

S23 NSERC USRA Simulation Studies for Underreported Infectious Disease Time Series Models

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```
library(ggplot2)
library(tinytex)
library(abind)

# Set parameter values
phi <- 0.8 # Autoregressive parameter
nu <- 5 # Mean parameter

# create sequence of pi's
pi_seq <- seq(0.1, 1, 0.1)

# initialize the objects that will be used to store all the results
df_pi <- c()
df_phi <- c()
df_nu <- c()

# for each pi...
for (pi in pi_seq) {

  # Define number of simulations and time series lengths
  num_simulations <- 1000
  T_values <- c(30, 50, 100, 500, 1000) # Updated values of T

  # Function to compute Method of Moments estimators
  compute_estimators <- function(z) {
    rho_1 <- acf(z, lag.max = 1, plot = FALSE)$acf[2] # Autocorrelation at lag 1

    # Handle missing or NaN values
    if (is.na(rho_1) || is.nan(rho_1)) {
      rho_1 <- 0.01
    }

    # Method of Moments estimators
    if (abs(1 - mean(z) / var(z)) < 0.001 || var(z) <= 0.001) {
      phi_hat <- 0.001 # Assign phi directly as 0.001
    } else {
      phi_hat <- (1 / rho_1) * (1 - mean(z) / var(z))
    }

    # Check if variance is close to zero
```

```

if (var(z) <= 0.001) {
  pi_hat <- 0.001 # Assign a small positive value to pi_hat
} else {
  pi_hat <- 1 - var(z) / mean(z) * (1 - rho_1 / phi_hat)
}

nu_hat <- (1 - phi_hat) * mean(z) / pi_hat

return(c(pi_hat, phi_hat, nu_hat))
}

# Function to simulate and estimate
simulate_and_estimate <- function(pi, phi, nu, T) {
  z <- rep(0, T) # Initialize time series
  z[1] <- 5 # Initialize Z_1 at 5

  # Simulate time series
  for (i in 2:T) {
    z[i] <- rpois(1, lambda = nu + phi * z[i-1])
  }

  # Thin the time series using binomial thinning
  y <- rbinom(T, z, pi)

  # Check for missing or NaN values
  if (any(is.na(y)) || any(is.nan(y))) {
    return(c(NA, NA, NA))
  }

  # Compute Method of Moments estimators
  estimators <- compute_estimators(y)

  return(estimators)
}

# Initialize matrix to store Method of Moments estimates and quantiles
estimates_matrix <- matrix(0, nrow = length(T_values), ncol = 3)
quantiles_matrix <- array(0, dim = c(length(T_values), 3, 3))

# Perform simulation study for different values of T
for (j in 1:length(T_values)) {
  T <- T_values[j]
  mom_estimators <- matrix(0, nrow = num_simulations, ncol = 3)

  # Run simulations
  for (i in 1:num_simulations) {
    mom_estimators[i, ] <- simulate_and_estimate(pi, phi, nu, T)
  }

  # Calculate average error
  avg_error <- abs(colMeans(mom_estimators) - c(phi, pi, nu))

  # Check for "Inf" or "NaN" average error estimates and replace with numeric values

```

```

avg_error[is.infinite(avg_error)] <- NA
avg_error[is.nan(avg_error)] <- NA

# Calculate quantiles
valid_simulations <- complete.cases(mom_estimators)
quantiles <- apply(mom_estimators[valid_simulations, ], 2, quantile, probs = c(0.025, 0.5, 0.975))

# Store Method of Moments estimates and quantiles in the matrices
estimates_matrix[j, ] <- colMeans(mom_estimators, na.rm = TRUE)
quantiles_matrix[j, , ] <- quantiles

# Print average error and quantiles
#cat("T =", T, "Average Error:", avg_error, "\n")
#cat("T =", T, "Quantiles:", quantiles, "\n")
}

# Create data frames for plotting
df_pi_pi <- data.frame(T = as.character(T_values), Estimate = quantiles_matrix[, 2, 1],
                      Lower = quantiles_matrix[, 1, 1], Upper = quantiles_matrix[, 3, 1])
df_phi_pi <- data.frame(T = as.character(T_values), Estimate = quantiles_matrix[, 2, 2],
                      Lower = quantiles_matrix[, 1, 2], Upper = quantiles_matrix[, 3, 2])
df_nu_pi <- data.frame(T = as.character(T_values), Estimate = quantiles_matrix[, 2, 3],
                      Lower = quantiles_matrix[, 1, 3], Upper = quantiles_matrix[, 3, 3])

# Append the results for this pi value to the corresponding data frame
df_pi <- abind(df_pi, df_pi_pi, along = 3) # added along = 3
df_phi <- abind(df_phi, df_phi_pi, along = 3) # along = 3
df_nu <- abind(df_nu, df_nu_pi, along = 3) # along = 3
}

# Now we have 1 matrix for each value of pi.
# We will plot each one separately.

# Plot for estimate of pi for each value of pi
pi_plots <- list()
for (i in 1:length(pi_seq)) {
  pi_plots[[i]] <- ggplot(as.data.frame(df_pi[, , i]), aes(x = T, y = as.numeric(Estimate))) + # added as
    geom_point() +
    geom_errorbar(aes(ymin = as.numeric(Lower), ymax = as.numeric(Upper)), width = 0.3) + #as.numeric t
    geom_hline(yintercept = pi_seq[i], linetype = "dashed", color = "red") +
    labs(title = paste("Method of Moments Estimates for the Reporting Probability (pi =", pi_seq[i], ")
      x = "Time Series Length (T)", y = "Estimate of the Reporting Probability (pi)") +
    theme_minimal() +
    scale_x_discrete(limits = as.character(T_values))
}

# Plot for estimate of phi for each value of pi
phi_plots <- list()
for (i in 1:length(pi_seq)) {
  phi_plots[[i]] <- ggplot(as.data.frame(df_phi[, , i]), aes(x = T, y = as.numeric(Estimate))) +
    geom_point() +
    geom_errorbar(aes(ymin = as.numeric(Lower), ymax = as.numeric(Upper)), width = 0.3) +
    geom_hline(yintercept = phi, linetype = "dashed", color = "red") +

```

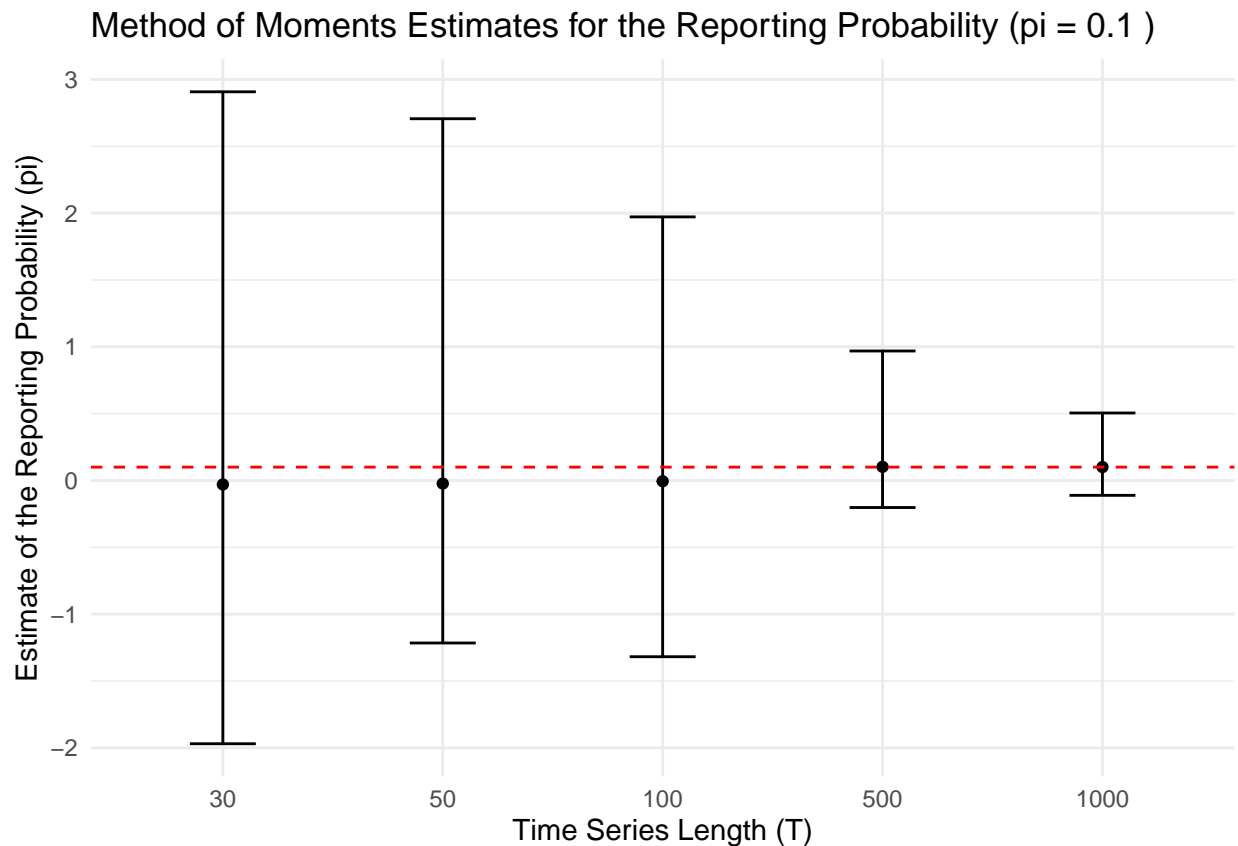
```

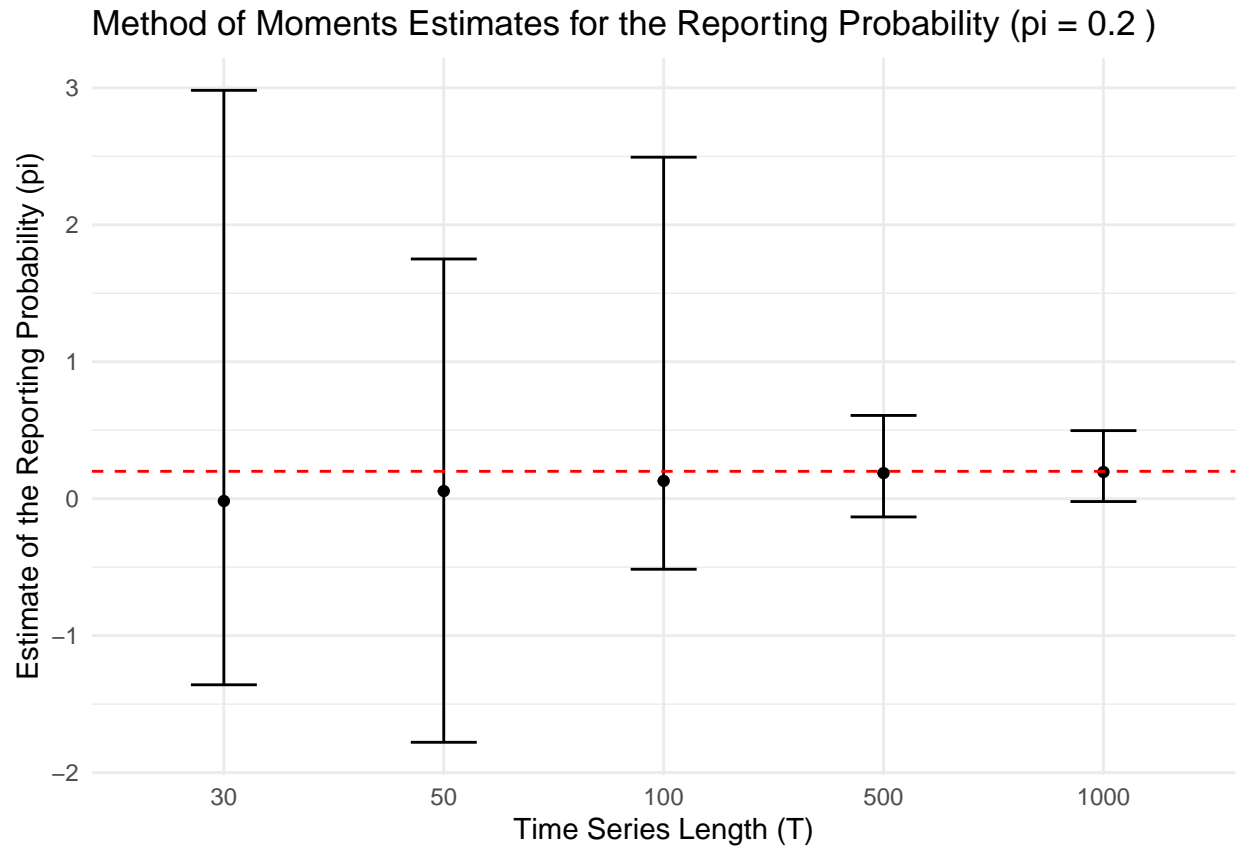
labs(title = paste("Method of Moments Estimates for the Autoregressive Parameter ( $\phi$ ) for  $\pi$  =", pi),
     x = "Time Series Length (T)", y = "Estimate of the Autoregressive Parameter ( $\phi$ )") +
theme_minimal() +
scale_x_discrete(limits = as.character(T_values))
}

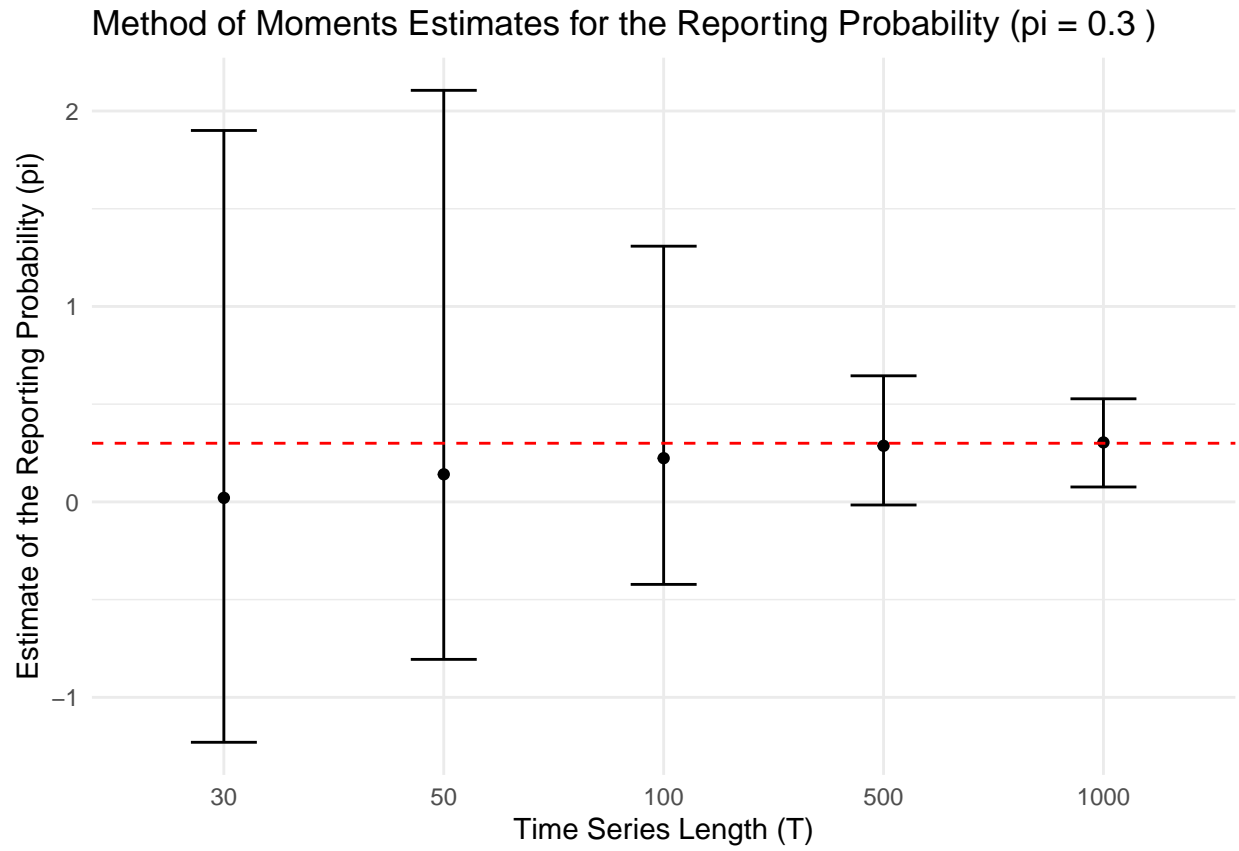
# Plot for estimate of nu for each value of pi
nu_plots <- list()
for (i in 1:length(pi_seq)) {
  nu_plots[[i]] <- ggplot(as.data.frame(df_nu[, , i]), aes(x = T, y = as.numeric(Estimate))) +
    geom_point() +
    geom_errorbar(aes(ymin = as.numeric(Lower), ymax = as.numeric(Upper)), width = 0.3) +
    geom_hline(yintercept = nu, linetype = "dashed", color = "red") +
    labs(title = paste("Method of Moments Estimates for the Mean Parameter ( $\nu$ ) for  $\pi$  =", pi_seq[i]),
         x = "Time Series Length (T)", y = "Estimate of the Mean Parameter ( $\nu$ )") +
    theme_minimal() +
    scale_x_discrete(limits = as.character(T_values))
}

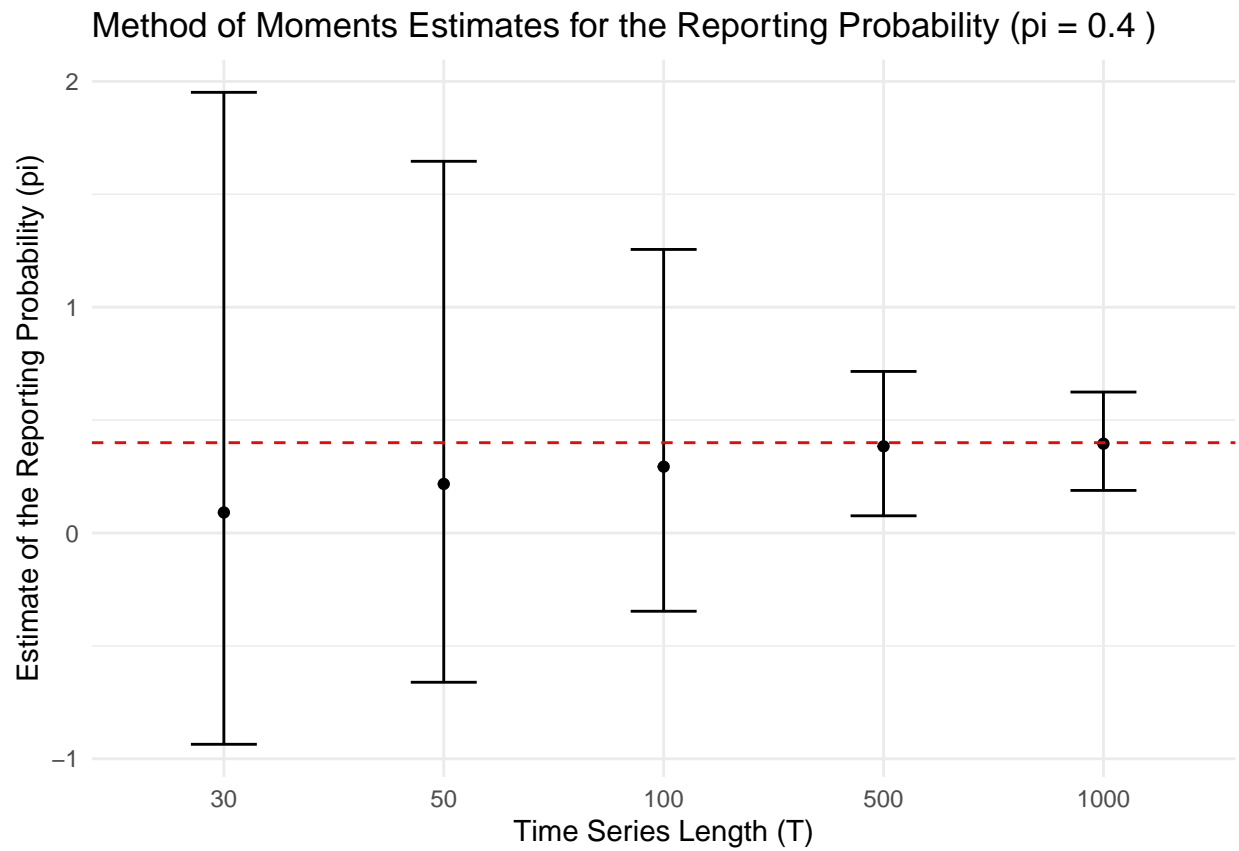
# Display the plots for each value of pi
for (i in 1:length(pi_plots)) {
  print(pi_plots[[i]])
}

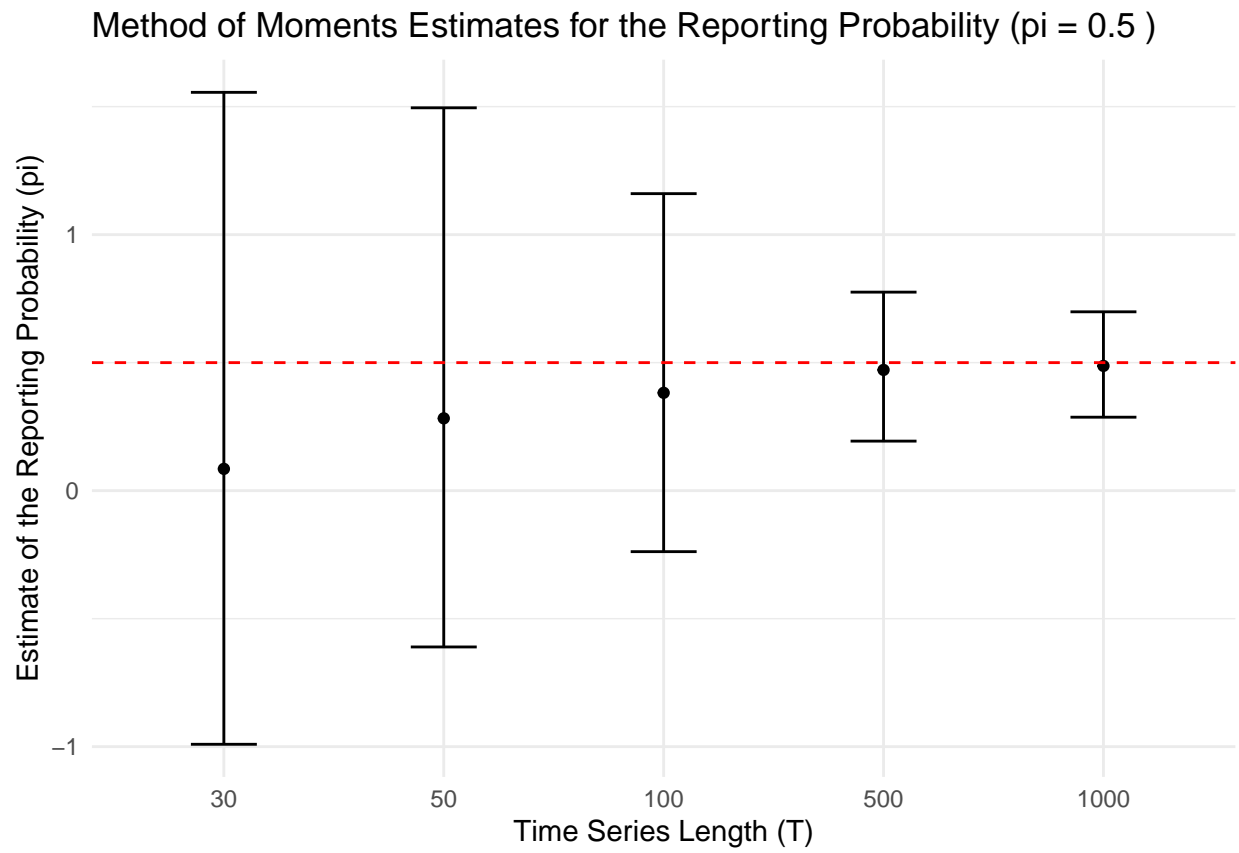
```

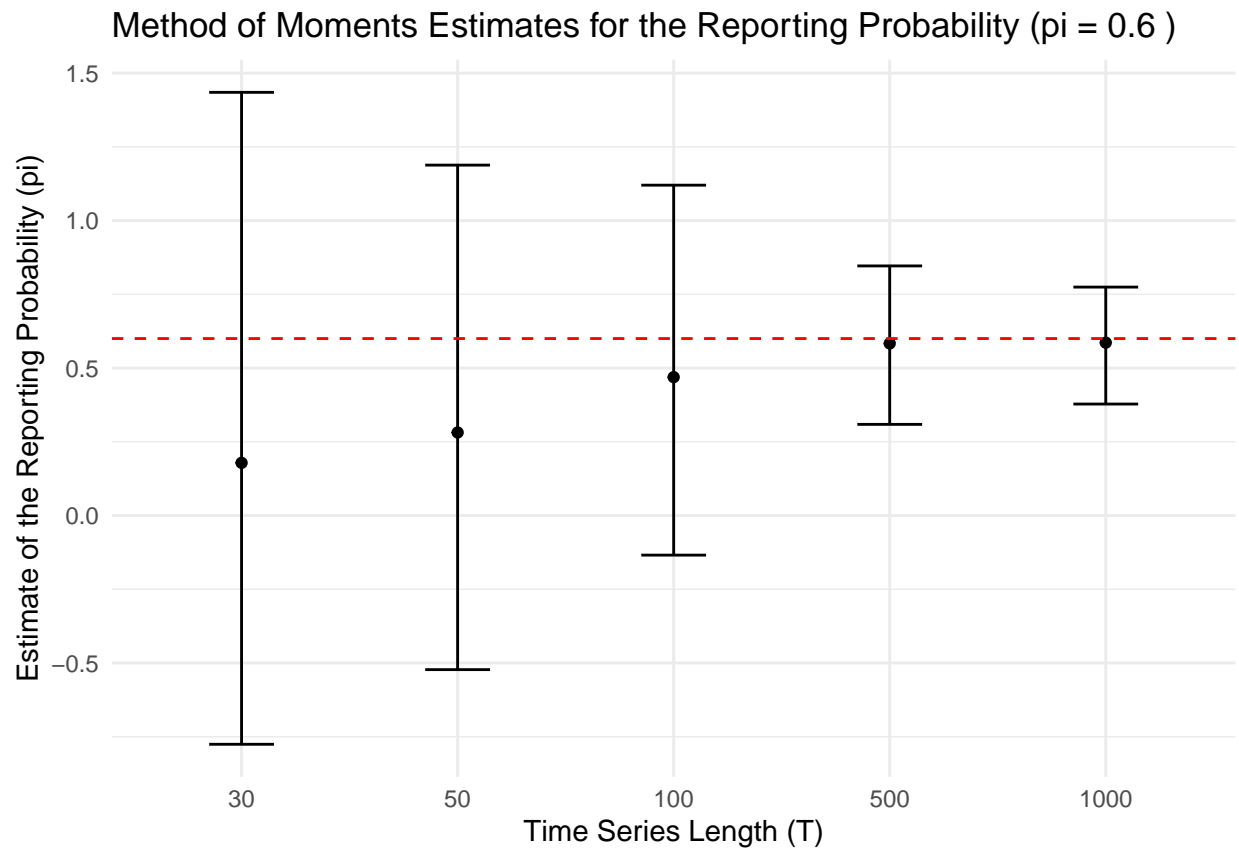


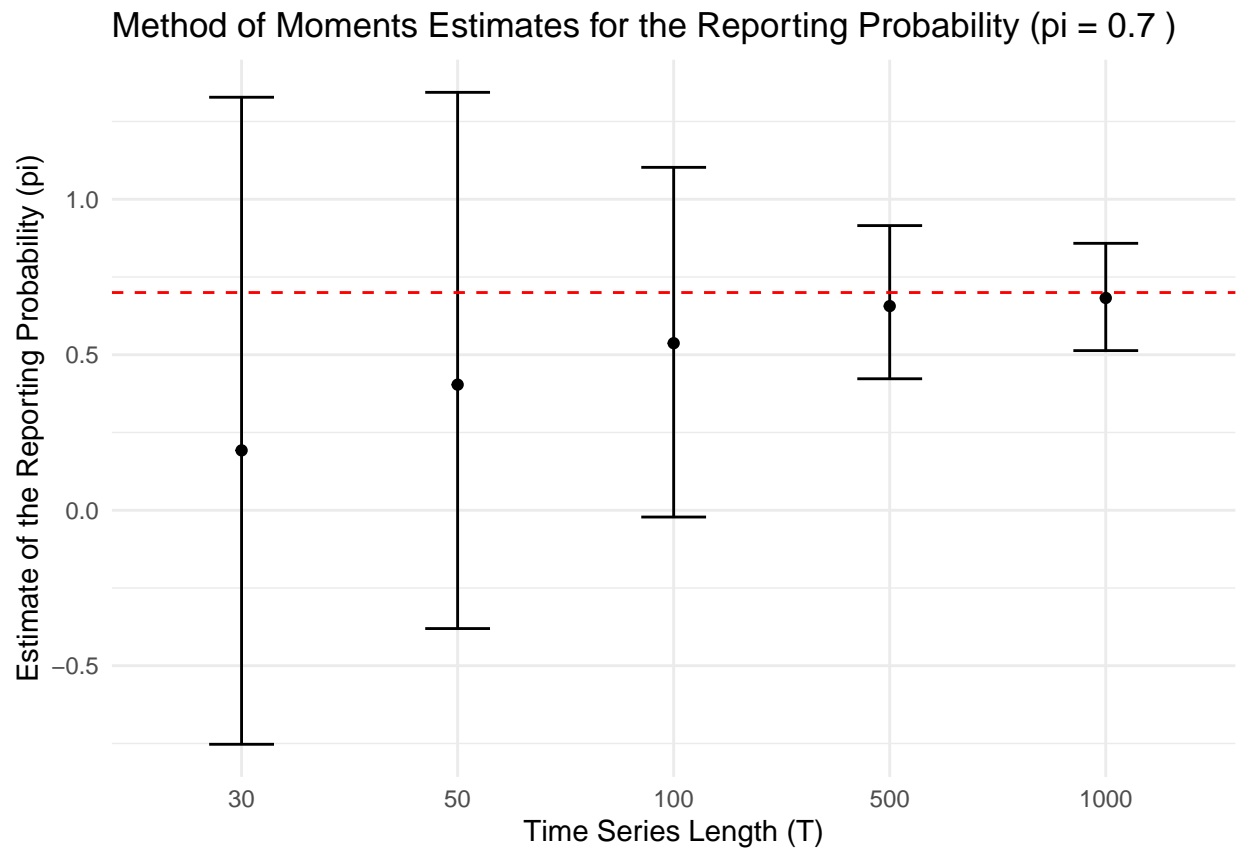


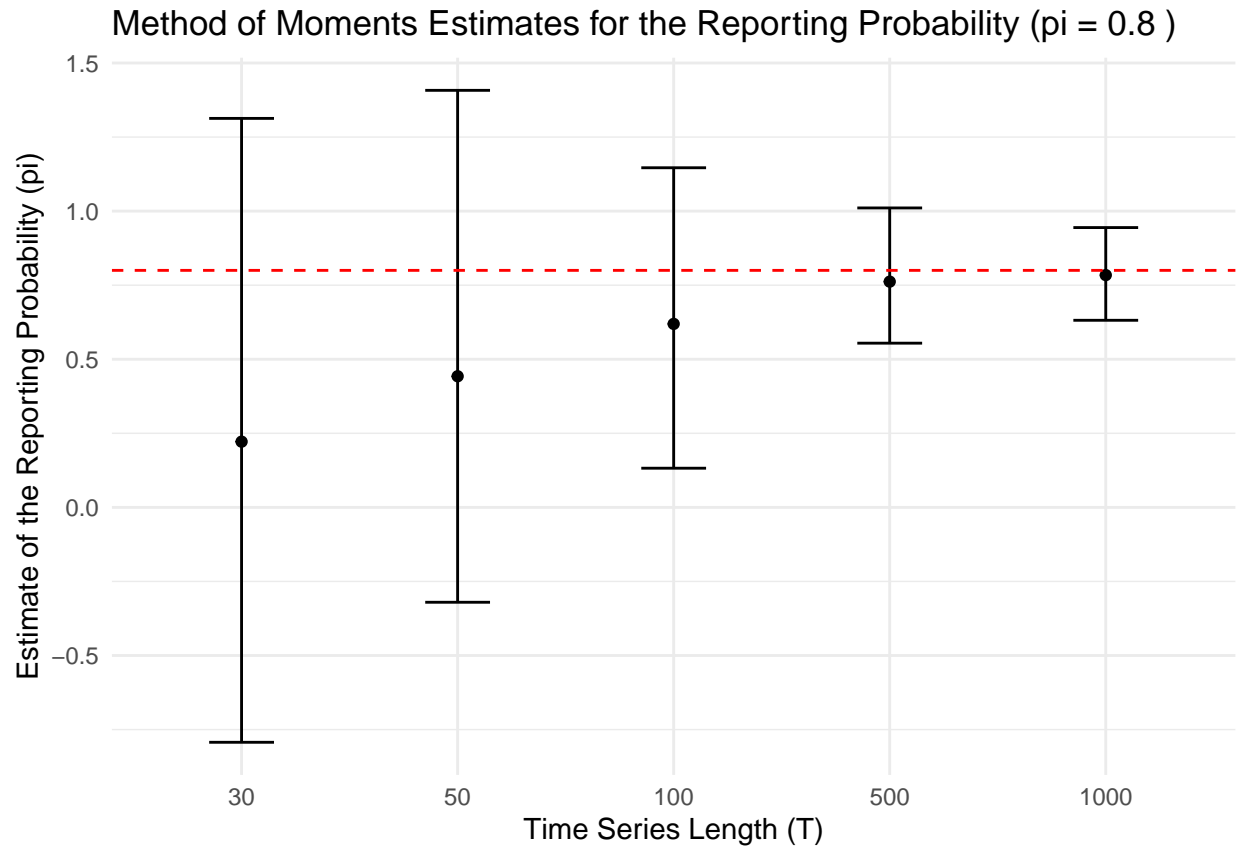


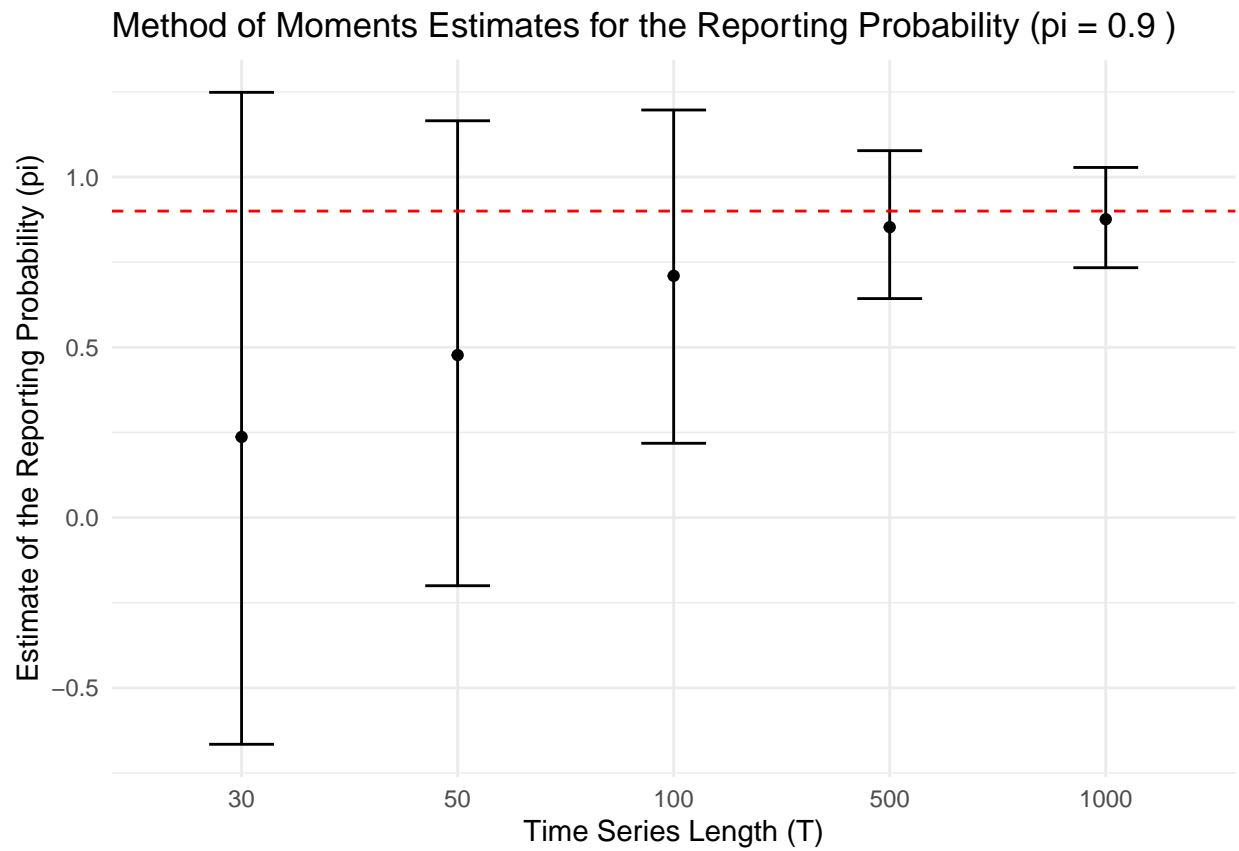


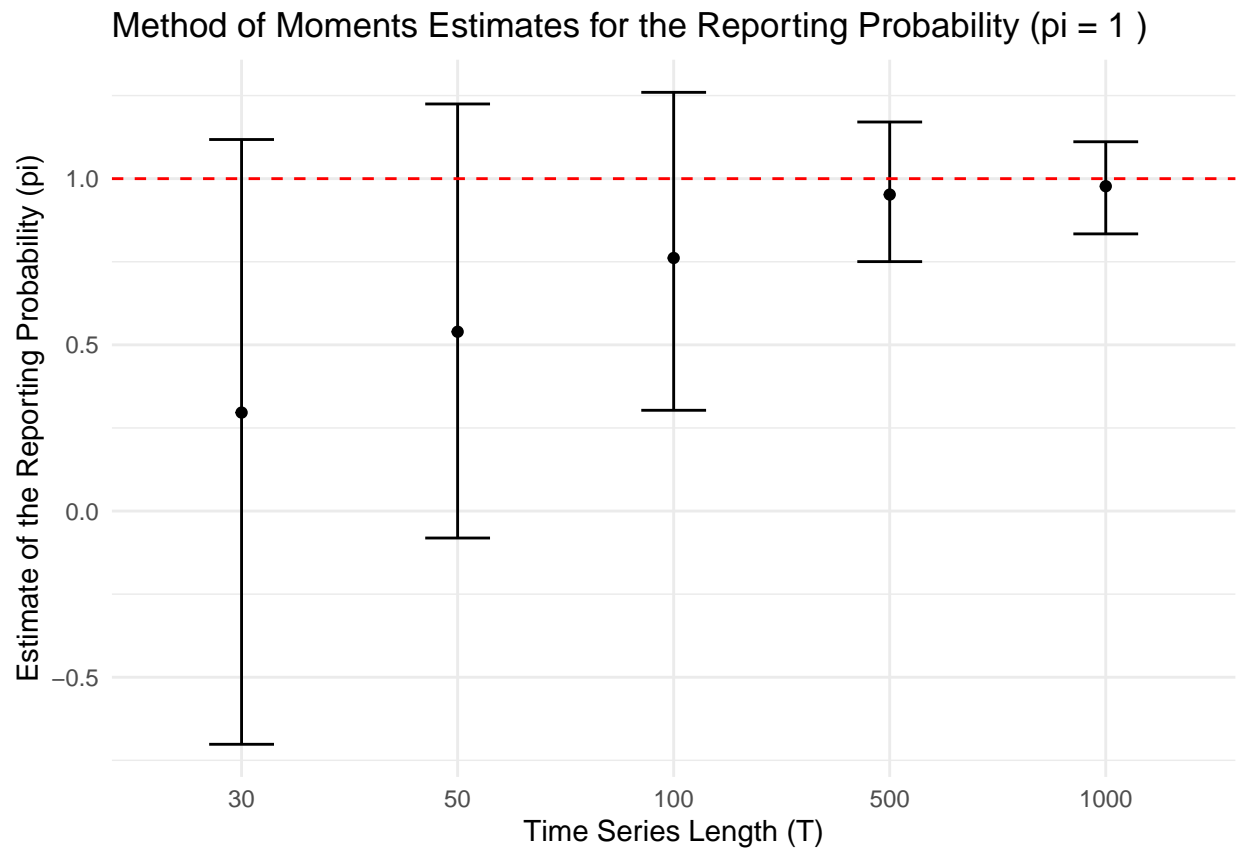




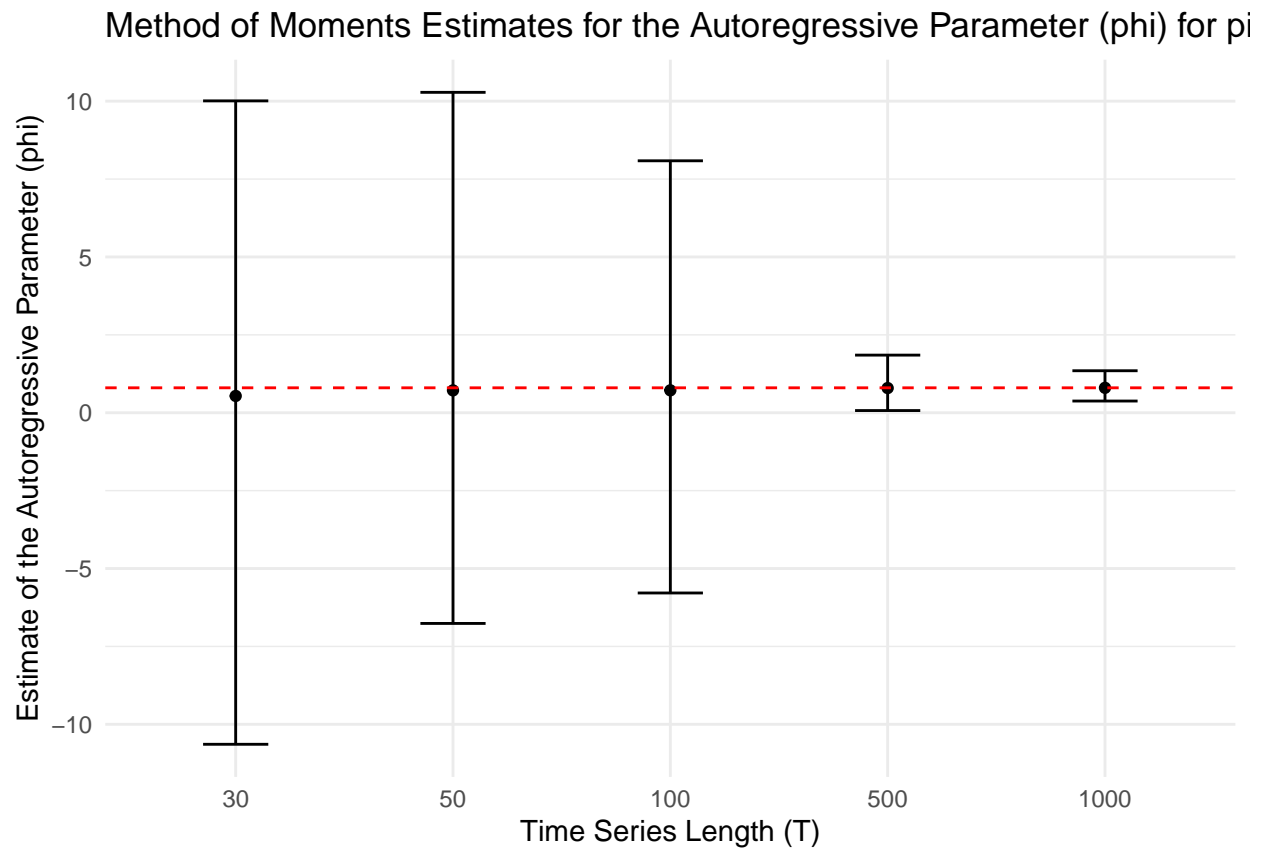


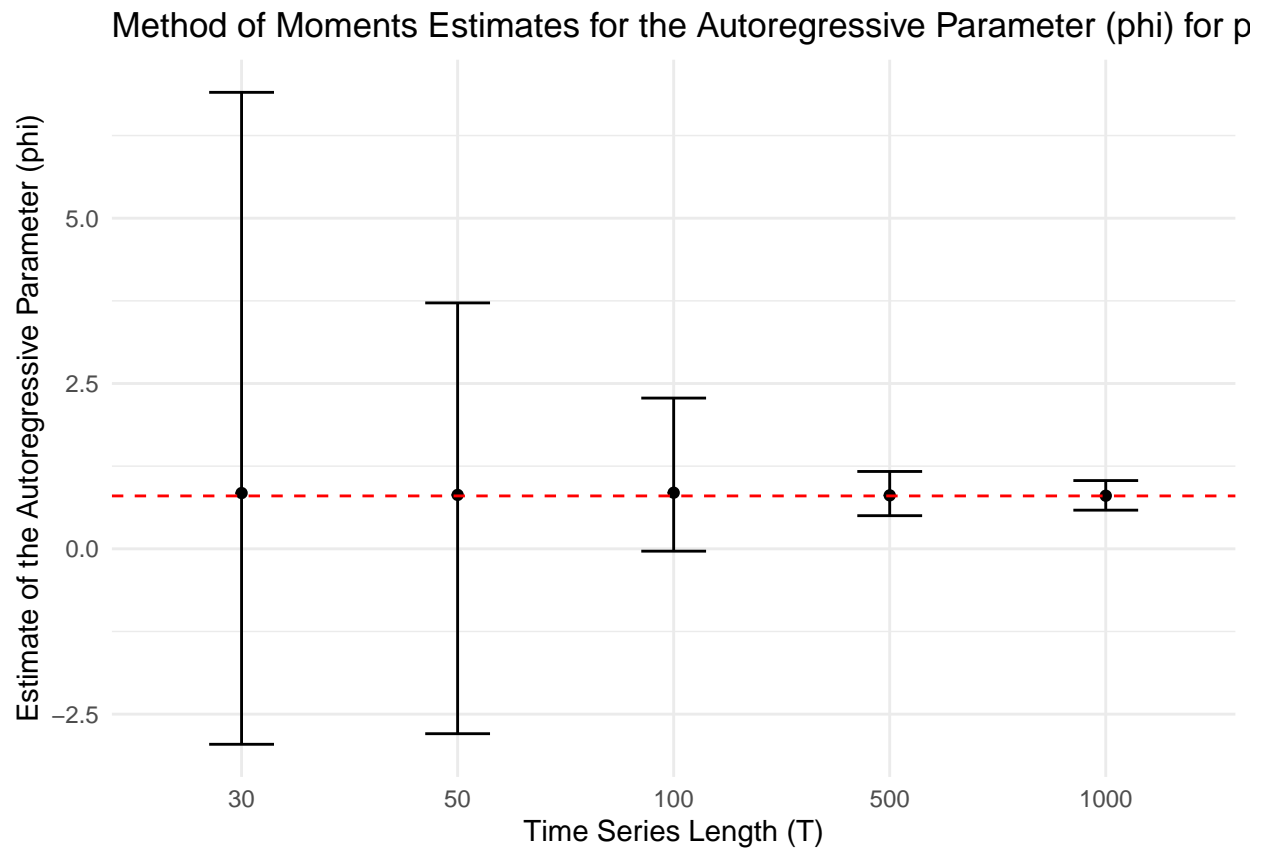


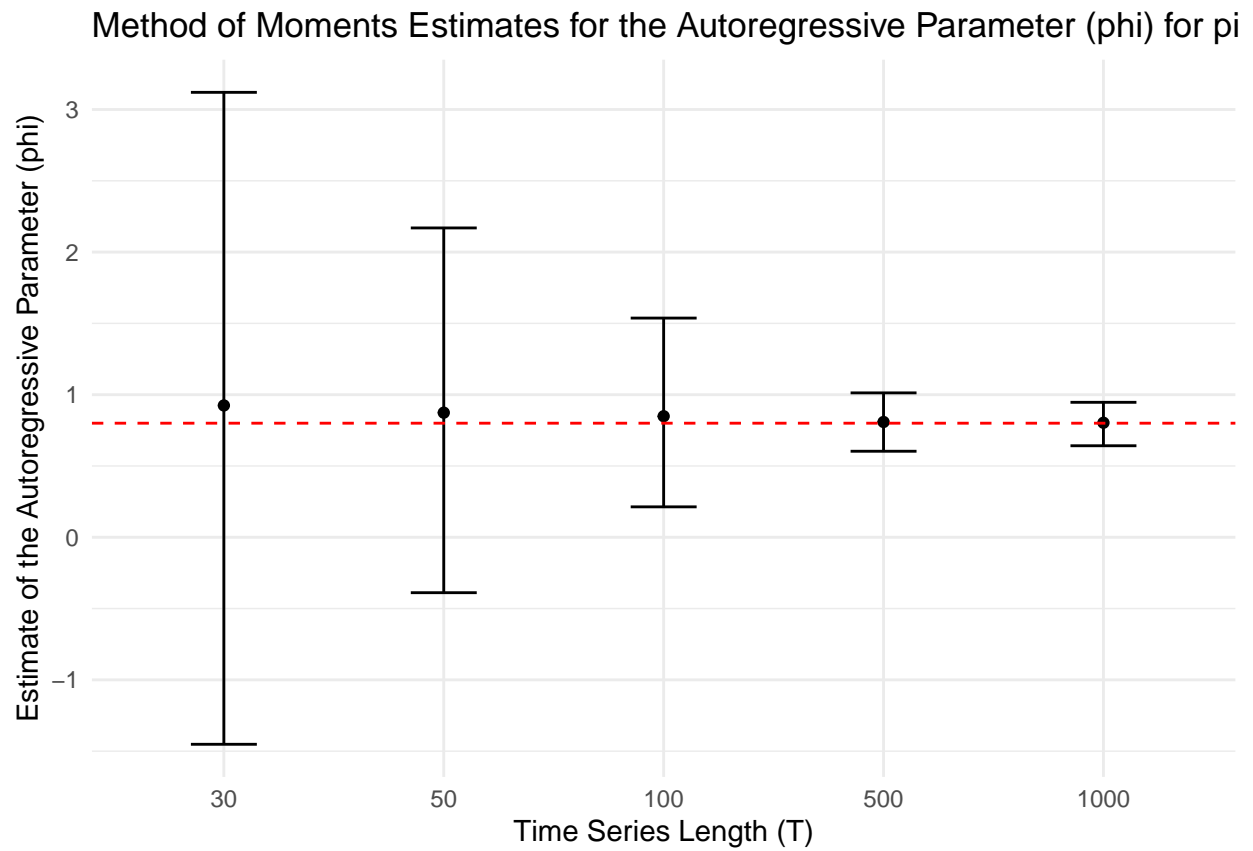


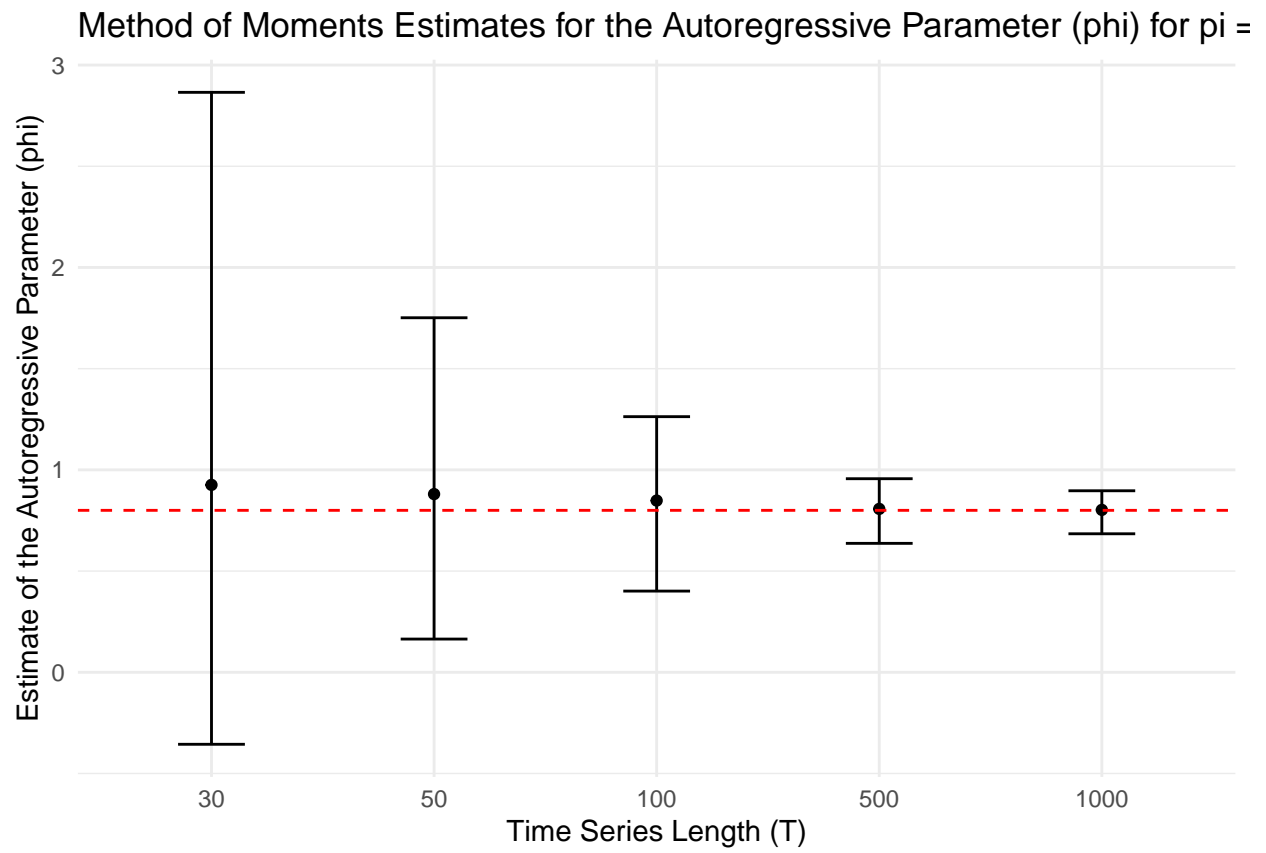


```
for (i in 1:length(phi_plots)) {  
  print(phi_plots[[i]])  
}
```

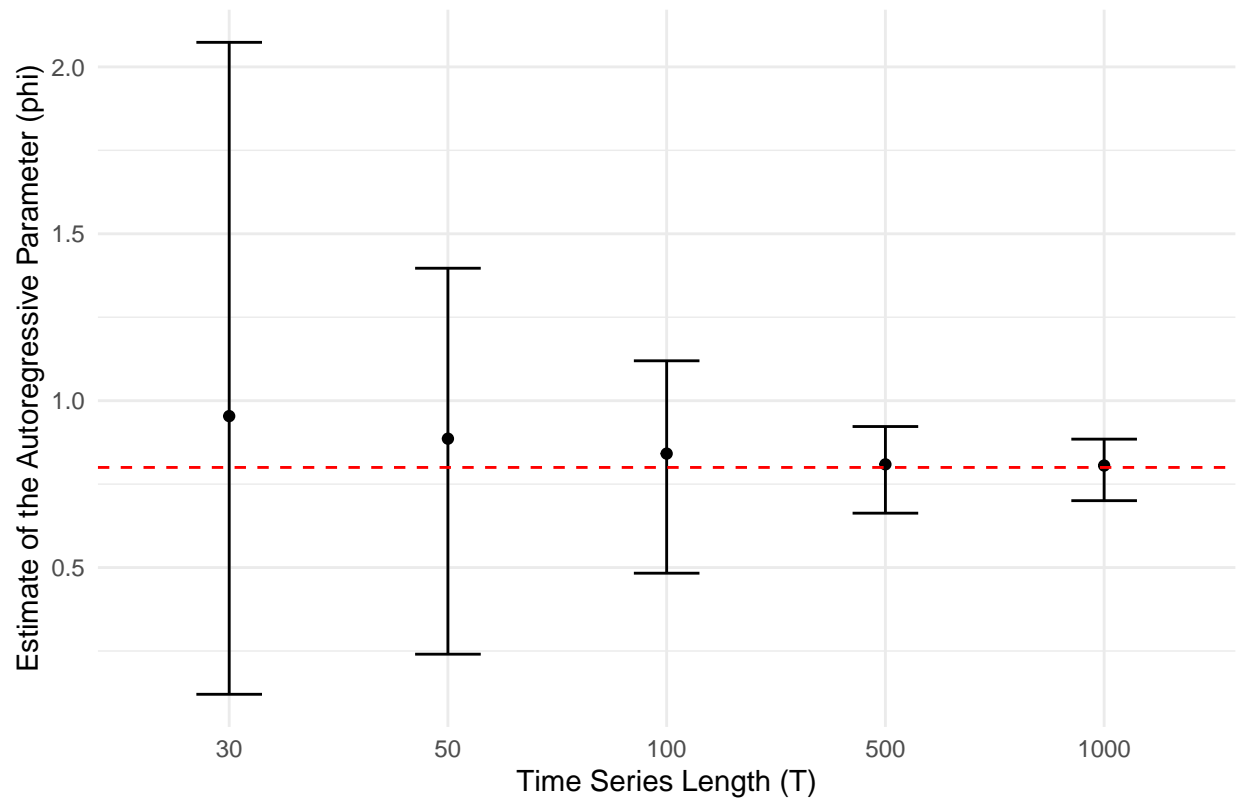




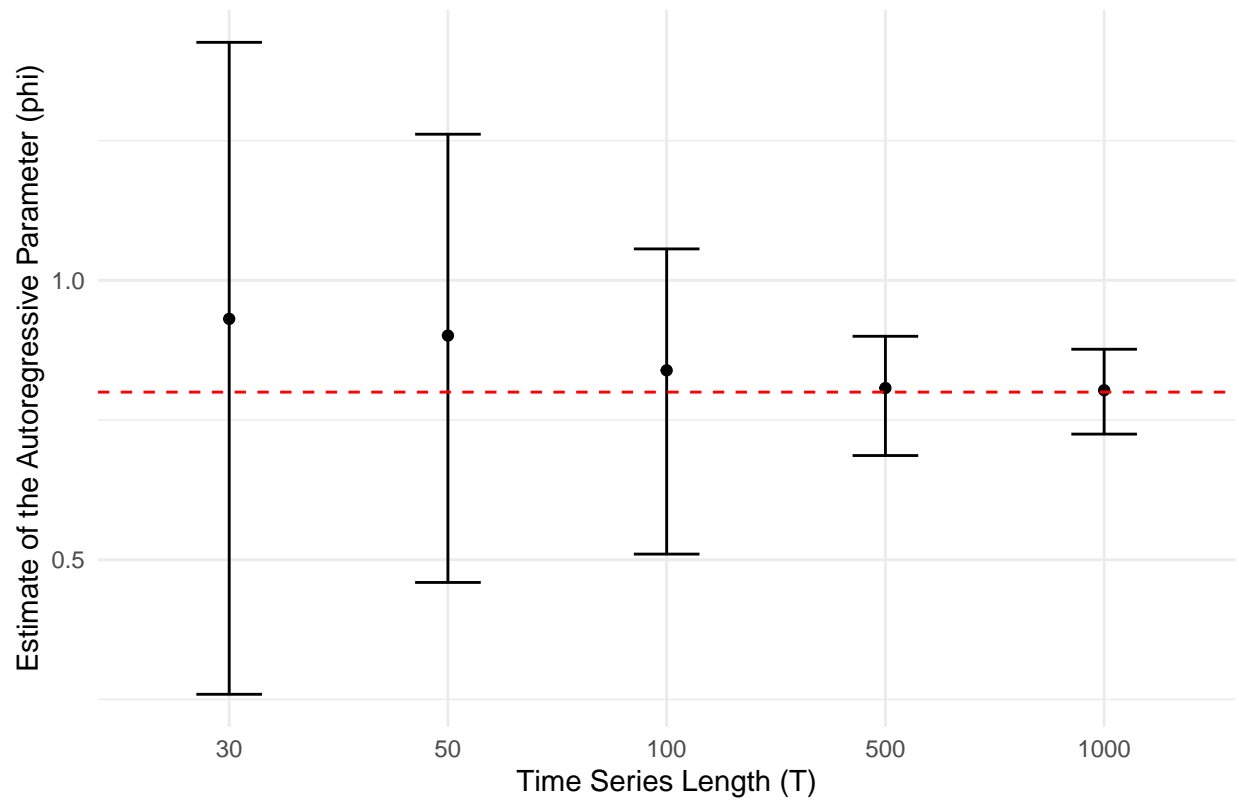


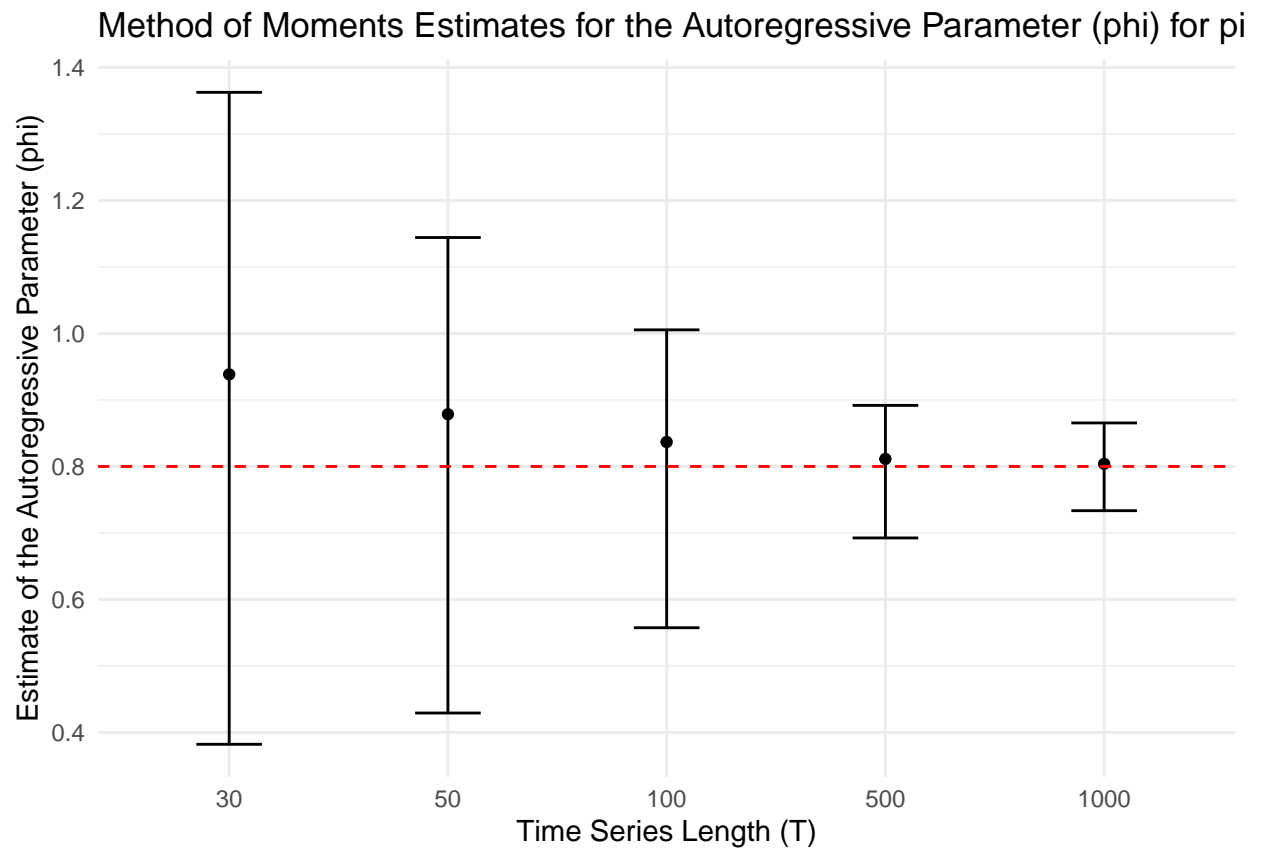


Method of Moments Estimates for the Autoregressive Parameter (ϕ) for π

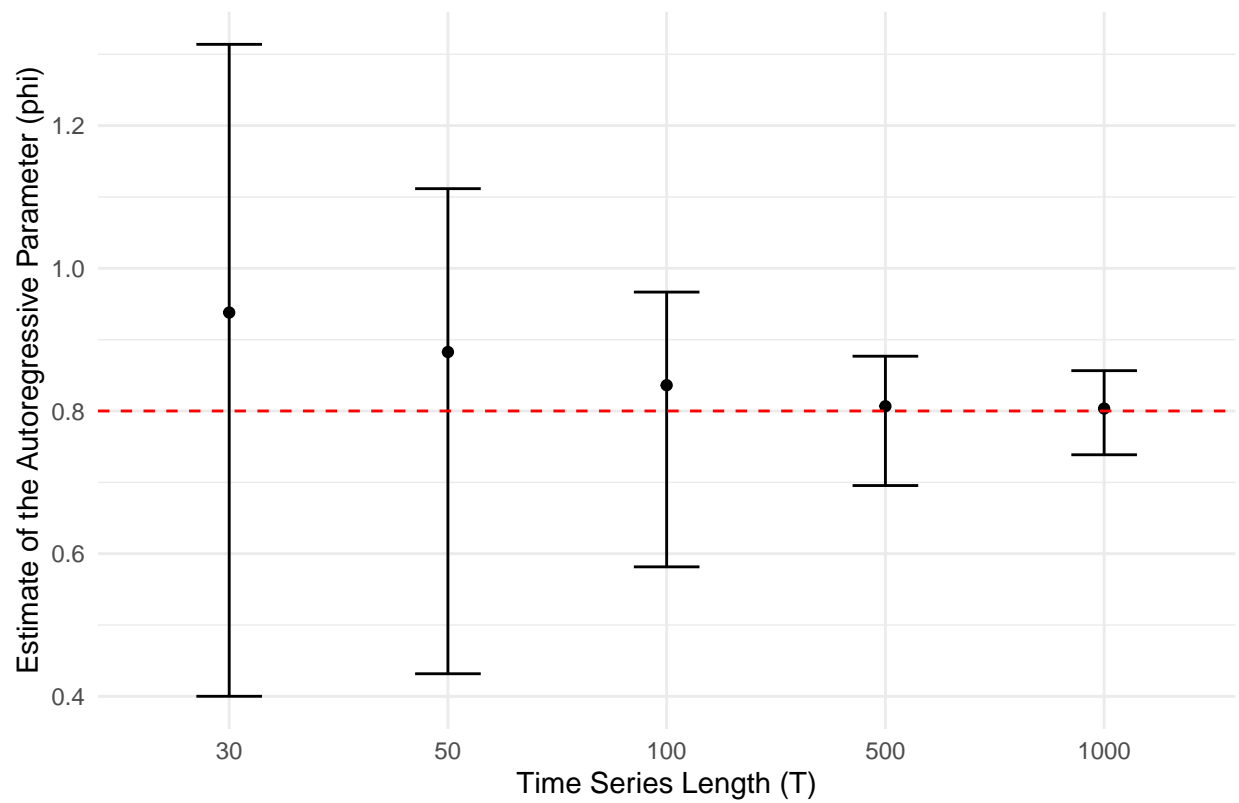


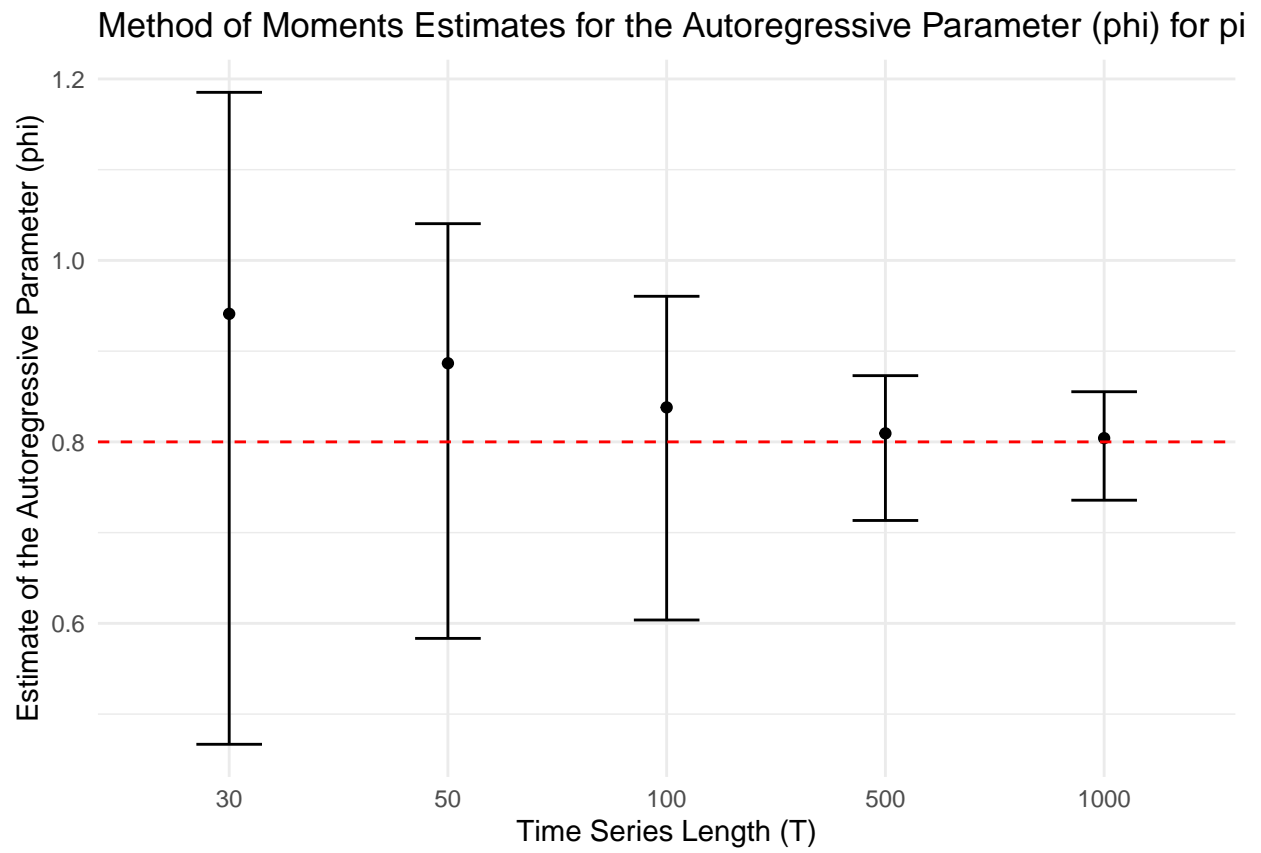
Method of Moments Estimates for the Autoregressive Parameter (ϕ) for π



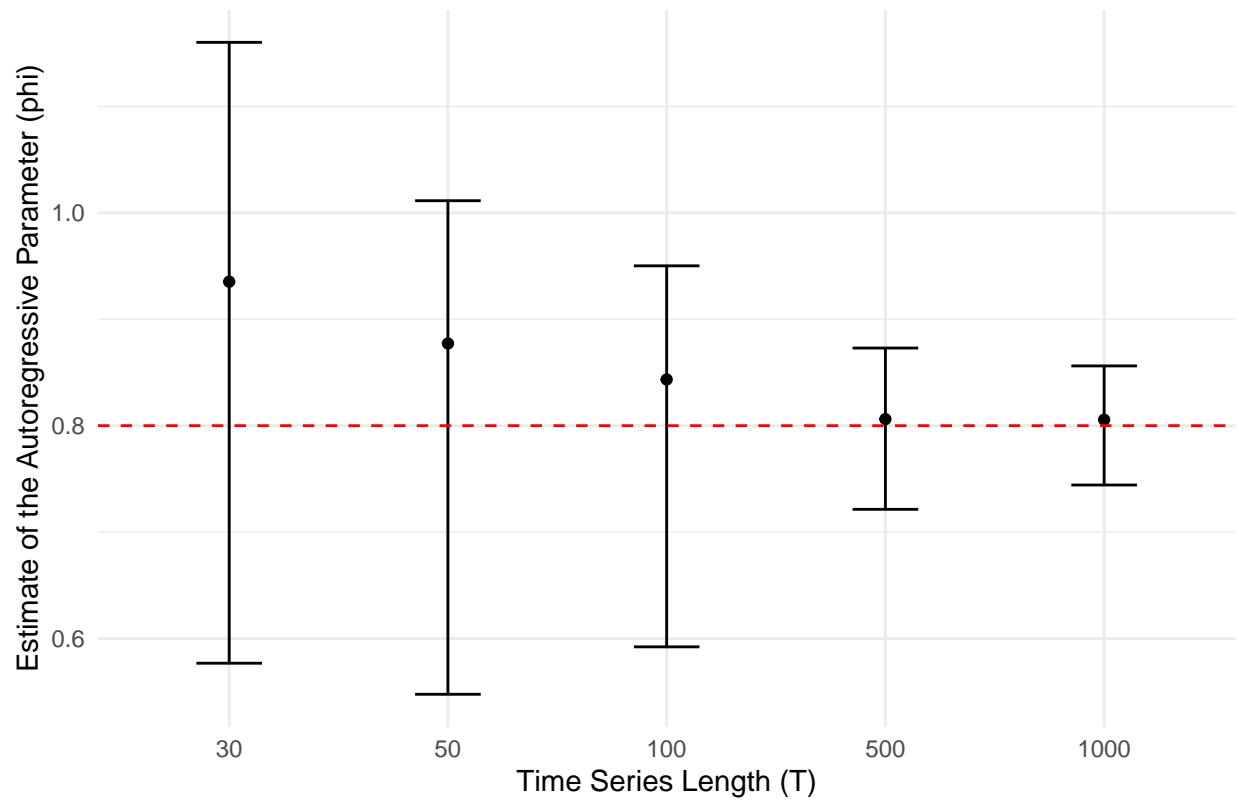


Method of Moments Estimates for the Autoregressive Parameter (ϕ) for π

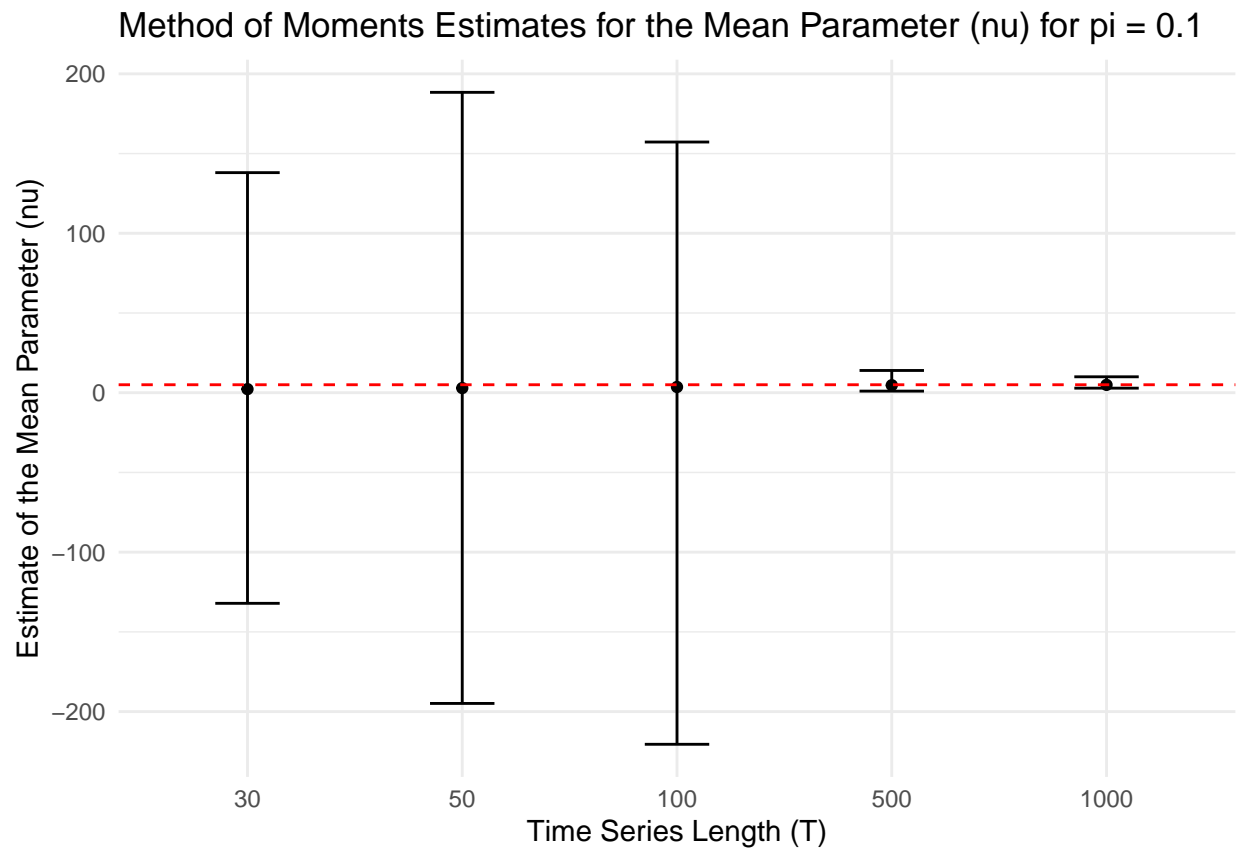


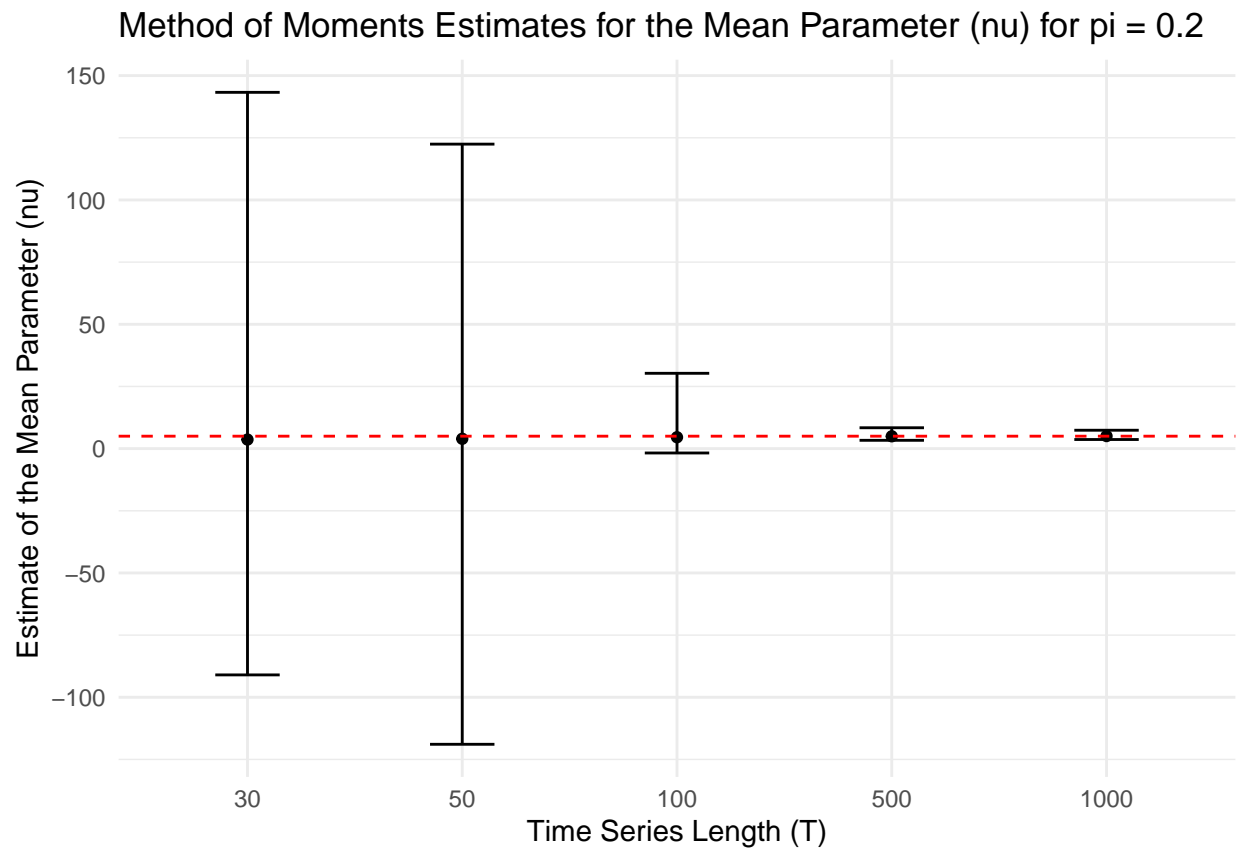


Method of Moments Estimates for the Autoregressive Parameter (ϕ) for π

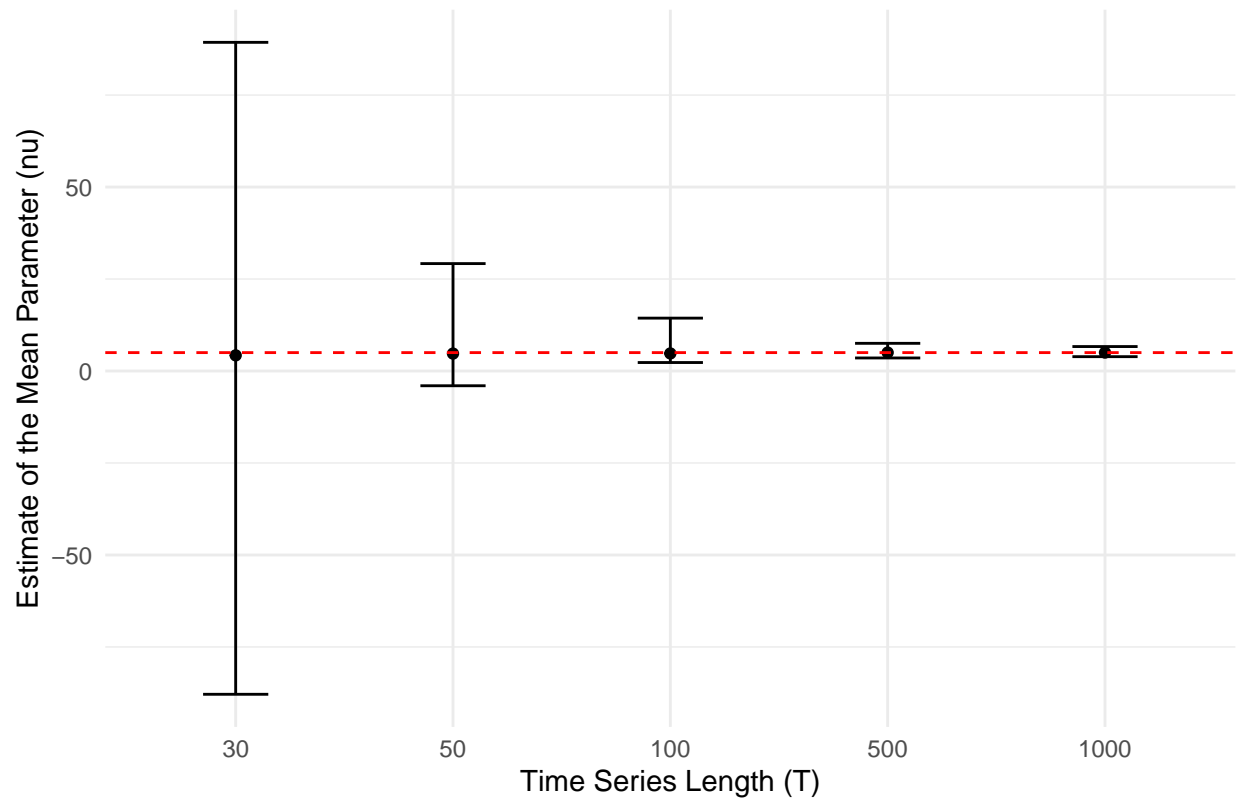


```
for (i in 1:length(nu_plots)) {  
  print(nu_plots[[i]])  
}
```

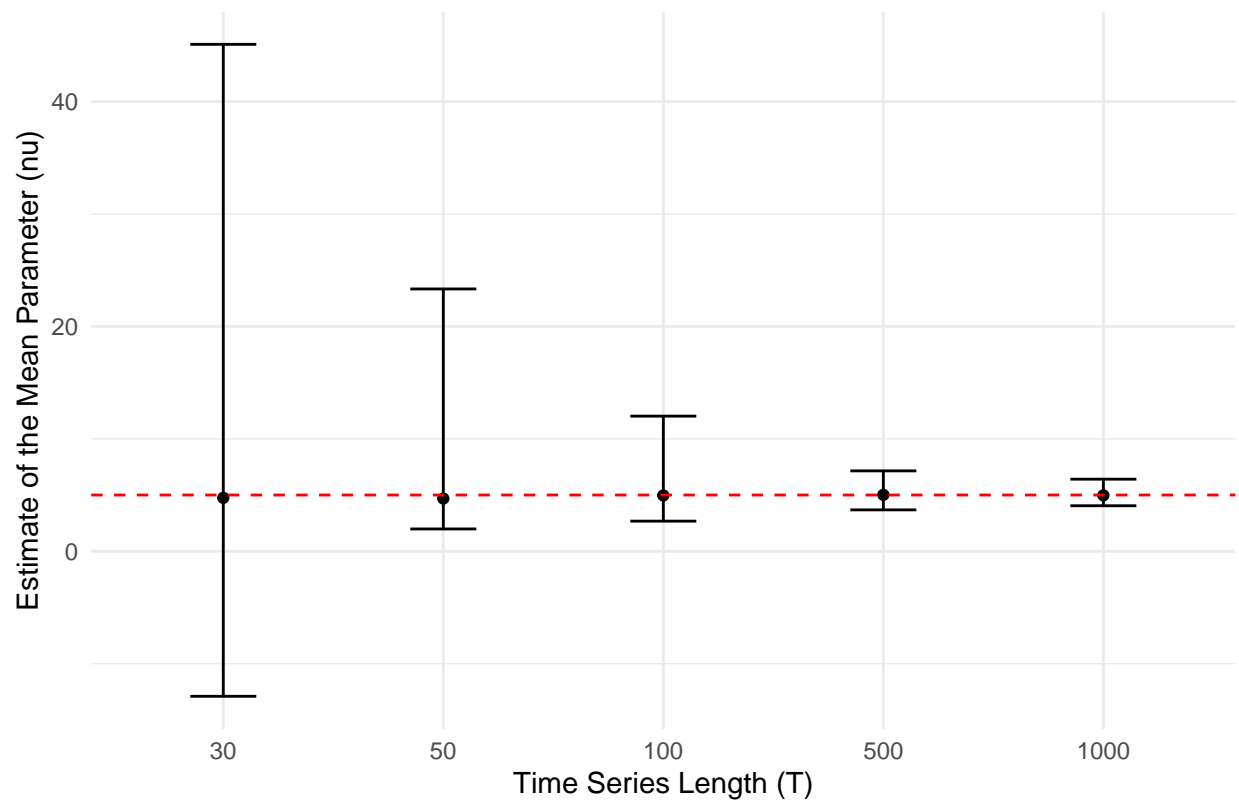




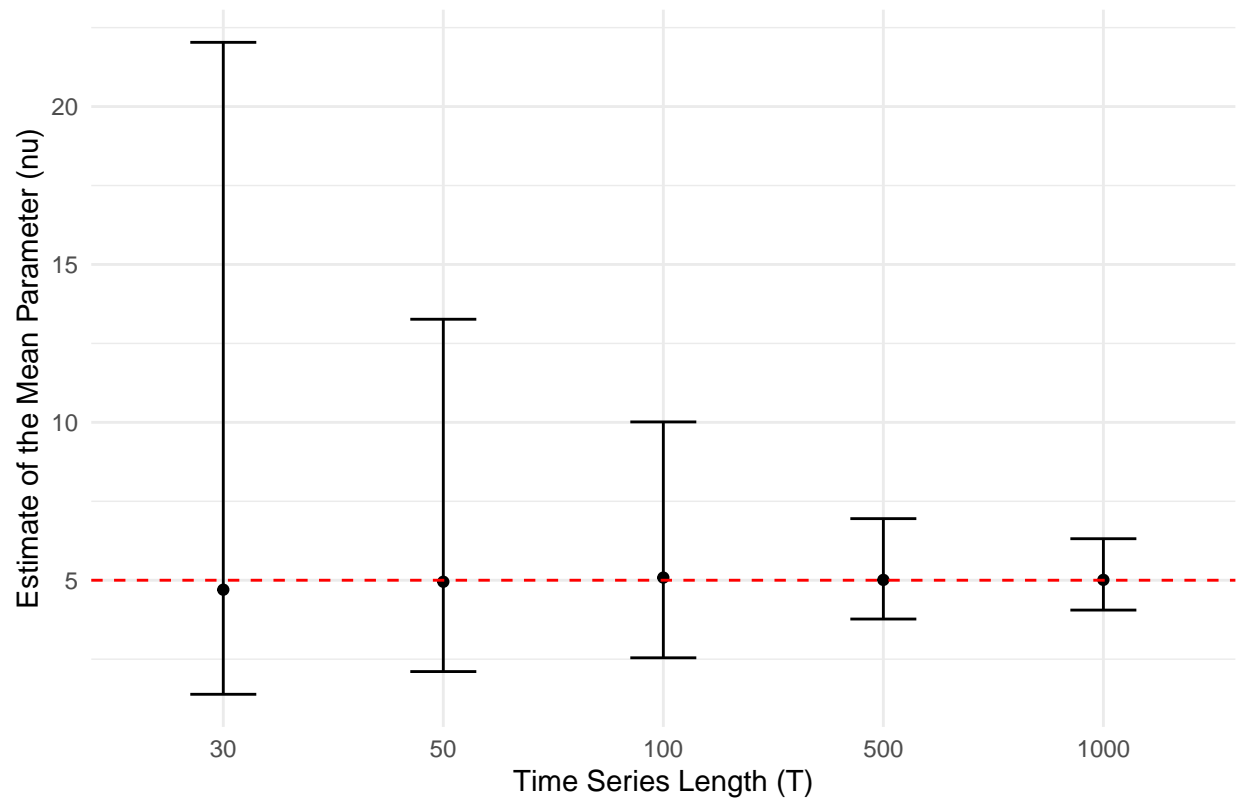
Method of Moments Estimates for the Mean Parameter (ν) for $\pi = 0.3$



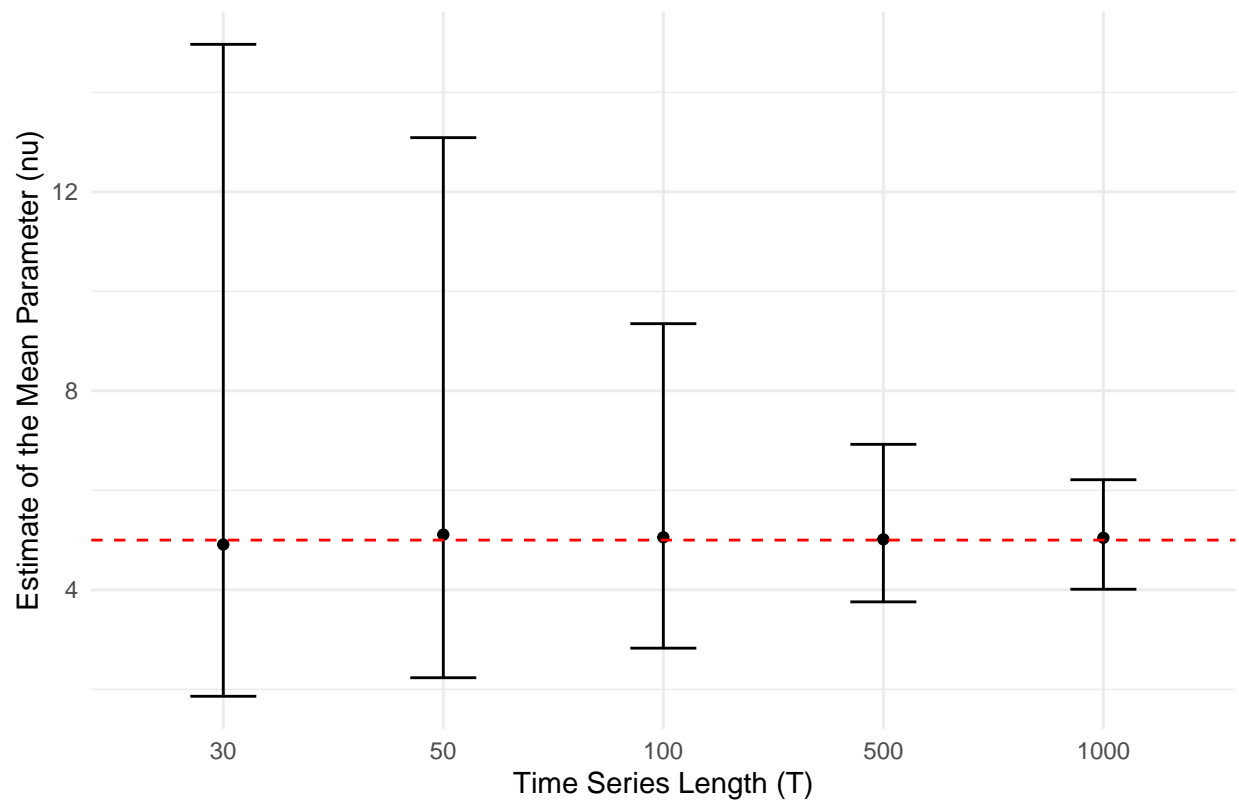
Method of Moments Estimates for the Mean Parameter (ν) for $\pi = 0.4$



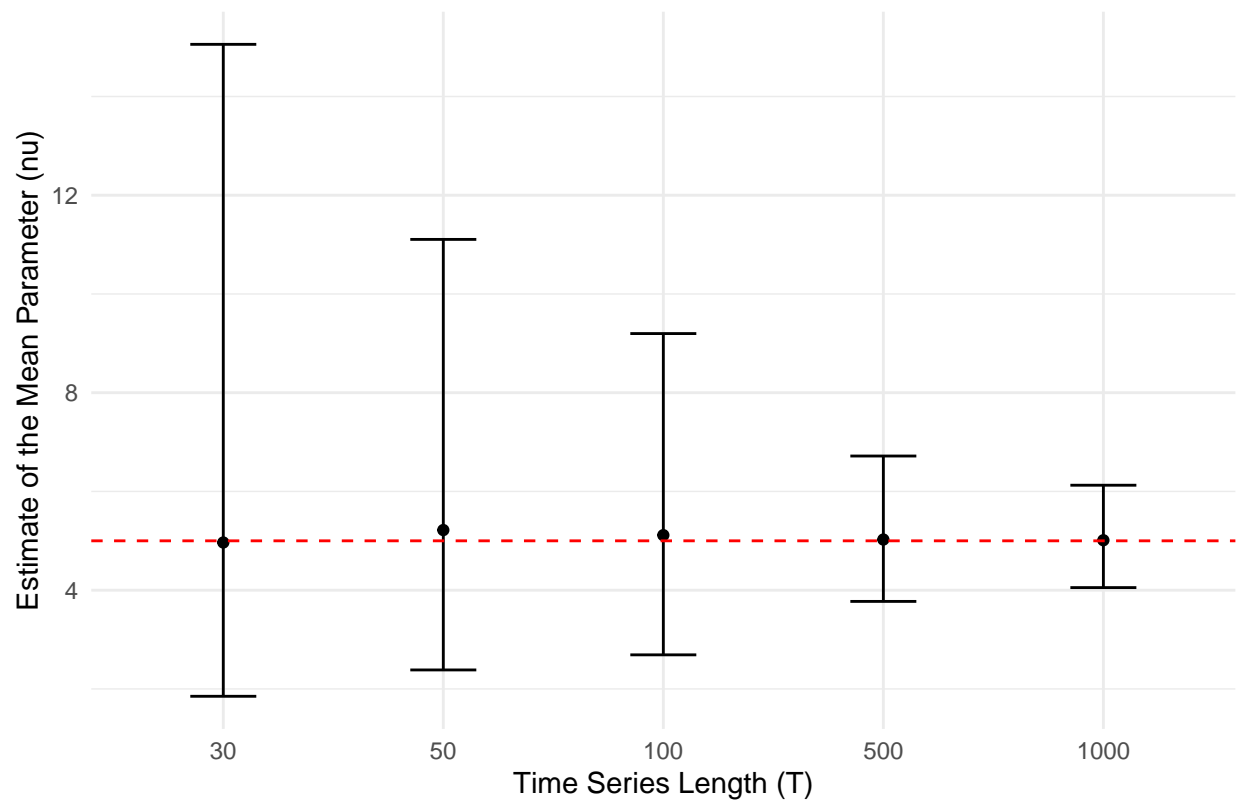
Method of Moments Estimates for the Mean Parameter (ν) for $\pi = 0.5$



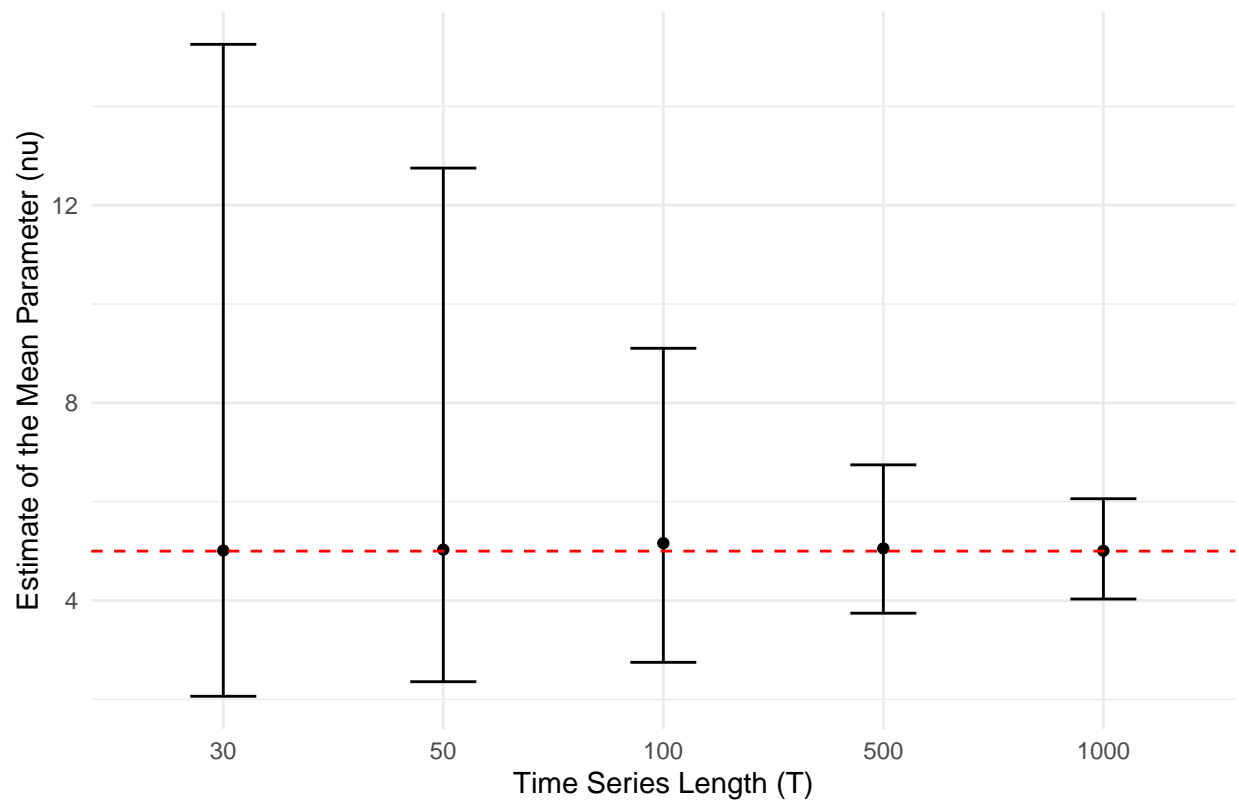
Method of Moments Estimates for the Mean Parameter (ν) for $\pi = 0.6$



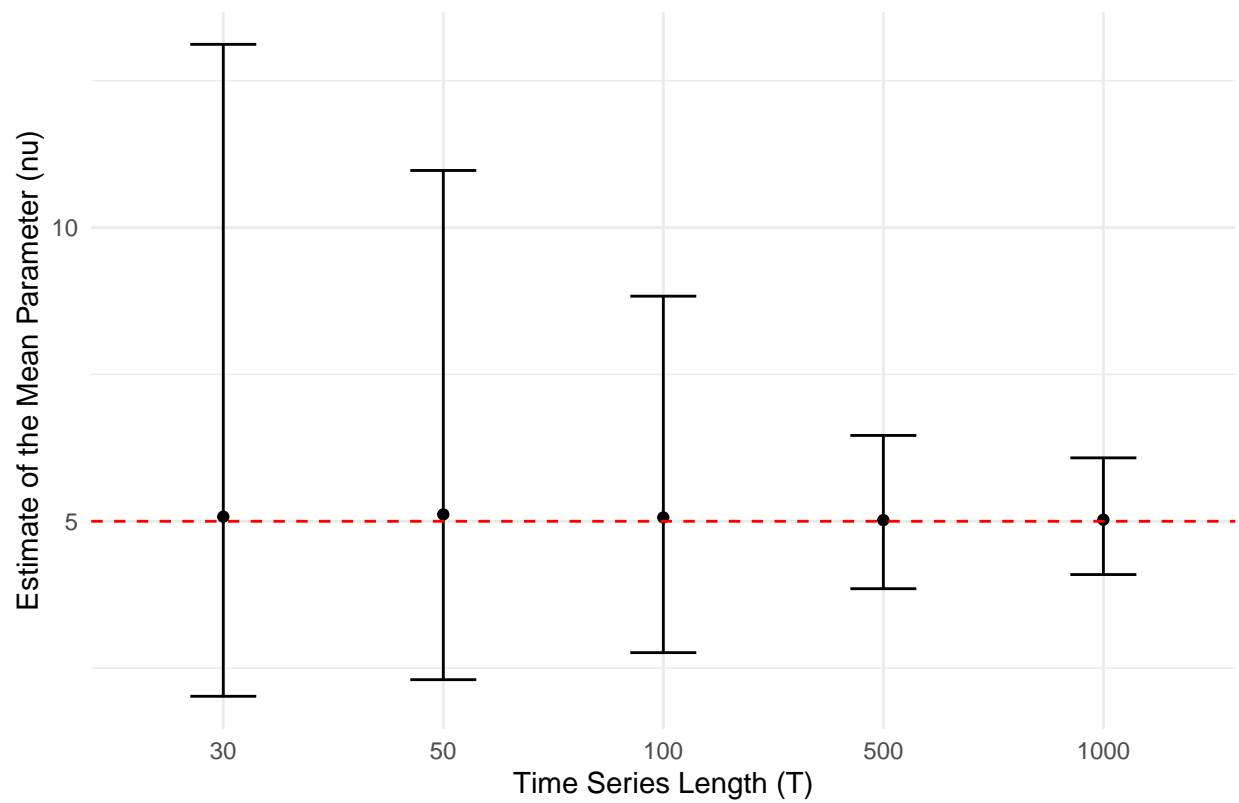
Method of Moments Estimates for the Mean Parameter (ν) for $\pi = 0.7$



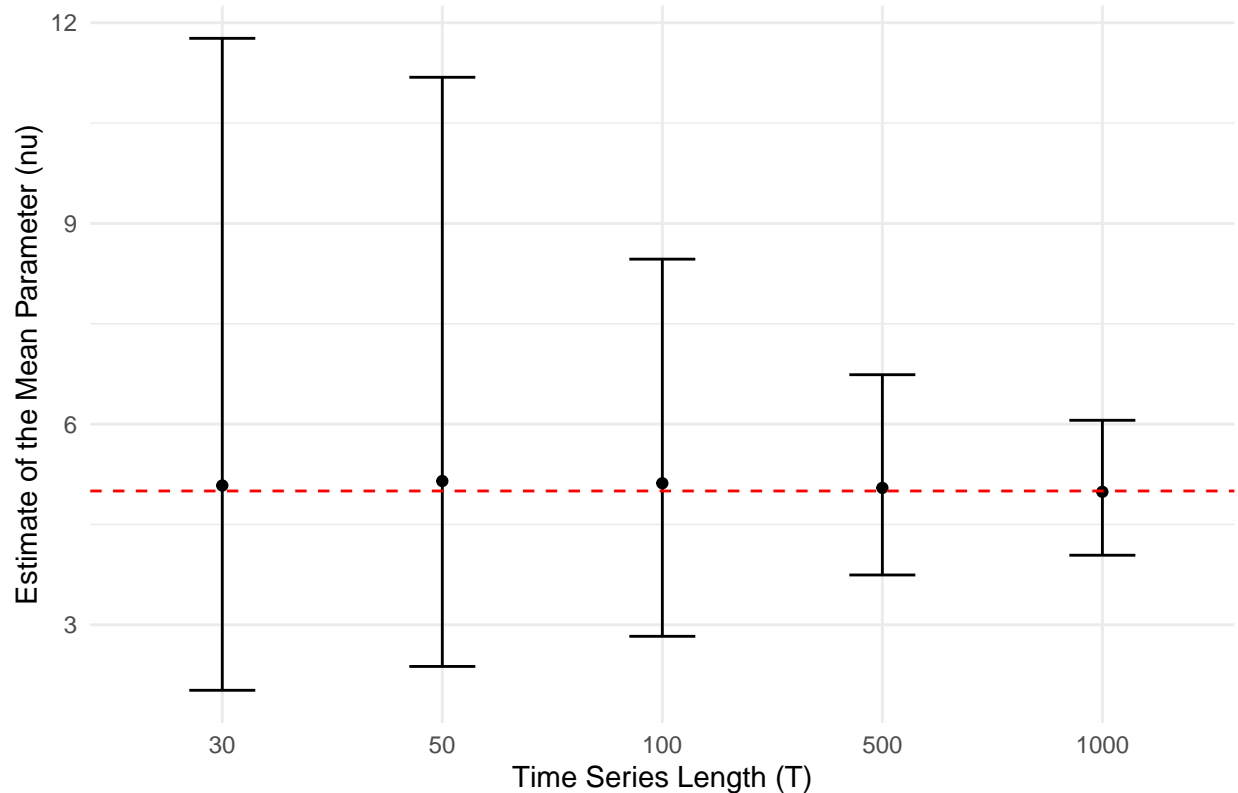
Method of Moments Estimates for the Mean Parameter (ν) for $\pi = 0.8$



Method of Moments Estimates for the Mean Parameter (ν) for $\pi = 0.9$



Method of Moments Estimates for the Mean Parameter (ν) for $\pi = 1$



```
# Load the "surveillance" package
library(surveillance)
```

```
## Loading required package: sp
```

```
## Loading required package: xtable
```

```
## This is surveillance 1.24.1; see 'package?surveillance' or
## https://surveillance.R-Forge.R-project.org/ for an overview.
```

```
# Set parameter values
phi <- 0.8 # Autoregressive parameter
nu <- 5 # Mean parameter
pi <- 0.1
T <- 1000

# Function to simulate and estimate
simulate_and_estimate <- function(pi, phi, nu, T) {
  z <- rep(0, T) # Initialize time series
  z[1] <- 5 # Initialize Z_1 at 5

  # Simulate time series
  for (i in 2:T) {
    z[i] <- rpois(1, lambda = nu + phi * z[i-1])
  }
}
```

```

}

# Thin the time series using binomial thinning
y <- rbinom(T, z, pi)

# Check for missing or NaN values
if (any(is.na(y)) || any(is.nan(y))) {
  return(c(NA, NA))
}

# Convert y to "sts" object
sts_object <- sts(y)

# Specify the "endemic" and "epidemic" components using control parameter
hhh_result <- hhh4(sts_object, control = list(end = list(f = ~ 1), ar = list(f = ~1),
  family = "Poisson"))

# Get parameter estimates and standard errors using summary()
hhh_summary <- summary(hhh_result)
phi_hat <- exp(hhh_summary$fixef["ar.1", "Estimate"])
nu_hat <- exp(hhh_summary$fixef["end.1", "Estimate"])

# Get standard errors
phi_se <- hhh_summary$fixef["ar.1", "Std. Error"]
nu_se <- hhh_summary$fixef["end.1", "Std. Error"]

return(list(phi_hat = phi_hat, nu_hat = nu_hat, phi_se = phi_se, nu_se = nu_se))
}

# We will run the above function for different values of pi, and plot the estimates of nu vs pi, and ph
# Knit the Markdown file

estimates <- simulate_and_estimate(pi = pi, phi = phi, nu = nu, T = 1000)

phi_estimates = c()
nu_estimates = c()

phi_estimates[1] = estimates$phi_hat
nu_estimates[1] = estimates$nu_hat

#exp(phi_hat)
#exp(nu_hat)

# Print the estimates
cat("Estimates for pi =", pi, "=> phi_hat =", estimates$phi_hat, " nu_hat =", estimates$nu_hat, "\n")

## Estimates for pi = 0.1 => phi_hat = 0.1839054 nu_hat = 2.113534

cat("Standard Errors => phi_se =", estimates$phi_se, " nu_se =", estimates$nu_se, "\n")

## Standard Errors => phi_se = 0.1625545 nu_se = 0.04132217

```

```

## Estimates for pi = 0.2 => phi_hat = 0.2961915  nu_hat = 3.719902

## Standard Errors => phi_se = 0.1028398  nu_se = 0.04498947

## Estimates for pi = 0.3 => phi_hat = 0.4358681  nu_hat = 4.215075

## Standard Errors => phi_se = 0.05919022  nu_se = 0.04628825

## Estimates for pi = 0.4 => phi_hat = 0.4730341  nu_hat = 5.240797

## Standard Errors => phi_se = 0.05390644  nu_se = 0.04852893

## Estimates for pi = 0.5 => phi_hat = 0.5904351  nu_hat = 4.999223

## Standard Errors => phi_se = 0.03838206  nu_se = 0.05443087

## Estimates for pi = 0.6 => phi_hat = 0.6314585  nu_hat = 5.755158

## Standard Errors => phi_se = 0.03383671  nu_se = 0.05680721

## Estimates for pi = 0.7 => phi_hat = 0.7314786  nu_hat = 4.645519

## Standard Errors => phi_se = 0.02790858  nu_se = 0.07342935

## Estimates for pi = 0.8 => phi_hat = 0.7171818  nu_hat = 6.139962

## Standard Errors => phi_se = 0.03020383  nu_se = 0.07486584

## Estimates for pi = 0.9 => phi_hat = 0.7896081  nu_hat = 4.823443

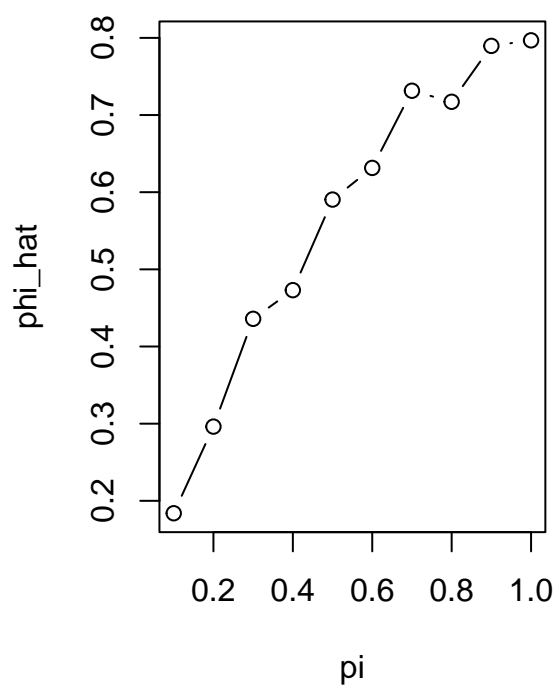
## Standard Errors => phi_se = 0.02489187  nu_se = 0.08992725

## Estimates for pi = 1 => phi_hat = 0.7967919  nu_hat = 4.927901

## Standard Errors => phi_se = 0.02381891  nu_se = 0.08975039

```

Estimates of ϕ vs. π



Estimates of ν vs. π

