#Read the data

data1 <- read.csv("XXX1.csv",row.names=1)

data2 <- read.csv (“XXX2.csv”,row.names=1)

m <- ncol(data1)

n <- ncol(data2)

#calculate r value

result1 <- matrix(nrow=m,ncol=n)

for (i in (1:m)){

for(j in (1:n)){

result[i,j] = cor(data1[,i],data2[,j],method="*pearson*")

}

}

row.names(result1) <- colnames(data1)

colnames(result1) <- colnames(data2)

library(reshape2)

Cor1 <- melt(result1,id.vars=c("colnames(result1)", "rownames(result1)"), value.name= "cor")

#Calculate *P*-value

result2 <- NULL

for(i in (1:m)){

 p <- NULL

 for (j in (1:n)){

 a <- cor.test( data1[,i], data2[,j], method="*pearson*")

 p <- c(p, a$p.value)

 }

 result2 <- cbind(result2, p)

}

result2 <- t(result2)

row.names(result2) <- colnames(data1)

colnames(result2) <- colnames(data2)

Cor2 <- melt(result1,id.vars=c("colnames(result2)", "rownames(result2)"), value.name= "*P*-value")

#Combine data

Cor <- cbind(Cor1, Cor2[,3])