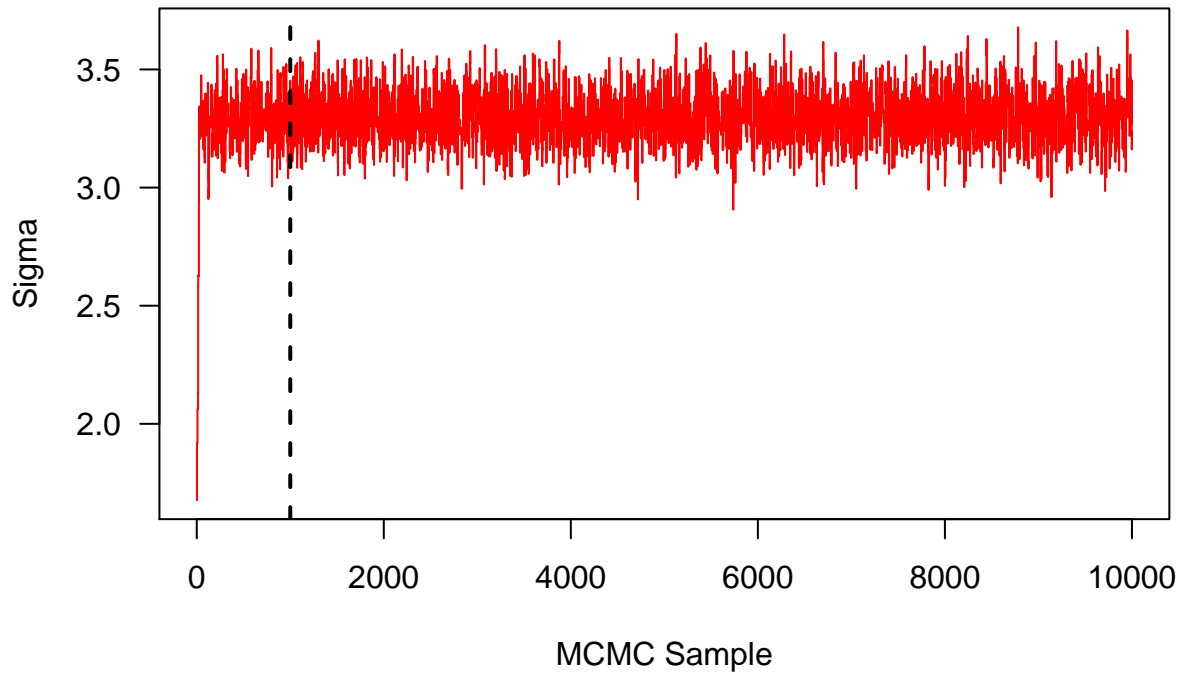
```
apecf10000.30=MCMCpostPsample2D(1.5,  
rep(0.5,1),o.gf,10000)
```

```
##      |                                     |      0%current sample:[1] mcm  
## : Done :[1] "***"  
## [1] "Time taken:"  
##      user  system elapsed  
## 651.87    1.19  663.73
```

```
PLOTpostvar2D(apecf10000.30, 1000,colu="red")
```



## (MCMC plot from Posterior of parameters

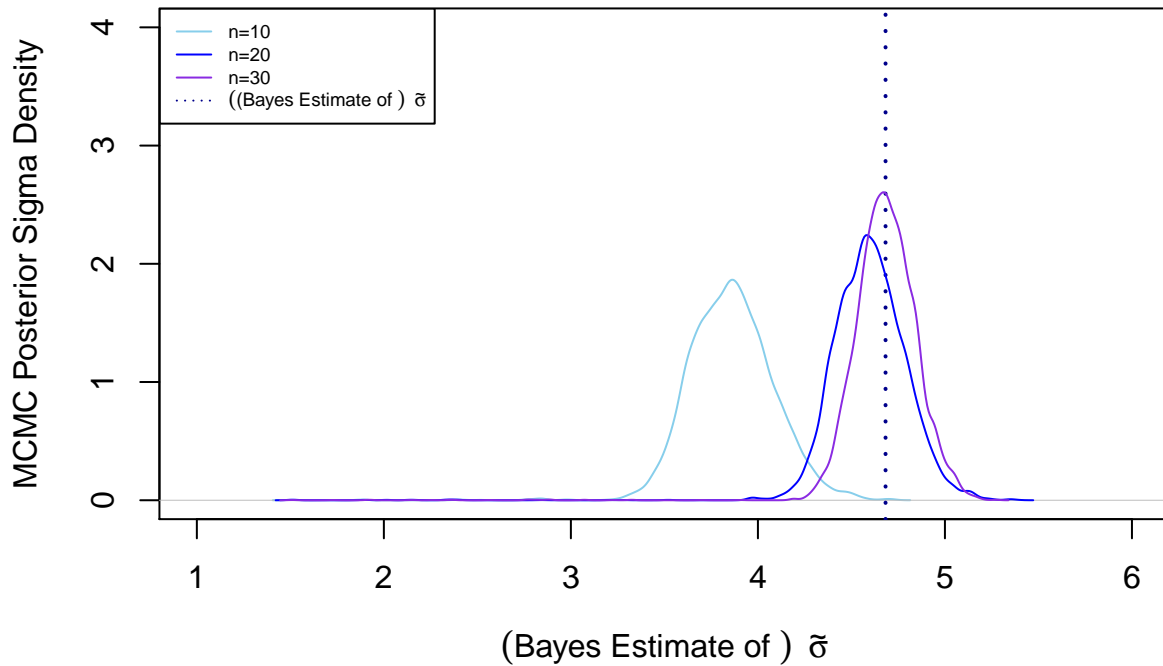


## Plot for Posterior Convergence of Gorilla Male

```
#plot(density(ress_10[1001:10000,2]), xlim=c(0,8),ylim=c(0,5),col="red",ylab="density (with vague prior)
plot(density(apem10000.10), xlim=c(1,6),ylim=c(0,4),col="skyblue", xlab=expression(("Bayes Estimate of 
#abline(v=theta, col="black",lwd=2, lty=1)
abline(v=mean(apem10000.30), col="darkblue",lwd=2, lty=3)
#lines(density(apem10000.20), ylim=c(0,12),col="blue")
lines(density(apem10000.20), ylim=c(0,12),col="blue")
lines(density(apem10000.30), ylim=c(0,12),col="blueviolet")

legend("topleft",cex=0.6, c("n=10","n=20","n=30", expression(("Bayes Estimate of ")~tilde(sigma))), 1
```

### MCMC posterior of Gorilla (male) $\tilde{\sigma}$ (+ uniform prior)

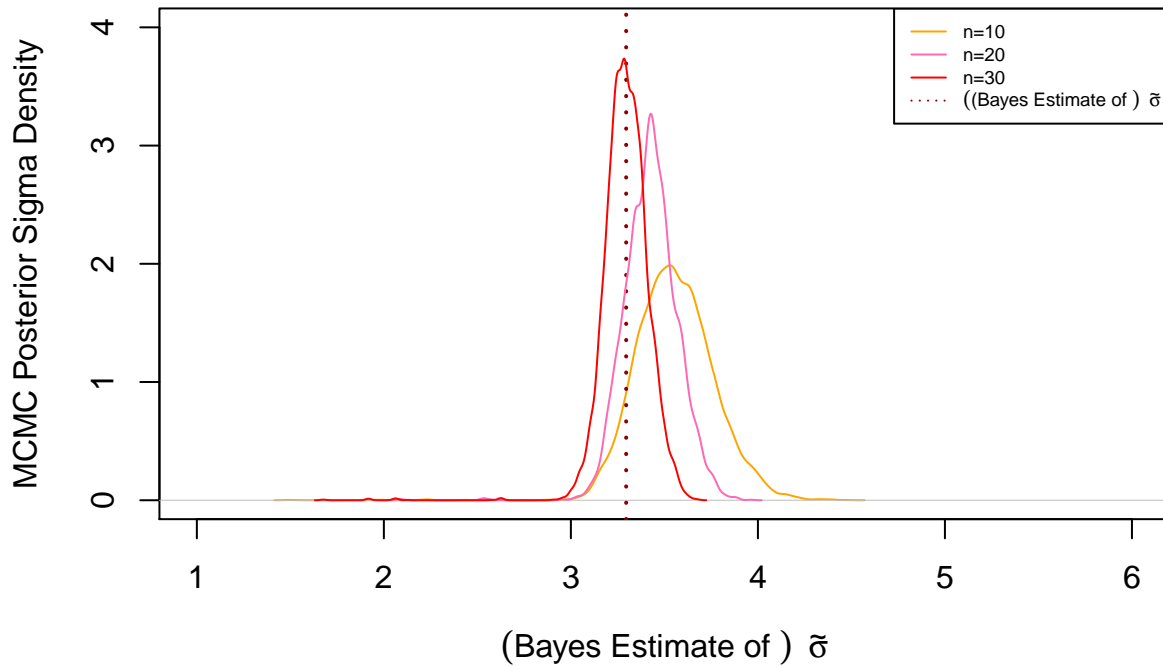


### Plot for Posterior Convergence of Gorilla Female

```
#plot(density(ress_10[1001:10000,2]), xlim=c(0,8),ylim=c(0,5),col="red",ylab="density (with vague prior)
plot(density(apef10000.10), xlim=c(1,6),ylim=c(0,4),col="orange", xlab=expression(("Bayes Estimate of "
#abline(v=theta, col="black",lwd=2, lty=1)
abline(v=mean(apef10000.30), col="red4",lwd=2, lty=3)
#lines(density(apem10000.20), ylim=c(0,12),col="blue")
lines(density(apef10000.20), ylim=c(0,12),col="hotpink")
lines(density(apef10000.30), ylim=c(0,12),col="red")

legend("topright",cex=0.6, c("n=10","n=20","n=30", expression(("(Bayes Estimate of ")~tilde(sigma))),
```

### MCMC posterior of Gorilla (female) $\sigma$ (+ uniform prior)



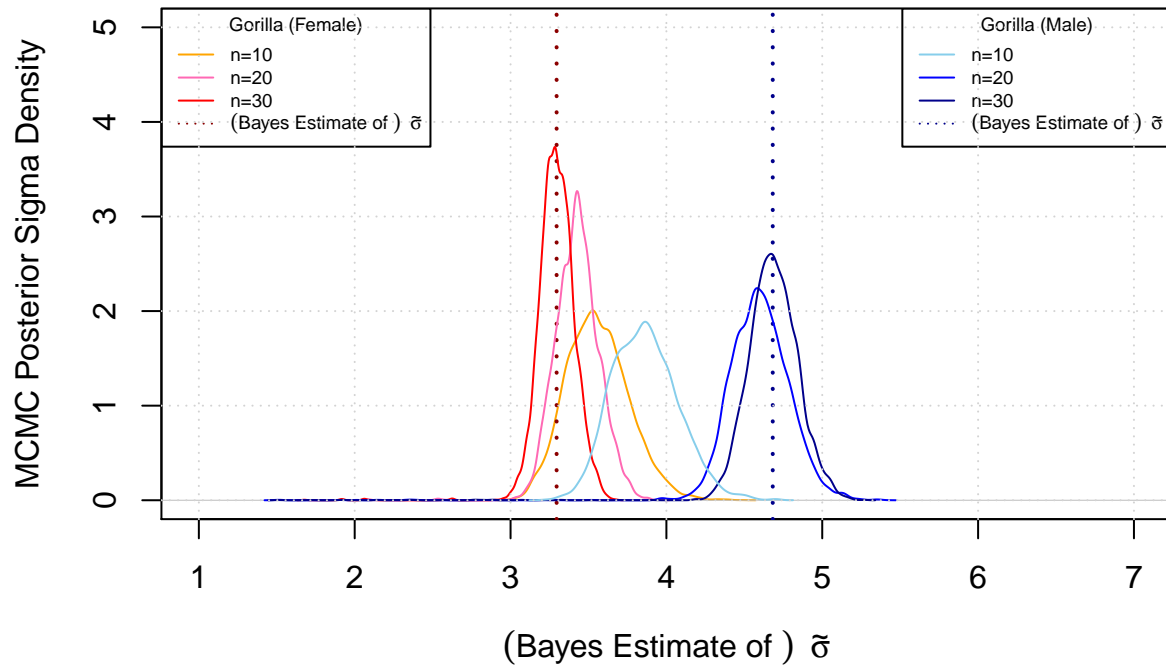
### Combined-Plot for Shape-Variance Comparison Plot of Gorilla Female vs. Male

*#COMBINED PLOT*

```
#plot(density(ress_10[1001:10000,2]), xlim=c(0,8),ylim=c(0,5),col="red",ylab="density (with vague prior)
plot(density(apef10000.10[1000:10000]), xlim=c(1,7),ylim=c(0,5),col="orange", xlab=expression(("Bayes E
#abline(v=theta, col="black",lwd=2, lty=1)
abline(v=mean(apef10000.30), col="red4",lwd=2, lty=3)
#lines(density(apem10000.20), ylim=c(0,12),col="blue")
lines(density(apef10000.20), ylim=c(0,12),col="hotpink")
lines(density(apef10000.30), ylim=c(0,12),col="red")

legend("topleft",cex=0.6,title="Gorilla (Female)", c("n=10","n=20","n=30", expression(("Bayes Estimate o
abline(v=mean(apem10000.30), col="darkblue",lwd=2, lty=3)
#lines(density(apem10000.20), ylim=c(0,12),col="blue")
lines(density(apem10000.20), ylim=c(0,12),col="blue")
lines(density(apem10000.30), ylim=c(0,12),col="blue4")
lines(density(apem10000.10[1000:10000]), ylim=c(0,12),col="skyblue");
legend("topright",cex=0.6,title="Gorilla (Male)", c("n=10","n=20","n=30", expression(("Bayes Estimate o
grid()
```

### MCMC posterior of Variance: Gorilla (Male vs. female) $\sigma^2$ (+ uniform prior)



### Conclusion

It is being observed from posterior Face-shape variance density comparison-plot that there are Bayesian evidence for more face-shape variability in Gorilla male than the same for Gorilla Female.