

STAT 443: Homework 4

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Question 1

(a & b)

```
knitr::include_graphics("q1.jpg")
```

1. Consider the AR(2) process from Assignment 2 (Q1):

$$X_t = 0.1X_{t-1} + 0.2X_{t-2} + Z_t, \quad \{Z_t\}_{t \in \mathbb{Z}} \sim WN(0, \sigma^2),$$

and recall that its autocorrelation function has the form

$$\rho(h) = \frac{15}{36}(-0.4)^{|h|} + \frac{21}{36}(0.5)^{|h|}, \quad h \in \mathbb{Z}.$$

- (a) Derive the normalized spectral density function of $\{X_t\}_{t \in \mathbb{Z}}$.
- (b) Write down the (power) spectral density function of $\{X_t\}_{t \in \mathbb{Z}}$.
- (c) Plot the spectral density and comment on its behaviour.

$$\sum_{h=-\infty}^{\infty} \rho(h) \cos(h\omega) = \frac{r^2(\cos(\omega) - r^2)}{(1 - 2r\cos(\omega) + r^4)\pi}$$

$$\begin{aligned} \text{a) Given } f^*(\omega) &= \frac{1}{\pi} \left(1 + 2 \sum_{k=1}^{\infty} \rho(k) \cos(k\omega) \right) \\ &= \frac{1}{\pi} \left[1 + 2 \sum_{k=1}^{\infty} \left(\frac{15}{36}(-0.4)^k + \frac{21}{36}(0.5)^k \right) \cos(k\omega) \right] \\ &= \frac{1}{\pi} \left(1 + \frac{5}{6} \sum_{k=1}^{\infty} (-0.4)^k \cos(k\omega) + \frac{7}{6} \sum_{k=1}^{\infty} (0.5)^k \cos(k\omega) \right) \\ &= \frac{1}{\pi} \left(1 + \frac{5}{6} \left(\frac{-0.4(\cos(\omega) - 0.16)}{(1 - 0.8\cos(\omega) + 0.16)} \right) + \frac{7}{6} \left(\frac{0.5(\cos(\omega) - 0.25)}{(1 - \cos(\omega) + 0.25)} \right) \right) \end{aligned}$$

$$\text{b) } \sigma_X^2 = \text{Var}(X_t)$$

Using Yule-Walker's Equation:

$$\gamma(0) = 0.1\gamma(1) + 0.2\gamma(2) + \sigma^2 \quad (1)$$

$$\gamma(1) = 0.1\gamma(0) + 0.2\gamma(1) \quad (2)$$

$$\gamma(2) = 0.1\gamma(1) + 0.2\gamma(0) \quad (3)$$

$$\begin{aligned} \text{In (2): } 0.8\gamma(1) &= 0.1\gamma(0) & \text{In (3) \& (2): } \gamma(2) &= 0.1(0.125\gamma(0)) + 0.2\gamma(0) \\ \gamma(1) &= 0.125\gamma(0) & \gamma(2) &= 0.2125\gamma(0) \end{aligned}$$

$$\begin{aligned} \text{In (1), } \gamma(0) &= 0.1(0.125\gamma(0)) + 0.2(0.2125\gamma(0)) + \sigma^2 \\ \gamma(0) &= \frac{\sigma^2}{(1 - 0.1 \times 0.125 + 0.2 \times 0.2125)} = \frac{\sigma^2}{0.945} \end{aligned}$$

$$\therefore f^*(\omega) = \frac{f(\omega)}{\sigma_X^2}$$

$$\begin{aligned} f(\omega) &= \sigma_X^2 f^*(\omega) = \frac{\sigma_X^2}{\pi} \left(1 + \frac{5}{6} \left(\frac{-0.4(\cos(\omega) - 0.16)}{(1 - 0.8\cos(\omega) + 0.16)} \right) + \frac{7}{6} \left(\frac{0.5(\cos(\omega) - 0.25)}{(1 - \cos(\omega) + 0.25)} \right) \right) \\ &= \frac{1}{0.945\pi} \left(1 + \frac{5}{6} \left(\frac{-0.4(\cos(\omega) - 0.16)}{(1 - 0.8\cos(\omega) + 0.16)} \right) + \frac{7}{6} \left(\frac{0.5(\cos(\omega) - 0.25)}{(1 - \cos(\omega) + 0.25)} \right) \right) \end{aligned}$$

(c)

```
# Calculate the spectral density
spectral_density <- function(omega) {
  term2 <- (15/18) * ((-0.4*cos(omega) - 0.16) / (1.16 + 0.8 * cos(omega)))
  term3 <- (21/18) * ((0.5*cos(omega) - 0.25) / (1.25 - cos(omega)))

  return((1/(0.95*pi)) * (1 + term2 + term3))
}

# Create a sequence of frequencies from 0 to pi
omega <- seq(0, pi, length.out = 1000)
```

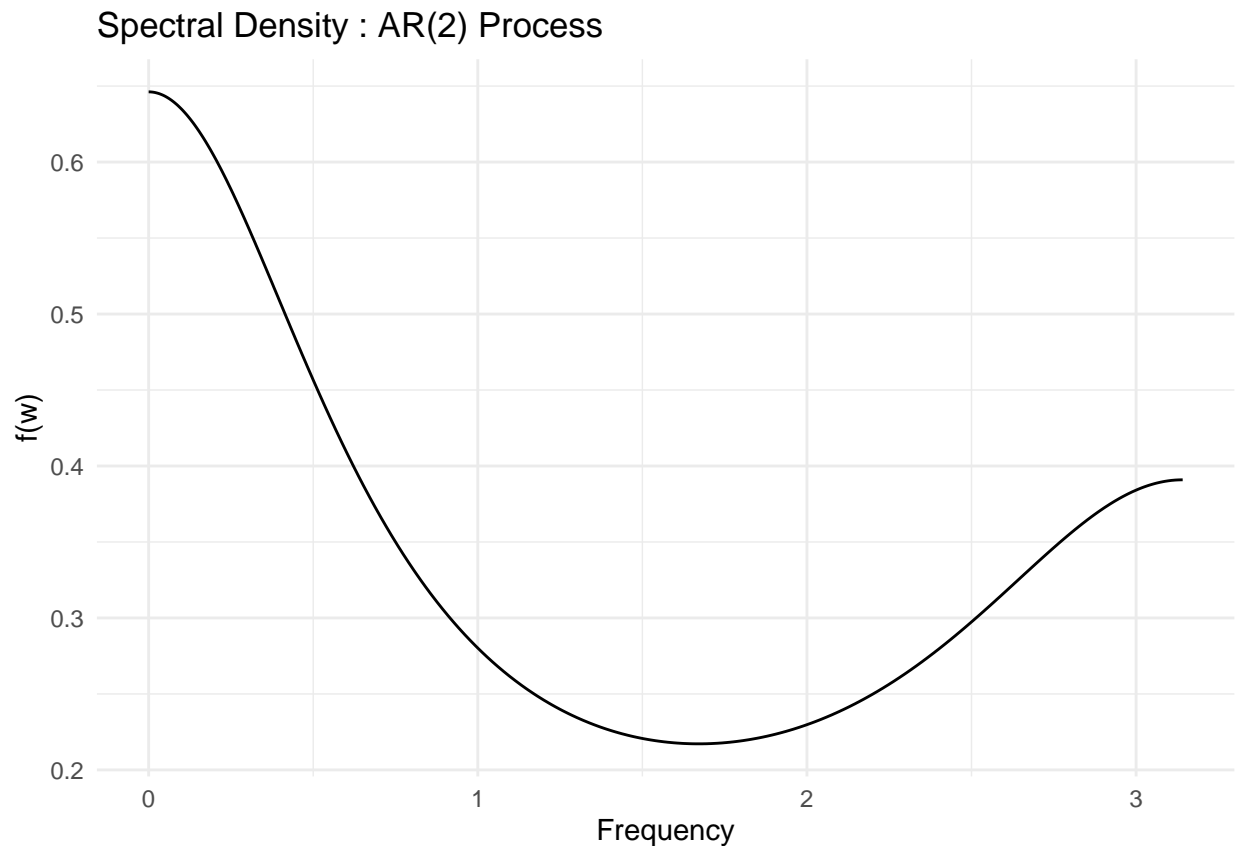
```

# Calculate the spectral density for each frequency
spec_values <- sapply(omega, spectral_density)

# Create a data frame for plotting
spec_df <- data.frame(
  omega = omega,
  frequency = omega/(2*pi),
  spectral_density = spec_values
)

# Plot using ggplot2
ggplot(spec_df, aes(x = omega, y = spec_values)) +
  geom_line() +
  labs(title = "Spectral Density : AR(2) Process",
       x = "Frequency",
       y = "f(w)") +
  theme_minimal()

```



We can see the spectral density being dominated by low frequencies, with the spectral density is at highest at frequency = 0 as it's peak. As we move further away from frequency 0, we observe a dip / decreasing spectral density and another smaller peak at approximately frequency 3. Overall, this suggests that our AR(2) process has long term dependencies.

Question 2

```
knitr::include_graphics("q2.jpg")
```

2. Given the power spectral density function

$$f(\omega) = \frac{1}{\pi} (1.1 + 0.54 \cos(\omega) - 0.2 \cos(2\omega)), \quad \omega \in (0, \pi),$$

compute the autocovariance function $\gamma(k)$ of the underlying stochastic process for $k = 0, 1, 2$ and $k > 2$.

$$\text{Given } f(\omega) = \frac{1}{\pi} (\gamma(0) + 2 \sum_{k=1}^{\infty} \gamma(k) \cos(k\omega)) = \frac{1}{\pi} (\gamma(0) + 2\gamma(1)\cos(\omega) + 2\gamma(2)\cos(2\omega))$$

\therefore

$$\gamma(k) = \begin{cases} 1.1 & \text{if } k=0 \\ 0.27 & \text{if } k=1 \\ -0.1 & \text{if } k=2 \\ 0 & \text{if } k>2 \end{cases}$$

$2\gamma(1) = 0.54$
 $\gamma(1) = 0.27$
 $2\gamma(2) = -0.2$
 $\gamma(2) = -0.1$

Question 3

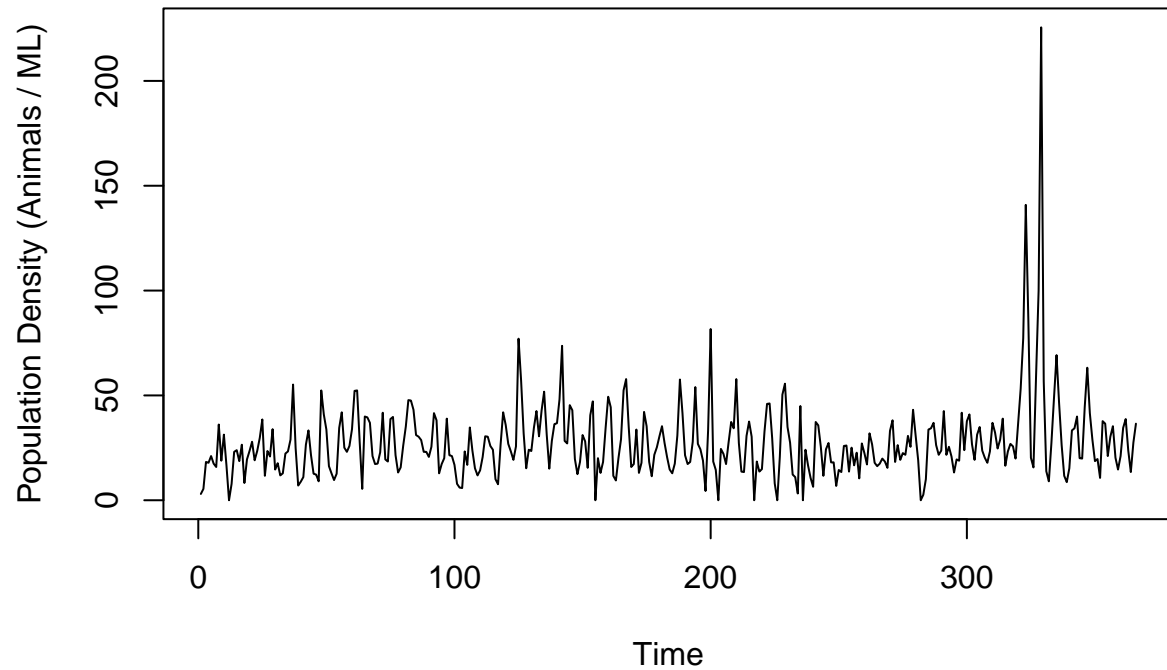
(a)

```
predator <- read.csv("predator_preys.csv")

rot_ts <- ts(predator$rotifers..animals.ml.)

plot(rot_ts,
     main = "Rotifer Population Over Time",
     ylab = "Population Density (Animals / ML)"
)
```

Rotifer Population Over Time

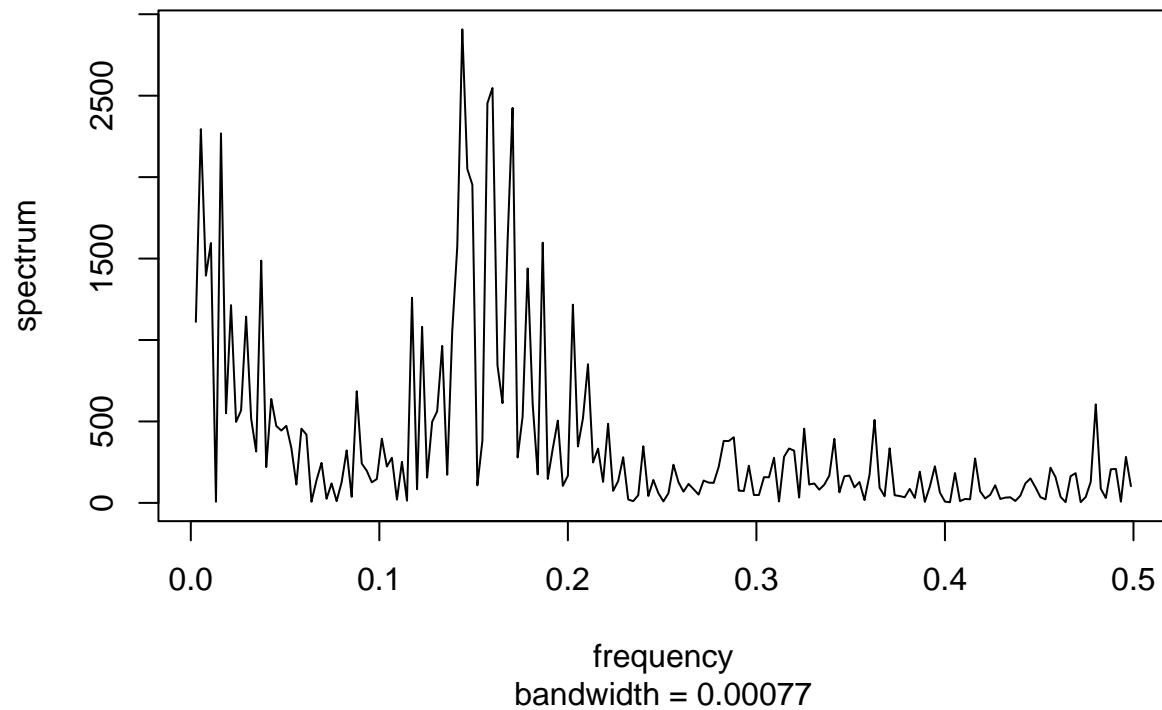


We observe a consistent fluctuation until at time ~350 where we observe a large spike / peak before dropping back to the consistent fluctuation.

(b)

```
raw_spec <- spectrum(rot_ts,  
                      log = "no",  
                      main = "Raw Periodogram of Rotifer Time Series")
```

Raw Periodogram of Rotifer Time Series



```
frequencies <- raw_spec$freq
spectrum_values <- raw_spec$spec

# Get the max
max_index <- which.max(spectrum_values)

dominant_freq <- frequencies[max_index]
dominant_freq
```

```
## [1] 0.144
```

```
angular_freq <- 2 * pi * dominant_freq
angular_freq
```

```
## [1] 0.9047787
```

```
wavelength_days <- 1 / dominant_freq
wavelength_days
```

```
## [1] 6.944444
```

For our dominating frequency 0.144, we get wavelength (days) ~ 6.944 and angular frequency ~ 0.905

(c)

```
N <- length(rot_ts)

num_cycles <- N / wavelength_days
num_cycles
```

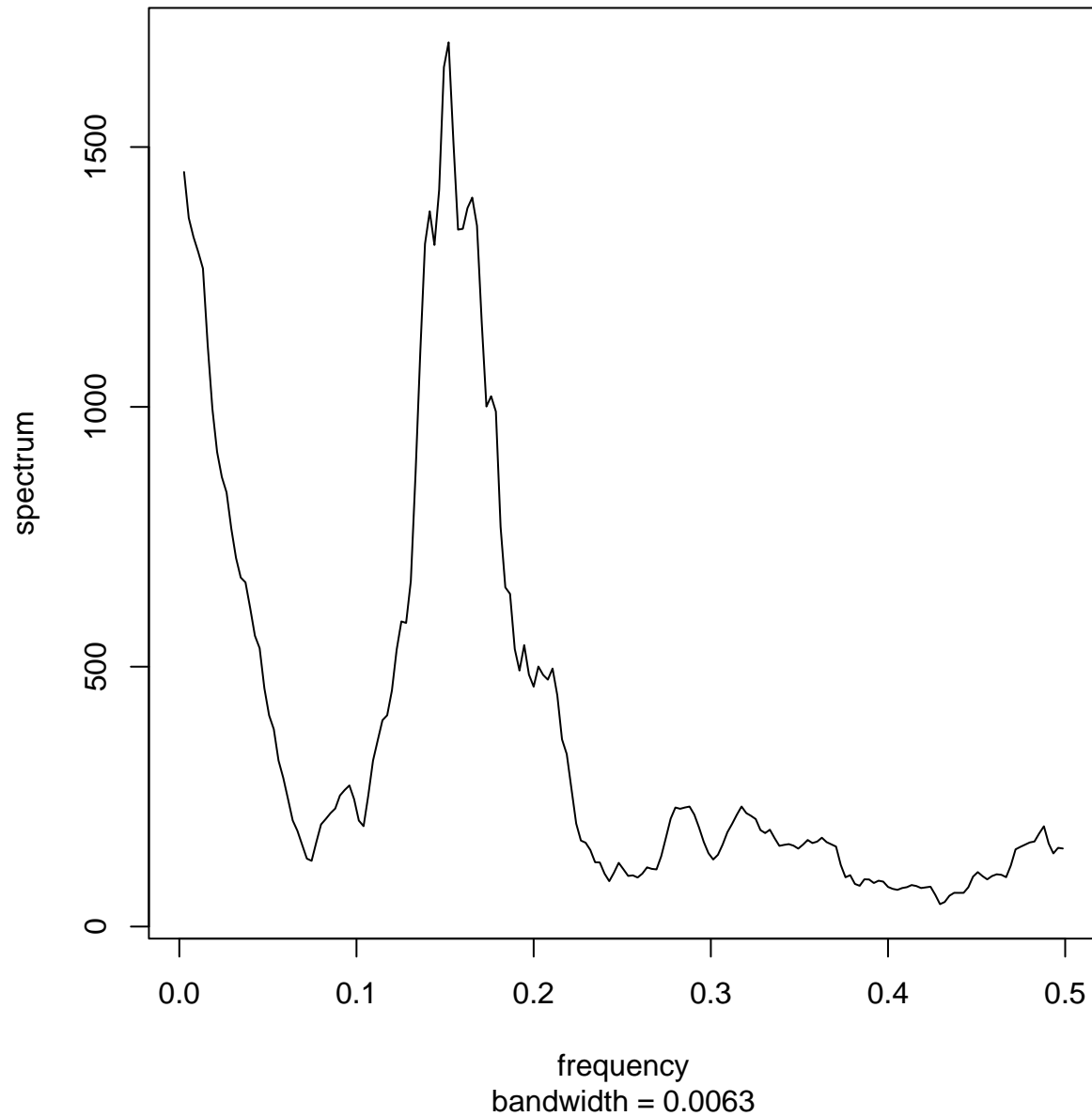
```
## [1] 52.704
```

On a cycle length of approximately 7 days, we get the predator prey cycles to be approximately 52.7

(d)

```
spec_smooth <- spec.pgram(rot_ts,
  spans = c(366 / 40), # As per piazza response @116
  log = "no",
  main = "Smoothed Periodogram Rotifer Population")
```

Smoothed Periodogram Rotifer Population



We may want to change the raw periodogram since we know it is not a consistent estimator of the spectral density function and change it to an estimator that is more consistent. As per piazza response @116, I chose $N / 40$ over $2 * \sqrt{N}$ for my span smoothing because I believe it includes several peaks and dips relatively better.

(e)

```
N <- length(rot_ts) # 366  
  
omega_59 <- (2 * pi * 59) / N  
omega_59
```



```
## [1] 1.012863
```

Our Fourier Frequency for $p = 59$ is 1.012863.

(f)

```
N <- length(rot_ts) # 366

# Create an empty data frame to store significant frequencies
significant_freqs <- data.frame(
  p = integer(),
  freq = numeric(),
  angular_freq = numeric(),
  wavelength = numeric(),
  F_stat = numeric(),
  p_value = numeric()
)

# Loop through potential frequencies (Fourier components)
for (p in 1:(N/2)) {
  t <- 1:N # Time vector
  omega_p <- 2 * pi * p / N # angular frequency

  # Fit the linear model
  model <- lm(rot_ts ~ cos(omega_p * t) + sin(omega_p * t))

  # Extract F-statistic and p-value from model summary
  model_summary <- summary(model)
  f_stat <- model_summary$fstatistic[1]
  p_value <- pf(f_stat,
                model_summary$fstatistic[2],
                model_summary$fstatistic[3],
                lower.tail = FALSE)

  # Store results if significant at 99% confidence level (p < 0.01)
  if (p_value < 0.01) {
    significant_freqs <- rbind(significant_freqs, data.frame(
      p = p,
      freq = p / N,
      angular_freq = omega_p,
      wavelength = N / p,
      F_stat = f_stat,
      p_value = p_value
    ))
  }
}

# Sort significant frequencies by F-statistic (DESC)
significant_freqs <- significant_freqs[order(-significant_freqs$F_stat), ]

print(significant_freqs)
```

```
##           p           freq angular_freq wavelength  F_stat      p_value
```

```
## value 2 0.005464481 0.03433435 183.000000 9.261036 0.0001194628
## value5 59 0.161202186 1.01286321 6.203390 8.744184 0.0001954750
## value2 4 0.010928962 0.06866869 91.500000 6.947354 0.0010942468
## value4 54 0.147540984 0.92702734 6.777778 6.495157 0.0016923460
## value3 52 0.142076503 0.89269299 7.038462 6.255040 0.0021341862
## value6 63 0.172131148 1.08153190 5.809524 6.178262 0.0022986471
## value1 3 0.008196721 0.05150152 122.000000 5.921789 0.0029462120
```

(g)

```
top_freqs <- head(significant_freqs, 3) # top 3 significant frequencies
N <- length(rot_ts) # 366
t <- 1:N # Time index

# Start building the formula as a string
formula_terms <- c()

# Loop through each top frequency and construct sin-cos terms
for (i in 1:nrow(top_freqs)) {
  p <- top_freqs$p[i]
  omega_p <- 2 * pi * p / N # angular frequency

  formula_terms <- c(
    formula_terms,
    paste0("cos(", omega_p, " * t)"),
    paste0("sin(", omega_p, " * t)")
  )
}

formula_str <- paste("rot_ts ~", paste(formula_terms, collapse = " + "))
data_frame <- data.frame(t = t, rot_ts = rot_ts)

# Fit linear model
model_fit <- lm(as.formula(formula_str), data = data_frame)

summary(model_fit)
```

```
##
## Call:
## lm(formula = as.formula(formula_str), data = data_frame)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -42.302 -10.218  -1.337   7.060 181.717
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    27.3898     0.9274  29.533 < 2e-16 ***
## cos(0.0343343459408721 * t)  1.1511     1.3116   0.878 0.380713
## sin(0.0343343459408721 * t) -5.7612     1.3116  -4.393 1.48e-05 ***
## cos(1.01286320525573 * t)   5.7107     1.3116   4.354 1.75e-05 ***
## sin(1.01286320525573 * t)  -0.2595     1.3116  -0.198 0.843252
```

```
## cos(0.0686686918817441 * t)  -4.5184      1.3116  -3.445 0.000639 ***
## sin(0.0686686918817441 * t)  -2.4074      1.3116  -1.835 0.067263 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 17.74 on 359 degrees of freedom
## Multiple R-squared:  0.1314, Adjusted R-squared:  0.1169
## F-statistic:  9.05 on 6 and 359 DF,  p-value: 3.187e-09
```

(h)

```
# Prediction
pred_vals <- predict(model_fit)

plot(rot_ts,
     ylab="Population Density (animals / ML)",
     main="Rotifer Population: Original vs Fitted Model")

lines(pred_vals, col="red", lwd=2)
legend("topright", legend=c("Original Data", "Fitted Model"),
     col=c("black", "red"), lty=1)
```

Rotifer Population: Original vs Fitted Model

