

CH5350 Applied Time Series Analysis

Final Project

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Libraries used:

- MTS
 - vars
 - complexplus
-

Introduction

Causality

Causality is an idea of cause-and-effect, although it isn't exactly the same. Causality can be defined as an intervention of one component of a multivariate data on another component at some other time point. A variable X is causal to variable Y if X is the cause of Y or Y is the cause of X.

Project PS

In this project, we try to understand widely used method of empirical measure of **Granger causality**, which stems from econometrics, but finds wide-applicability in several scientific areas. Which involves the implementation of VAR *vector auto-regressive* models, for checking the relation between two variables.

Mathematics

Time Domain

$$\mathbf{z}[k] = \Sigma \mathbf{A}_r \mathbf{z}[k-r] + \mathbf{e}[k]$$

where $\mathbf{z} = [z_1[k], z_2[k], z_3[k], z_4[k], \dots, z_M[k]]^T$ and $\mathbf{A}_r, r = 1, 2, 3, \dots, P$ are $M \times M$ coefficient matrices. \

Granger-causal between $z_i \rightarrow z_j$ doesn't exist iff

$$a_{ij}(r) = 0 \quad \forall r$$

In practice, the estimated VAR model coefficient matrices have to be used; in which case, the existence of Granger-causal relationship from $z_j \rightarrow z_i$ can be tested by conducting a hypothesis test using the statistic.

$$S_{ij} = N \hat{a}_{ij}^T \hat{V}_{ij}^{-1} \hat{a}_{ij}$$

where N is number of observations, $\hat{\mathbf{a}}_{ij} = (a_{ij}(1), a_{ij}(2), a_{ij}(3), \dots, a_{ij}(P))^T$ \mathbf{V}_{ij} is $P \times P$ matrix, obtained by covariance matrix of VAR model.

Frequency Domain

The frequency-domain implementations of Granger causality, of which the direct pathway function (DPF) is an effective measure. The DRF, denoted by $\psi(\omega)$, connects directionally with innovations e_i to variable z_i at each frequency.

$$\psi_{ij}(\omega) = \frac{h_{D,ij}(\omega)}{\sqrt{\sum |h_{D,ij}(\omega)|^2}}$$

where $h_{D,ij}(\omega)$ is the frequency response function of the direct pathway from e_j to z_i .

A variable z_j does not Granger-cause z_i iff

$$|\hat{\psi}_{ij}(\omega)|^2 = 0 \quad \forall \omega$$

R Functions for all equations

a) These functions are for finding S_{ij} time-domain Granger-causality check.

if the returned histogram of S_{ij} realization function is gaussian that implies that i is Granger-cause of j

```
Vhat <- function(Data, Order)
{
  # Read input arguments
  Z = Data
  P = Order

  # Size of the data
  Size = dim(Z)
  N = Size[1]
  M = Size[2]

  # The regressor matrix
  BigZ = {}
  BigZ1 = {}

  for (k in 1:P){
    BigZ = cbind(BigZ, Z[k:(N-P+k-1),])
  }

  # Re-arrange the regressor matrix
  for (j in 1:M){
    BigZ1 = cbind(BigZ1, BigZ[, (seq(j, M*P, M))])
  }

  # The Y matrix
  Y = Z[(P+1):N,]

  # Regressors covariance matrix and its inverse
  SigmaR = (t(BigZ1)%*%BigZ1)/N
  invSigmaR = qr.solve(SigmaR)
```

```

# Estimate VAR model coefficient matrices
Ahat = (qr.solve(t(BigZ1)%*%BigZ1)%*%t(BigZ1))%*%Y

# Innovations covariance matrix
SigmaE = (1/(N-P))*t((Y-BigZ1%*%Ahat))%*%(Y-BigZ1%*%Ahat)

# Covariance matrix of VAR model coefficients
SigmaA = kronecker(invSigmaR,SigmaE)

# Estimation of V matrix
Vmatrix = array(0,dim = c(P,P,M*M))
for (i in seq(1,M*M)){
  Vmatrix[,i] = SigmaA[((i-1)*P+1):(i*P),((i-1)*P+1):(i*P)]
}
return(Vmatrix)
}

#####

aij <- function(Avec, i1, j1){
  # A is vector of matrices each of PXP dimensions
  # which form a coefficients VAR model

  aij_vector <- c()

  for(i in 1:dim(Avec)[3]){
    # paste(c(i,i1, j1))
    aij_vector <- c(aij_vector, Avec[i1,j1,i])
  }

  return(aij_vector)
}

#####

Sij <- function(data,order){
  # aij is vector of A's at given i,j
  # N is total number of observations
  # Vij is output of vhat function
  Vhat_vector <- Vhat(data, order) # 3X3XM2 dimensions
  N <- dim(data)[1]
  M <- dim(data)[2]
  Avec <- model_coefficients(data, order)
  # print(Avec)

  Sij_vector <- c()
  for(i in 1:M){
    for(j in 1:M){
      aij_vec <- aij(Avec, i, j)
      # print(aij_vec)
      Sij_vector = c(Sij_vector, N*t(aij_vec)%*%qr.solve(Vhat_vector[, ,i])%*%aij_vec)
    }
  }
}

```

```

    return(array(Sij_vector, dim = c(M,M)))
}

#####

Sij_realizations <- function(data, order, R=1000){
  M <- dim(data)[2]
  Sij_matrix <- array(0, dim = c(M*M, R))

  for(r in 1:R){
    ek <- matrix(rnorm(dim(data)[1]*dim(data)[2]), nrow = dim(data)[1])
    Sij_vector <- c(Sij(data+ek, order))

    # print(Sij_vector)
    Sij_matrix[,r] <- Sij_vector
  }

  par(mfrow=c(M,M))
  for(i in 1:(M*M))
    hist(Sij_matrix[i,], main = "Sij Distribution")
}

#####

model_coefficients <- function(data, order){
  model <- MTS::VAR(data, order, output = F);
  M <- dim(data)[2]
  Avec <- array(model$Phi, dim = c(M,M, order))
  return(Avec)
}

# Sij(data,3)

```

b) These functions is for finding ψ_{ij} frequency domain Granger-causality check.

```

psi <-function(model, freq_band, plot=F){
  col_size <- dim(model$Phi)[2] # as all the coefficient matrix are stacked together
  M <- dim(model$Phi)[1] # row size of matrix A

  model_order <- col_size/M
  Avec <- array(model$Phi, dim = c(M, M, model_order))
  # print("AVEC TENSOR")
  # print(Avec)

  # Avec got updated
  # Avec is three dimensional tensor with coefficient matrix in order
  # Abar_matrix is matrix obtained from Avec as shown in given in above equation xx

  psi_matrix <- c()

  for(w in 1:length(freq_band)){
    Abar_matrix <- Abar(Avec, w)
    # print("ABAR MATRIX")
  }
}

```

```

# print(Abar_matrix)

psi_vector <- c()
for(psi_i in 1:M){
  for(psi_j in 1:M){
    # M is minor matrix of Abar obtained by eliminating ith row & jth col
    # Mbar is minor matrix of Abar obtained by eliminating ith & jth row & col
    h_denominator2 <- 0
    # for denominator calculation
    for(i in 1:M){
      Mij <- Abar_matrix[-c(i), -c(psi_j)]
      Mbarij <- Abar_matrix[-c(i,psi_j), -c(i,psi_j)]
      h_denominator2 <- h_denominator2 +
        abs(hij(Abar_matrix, Mij, Mbarij, i, psi_j))^2
    }
    h_denominator <- sqrt(h_denominator2)
    # print("h_denominator")
    # print(h_denominator)

    # for numerator calculation
    Mij <- Abar_matrix[-c(psi_i), -c(psi_j)]
    Mbarij <- Abar_matrix[-c(psi_i,psi_j), -c(psi_i,psi_j)]
    h_numerator <- hij(Abar_matrix, Mij, Mbarij, psi_i, psi_j)

    # psi_value calculation...
    psi_value <- h_numerator/h_denominator
    psi_vector <- c(psi_vector, psi_value)
  }
}
# print("PSI VECTOR")
# print(psi_vector)
psi_matrix <- rbind(psi_matrix, psi_vector)
}

psi_matrix <- array(psi_matrix, dim = c(length(freq_band),M*M))

# plot magnitude
if(plot == T){
  plot.ts(abs(psi_matrix)^2)
  #plot.ts(Arg(psi_matrix))
}
return(t(abs(psi_matrix)^2))
}

#####

Abar <- function(Avec, w){
  # Abar is matrix obtained by sum of Avec tensor(3D)
  # abar zero initialization

  abar <- 0
  M <- dim(Avec)[1][1] # Dimension of coefficient matrix of model
  I <- diag(M) # Identity matrix of order M

```

```

# abar updates...
for(r in 1:dim(Avec)[3]){
  abar <- abar + Avec[, ,r]*exp(-1i*r*w)
}

abar <- I - abar

# abar matrix return..
return(abar)
}

#####

hij <- function(Abar_vector, Mij, Mbarij, i, j){
  # equation implementation of h(w)
  dr <- Det(Abar_vector) # denominator of hij(w)

  if(i == j){
    if(length(Mij) == 1){
      nr <- Mij
    }
    else{
      nr <- Det(Mij) # numerator in case of i==j
    }
  }
  else{
    if(length(Mbarij) == 1){
      nr <- -1*Abar_vector[i,j]*Mbarij
    }
    else{
      nr <- -1*Abar_vector[i,j]*Det(Mbarij) # numerator in other cases
    }
  }
  # print("hijD")
  # print(nr/dr)
  return(nr/dr)
}

```

c & d) closed-form approximation for distribution of $|\hat{\psi}_{ij}(\omega)|^2$ functions involved are..

Quantile calculation for large number of realizations standard gaussian distribution is assumed

```

cform_distribution <- function(zk, R, order, freq_band){
  # R is simulation steps

  dpf_datasets <- gen_realization(zk, R)

  inference_matrix_realizations <- array(1,
                                         dim = c(1, dim(zk)[2]*dim(zk)[2]*length(freq_band)))
  inference_matrix <- array(0, dim = c(dim(zk)[2], dim(zk)[2]))

  psi_tensor <- array(0, dim = c(dim(zk)[2]^2, length(freq_band), R))

  for(i in 1:R){

```

```

    # print(zk_datasets[, , i])
    model <- MTS::VAR(dpf_datasets[, , i], order, output = FALSE)
    psi_tensor[, , i] <- psi(model, freq_band)
  }
  zetaij_matrix <- zetaij(psi_tensor)

  # print("ZETA MATRIX")
  # print(zetaij_matrix)

  # check on data...
  psi_matrix <- psi(MTS::VAR(zk, order, output = F), freq_band)

  # print("PSI MATRIX")
  # print(psi_matrix)

  # compare psi_matrix with zetaij_matrix
  # 1 if ej is granger-cause of zi else 0
  print("INFERENCE MATRIX")
  inference_matrix_realizations[which(psi_matrix < zetaij_matrix)] = 0
  inference_matrix_realizations <- array(inference_matrix_realizations,
                                         dim = c(dim(zk)[2], dim(zk)[2], length(freq_band)))

  for(i in 1:dim(zk)[2]){
    for(j in 1:dim(zk)[2]){
      if(sum(inference_matrix_realizations[i,j,]) > 0){
        inference_matrix[i,j] = 1
      }
    }
  }

  # print(inference_matrix_realizations)
  # print(inference_matrix)

  return(t(inference_matrix))
}

#####

gen_realization <-function(zk, R){
  # null initialized datasets
  zk_datasets <- array(0, dim = c(dim(zk)[1], dim(zk)[2], R))
  for(i in 1:R){
    # step 1 DFT of data
    zk_dft <- fft(zk)

    # step 2 Randomize the phase by replacing the phase with a randomly generated
    # sequence over the interval [0, 2pi]
    # set.seed(0)
    phase_random <- runif(dim(zk)[1], 0, 2*pi)

    # zk_dft updated with random phase
    zk_dft_phase <- zk_dft * exp(1i*phase_random)

```

```

    # check for ifft calculation only real part...
    zk_datasets[, , i] <- Re(fft(zk_dft_phase, inverse=T))/length(zk)
  }

  return(zk_datasets)
}

#####

zetaij <- function(psi_tensor, alpha=0.95){
  M2 <- dim(psi_tensor)[1]
  w <- dim(psi_tensor)[2]
  zeta_mat <- array(0, dim = c(M2, w))
  for(i in 1:M2){
    for(j in 1:w){
      ts <- psi_tensor[i, j, ]

      # hist_val <- hist(ts, plot=F)
      # alpha% quantile distribution calculation
      zeta_mat[i,j] <- qnorm(alpha)*sd(ts)/sqrt(length(ts))+mean(ts)
    }
  }
  return(zeta_mat)
}

```

e) Simulation using VARMAsim based on SIC

```

N <- 2000
A1 <- matrix(c(0.3, 0, 0, 0.6, 0.4, 0, 0, 0.4, 0.5), nrow=3, byrow = T)
A2 <- matrix(c(0.2, 0, 0, 0, 0.5, 0, 0, 0.3, 0.4), nrow=3, byrow = T)

sig <- diag(3)
# set.seed(20)
zksim <- MTS::VARMAsim(N, arlags = c(1, 2), phi = cbind(A1,A2), sigma = sig)

sim_model <- vars::VARselect(zksim$series, lag.max = 10)

# best model is...
best_order <- which.min(sim_model$criteria["SC(n)",])
cat("Best VAR model based on SIC is: ", best_order)

## Best VAR model based on SIC is: 2
# model is

best_model <- MTS::VAR(zksim$series, best_order)

## Constant term:
## Estimates: 0.01586428 0.04402045 0.000985901
## Std.Error: 0.02377998 0.02295323 0.02382566
## AR coefficient matrix
## AR( 1 )-matrix
##      [,1] [,2] [,3]
## [1,] 0.29096 0.031 -0.0025
## [2,] 0.62542 0.381 -0.0184

```

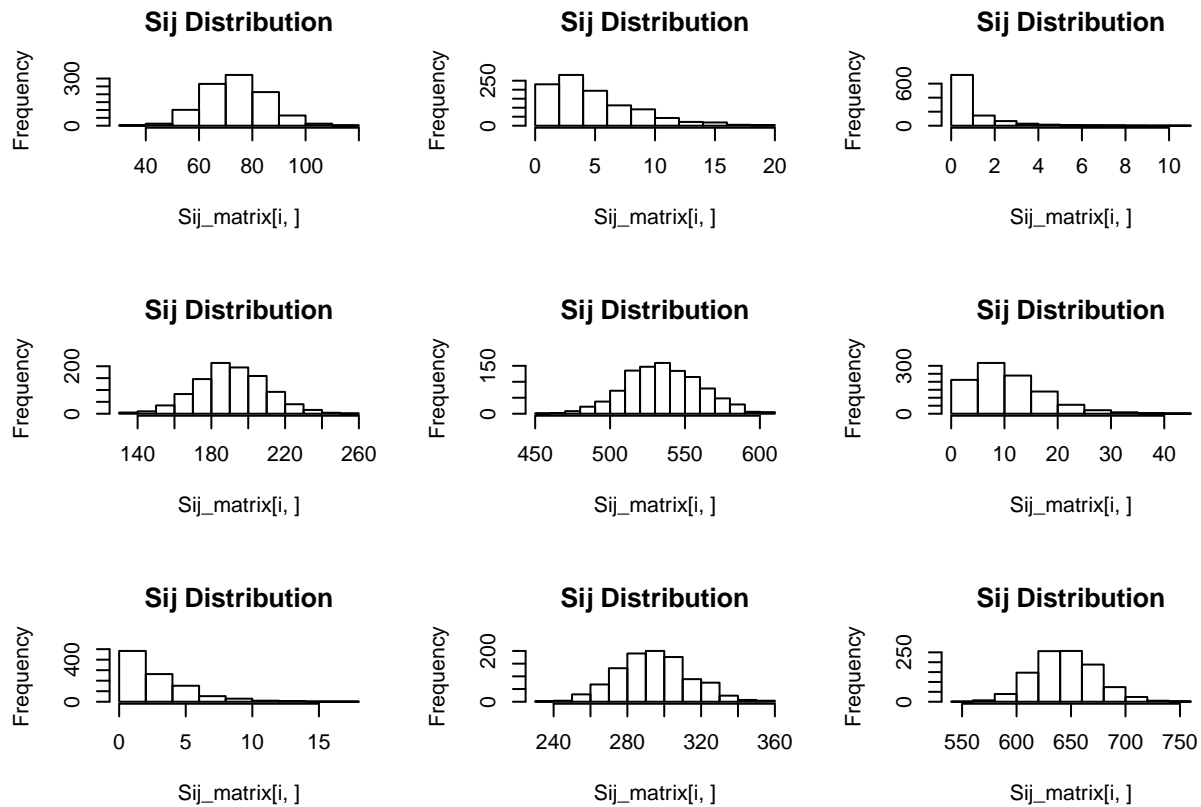


```

## [3,] 0.00442 0.423 0.5380
## standard error
##      [,1] [,2] [,3]
## [1,] 0.0221 0.0194 0.0205
## [2,] 0.0213 0.0187 0.0197
## [3,] 0.0221 0.0195 0.0205
## AR( 2 )-matrix
##      [,1] [,2] [,3]
## [1,] 0.1908 -0.0321 0.00348
## [2,] 0.0123 0.5158 0.01554
## [3,] -0.0247 0.2576 0.36570
## standard error
##      [,1] [,2] [,3]
## [1,] 0.0256 0.0206 0.0192
## [2,] 0.0247 0.0199 0.0185
## [3,] 0.0257 0.0207 0.0192
##
## Residuals cov-mtx:
##      [,1] [,2] [,3]
## [1,] 1.02942579 -0.01126937 0.03728825
## [2,] -0.01126937 0.95909089 -0.05372049
## [3,] 0.03728825 -0.05372049 1.03338467
##
## det(SSE) = 1.015884
## AIC = 0.03375876
## BIC = 0.08416688
## HQ = 0.05226757

# time domain causality test : inference based on Gaussian
# distributed histogram
Sij_realizations(zksim$series, best_order)

```



```
# frequency domain causality test
cform_distribution(zksim$series, 50, best_order, w)
```

```
## [1] "INFERENCE MATRIX"
```

```
##      [,1] [,2] [,3]
## [1,]    0    1    0
## [2,]    1    1    1
## [3,]    0    1    1
```

f)

- For large number of realizations both time and frequency domain methods yield correct results for *Granger-causality*
- Frequency domain *Granger-causality* will give more accurate results as it uses more number of observations
- Answer remains invariant with given realization, but is highly dependent on number of Realizations which can be observed:

```
cform_distribution(Zk, R=2, 2, w)
```

```
## [1] "INFERENCE MATRIX"
```

```
##      [,1] [,2] [,3]
## [1,]    1    0    0
## [2,]    0    1    0
## [3,]    1    0    1
```

```
cform_distribution(Zk, R=20, 2, w)
```

```
## [1] "INFERENCE MATRIX"
```

```
##      [,1] [,2] [,3]
```

```
## [1,] 0 0 0
## [2,] 1 1 0
## [3,] 0 0 1
```

```
cform_distribution(Zk, R=50, 2, w)
```

```
## [1] "INFERENCE MATRIX"
```

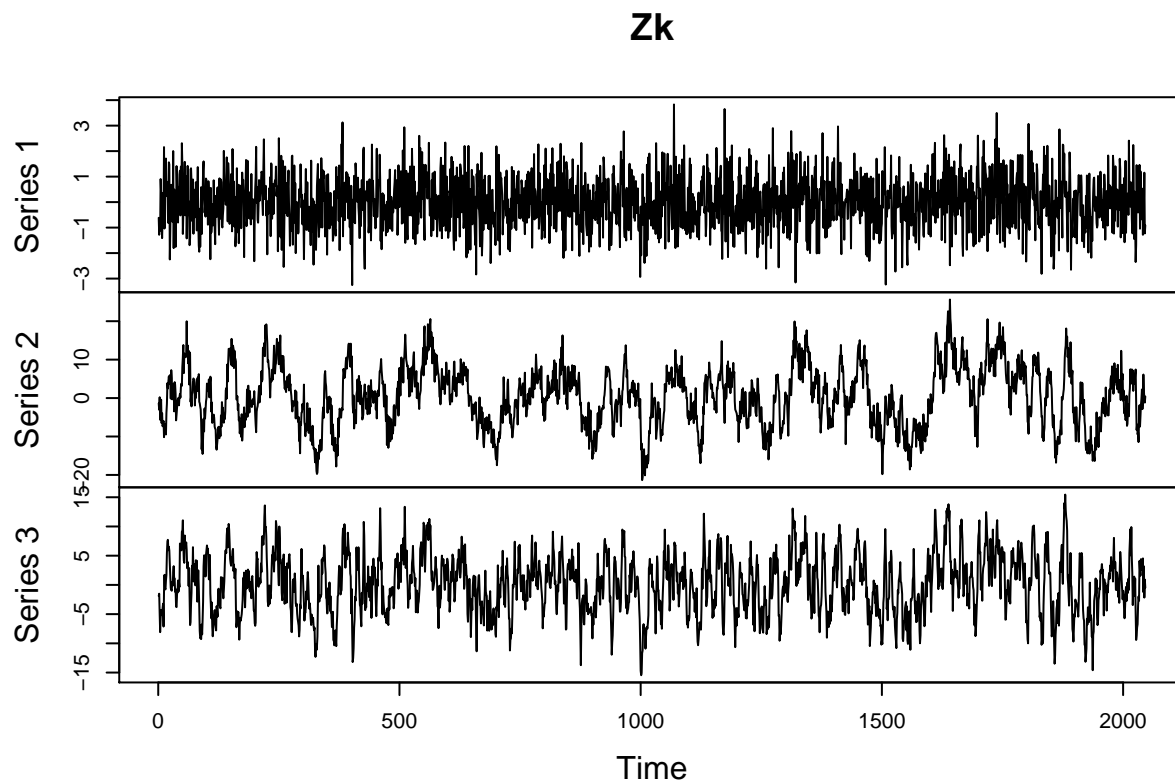
```
##      [,1] [,2] [,3]
## [1,] 0 0 0
## [2,] 0 1 0
## [3,] 0 0 1
```

2)

Data Pre-Processing & Model Building

Plots and nature of data

```
plot.ts(Zk)
```



Normalization

```
# normaliaztion to bring all series in same range
```

```
Zk[,1] <- (Zk[,1]-mean(Zk[,1]))/sd(Zk[,1])
```

```

Zk[,2] <- (Zk[,2]-mean(Zk[,2]))/sd(Zk[,2])
Zk[,3] <- (Zk[,3]-mean(Zk[,3]))/sd(Zk[,3])

## 1a & 1b implementation
data_model <- vars::VARselect(Zk, lag.max = 10)

# best model is...
data_order <- which.min(data_model$criteria["SC(n)",])

Sij(Zk, data_order)

```

```

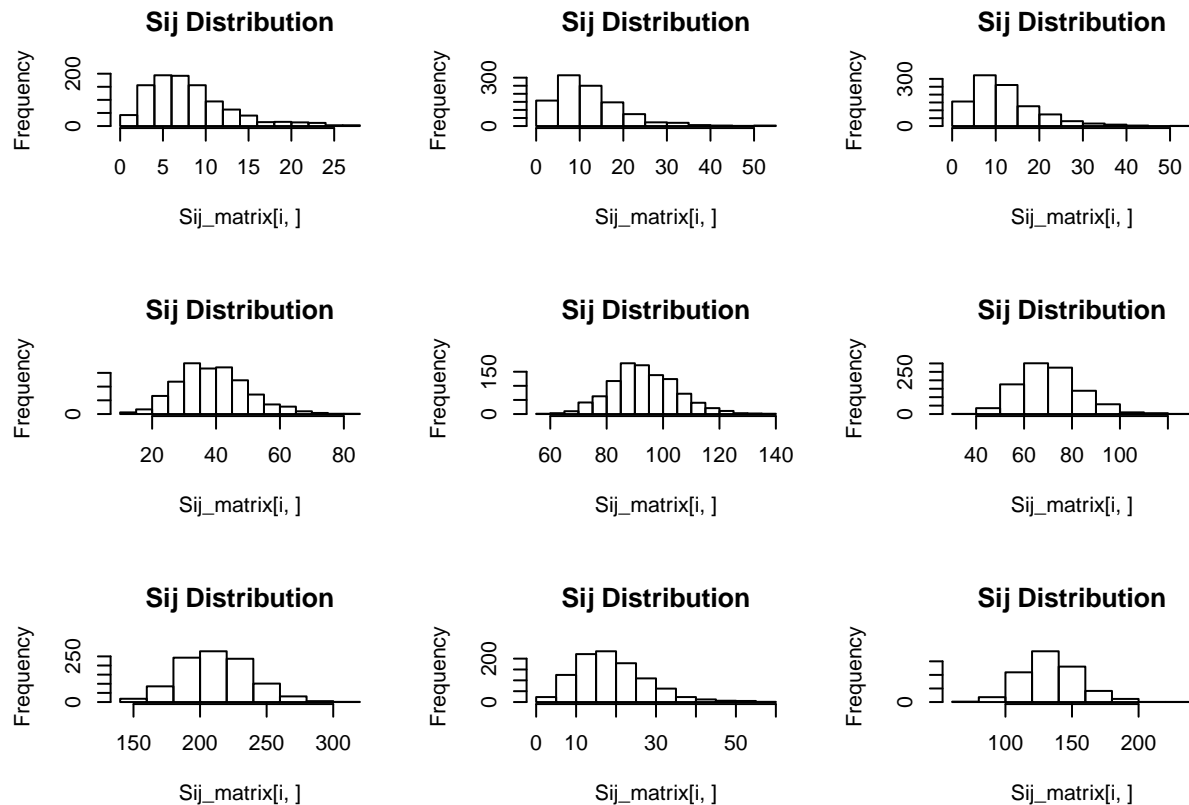
##           [,1]      [,2]      [,3]
## [1,]  32.54673 325.30671 1891.19592
## [2,]  37.02977 157.12525  14.48791
## [3,] 348.20709  38.20473   67.51265

```

```

# evaluation for realizations...
question1_a <- Sij_realizations(Zk, data_order, 1000)

```



```

# evaluation in freq domain
cform_distribution(Zk, R=50, data_order, w)

```

```

## [1] "INFERENCE MATRIX"
##           [,1] [,2] [,3]
## [1,]      0    0    0
## [2,]      1    1    0
## [3,]      1    0    1

```

```
model <- MTS::VAR(Zk, data_order, output = F)
```

```
question1_b <- psi(model, w, T)
```

$\text{abs}(\text{psi_matrix})^2$

