1) For ARMA(2,1) process

$$X_t = 0.2X_{t-1} + 0.08X_{t-2} + W_t + 0.7W_{t-1}.$$

I have at first tried 30 and 20 breaks, which didn't seem to give optimum look. Then I did a little research and used Sturges formula to determine breaks which gave me 15, and then tried Freedman-Diaconis formula which gave me 60 breaks (not included because it was also not great). I decided on 15 which can be seen on figure 3.

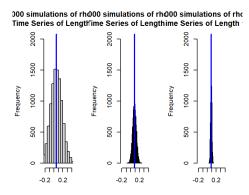


Figure 1: 30 breaks

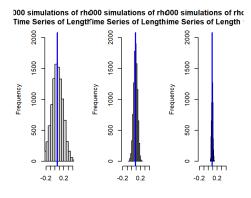


Figure 2: 20 breaks

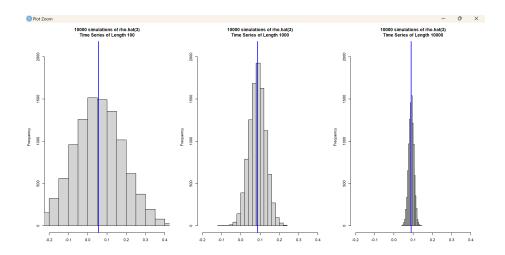


Figure 3: 15 breaks

R code:

```
# ARMA(2,1) process
phi <- c(0.2, 0.08) # AR coefficients
theta <- 0.7
                      # MA coefficient
n_values <- c(100, 1000, 10000)
n_simulations <- 10000
simulate_rho <- function(n) {</pre>
 rho_values <- numeric(n_simulations)</pre>
 for (i in 1:n_simulations) {
    # simulate ARMA(2,1)
    arma_process <- arima.sim(model = list(ar = phi,</pre>
    ma = theta), n = n)
    # compute rho(3)
    rho_values[i] <- acf(arma_process, plot = FALSE)$acf[4]</pre>
 return(rho_values)
}
# histograms
par(mfrow = c(1, 3)) # Arrange plots in a row
for (n in n_values) {
 rho_values <- simulate_rho(n)</pre>
 hist(rho_values, breaks = 15, main = paste("10000
 simulations of rho.hat(3) \n
 Time Series of Length", n),
       xlab = "", ylab = "Frequency", xlim = c(-0.2, 0.4),
```

```
ylim = c(0, 2100))
abline(v = mean(rho_values), col = "blue", lwd = 2)
}

# determining breaks
n <- length(rho_values)
sturges_breaks <- ceiling(log2(n) + 1)
print(sturges_breaks) # 15

bin_width <- 2 * IQR(rho_values) / (n^(1/3))
fd_breaks <- ceiling(diff(range(rho_values)) / bin_width)
print(fd_breaks) # 60 definitely not</pre>
```

The histograms indicating the distribution of the values of $\hat{\rho}(3)$;

For n=100, the distribution is wide, which tells us greater variability in the estimated autocorrelations.

For n=1000, the distribution becomes more concentrated around the true value of $\rho(3)$ which is 0.09.

For n=10000, even more closely concentrated with minimum variability.

Larger sample sizes lead to more accurate and precise estimates of the autocorrelation. For large n, the sample autocorrelation $\hat{\rho}(h)$ converges in probability to the true autocorrelation $\rho(h)$. This shows that increasing sample size reduces the variability in the estimates of autocorrelations.

2) For AR(2) process of

$$X_t = 0.7X_{t-1} - 0.6X_{t-2} + W_t$$

```
set.seed(4)
n <- 100
X <- arima.sim(model = list(ar = c(0.7, -0.6)), n)

par(mfrow = c(1, 2))
acf(X)
pacf(X)

rho.hat <- acf(X, plot = F)$acf[-1][1:2]
rho.hat

acf_values <- acf(X, plot = FALSE)$acf
acvf_lag0 <- var(X)</pre>
```

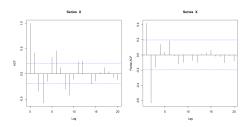


Figure 4: ACF and PACF

```
> rho.hat
[1] 0.4163640 -0.3480183
> acvf_lag0
[1] 1.740149
```

Figure 5: ACF, and sample ACVF at lag zero

```
R <- matrix(c(1, rho.hat[1], rho.hat[1], 1),2,2)
phi.hat <- solve(R,rho.hat)
phi.hat</pre>
```

$$\hat{\phi}_1 \approx 0.67897$$

$$\hat{\phi}_2 \approx -0.6307$$

These are respectable approximations to the 'true' values

$$\phi_1 = 0.7$$

and

$$\phi_1 = -0.6$$

that we started with.

Yule-Walker estimator of the variance

```
gamma.0 <- acf(X, type = 'covariance', plot = F)$acf[1]
var.W <- gamma.0*(1 - sum(rho.hat*phi.hat))
var.W</pre>
```

The results are $\hat{\gamma}(0)\approx 1.723,$ and, $\hat{\sigma}_W^2\approx 0.858.$

Maximum likelihood estimators with sarima:

```
X.model <- sarima(X, 2,0,0, no.constant = TRUE, details = FALSE)</pre>
```

Figure 6: sarima results

Figure 7: arima results

from sarima:

$$\hat{\phi}_1 \approx 0.67$$

$$\hat{\phi}_2 \approx -0.63$$

The fitted model from arima is:

$$Y_t = 0.6762Y_{t-1} - 0.6344Y_{t-2} - 0.0157 + W_t,$$

where the white noise term with variance $\hat{\sigma}^2 \approx 0.8449$.

From sarima:

$$Y_t = 0.6762Y_{t-1} - 0.6340Y_{t-2} + W_t,$$

where the white noise term with variance $\hat{\sigma}^2 \approx 0.8451$. Note: I have attached the sarima plots to the end since it is another pdf from R. The standardized residuals seem to concentrate around zero. ACF seems to have a small spike on lag 9. As for QQ plot, points on the plot lie mostly along a line so I would say it is normally distributed.

```
Recreated ljung box
X2 <- sarima(X, 2,0,0)  # fit an AR(2) model
X2.resid <- X2$fit$residuals  # store the residuals
X2.st.resid <- X2.resid/sqrt(X2$fit$sigma2)</pre>
```

LBtest = NULL

```
for(H in 3:20){
   LBtest[H]=Box.test(X2.st.resid, lag = H,
   type='Ljung-Box', fitdf = 2)$p.value
   }
plot(LBtest, ylim = c(0,1))
grid(nx = NULL, ny = NULL, col = 'lightgray',
lty = 'dotted',lwd = par('lwd'), equilogs = TRUE)
abline(h = 0.05, lty = 'dashed', col='blue')
```

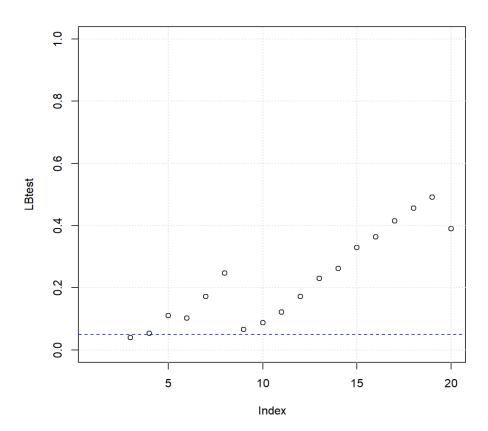


Figure 8: Ljung Box

Starting from lag 4 and on it looks like p values are above the treshold (which I also put in as 0.05) and this is the wanted result, we fail to reject.

For AR(2), as I calculated with sarima before AIC = 2.741791 and AICc = 2.743028 from;

```
> X.model \leftarrow sarima(X, 2,0,0, no.constant = TRUE, details = TRUE)
initial value 0.274042
     2 value -0.001054
iter
iter
      3 value -0.076077
     4 value -0.080720
iter
     5 value -0.081457
iter
     6 value -0.082074
iter
     7 value -0.082104
iter
     8 value -0.082106
iter
iter
     9 value -0.082113
iter 10 value -0.082113
iter 11 value -0.082113
iter 11 value -0.082113
final value -0.082113
converged
initial value -0.078038
iter 2 value -0.078042
iter 3 value -0.078043
iter 4 value -0.078043
iter 4 value -0.078043
iter 4 value -0.078043
final value -0.078043
converged
<><><><><>
Coefficients:
   Estimate
                SE t.value p.value
ar1 0.6762 0.0768 8.8009
                                0
ar2 -0.6340 0.0766 -8.2791
                                0
sigma^2 estimated as 0.8451468 on 98 degrees of freedom
AIC = 2.741791 AICc = 2.743028 BIC = 2.819946
```

Figure 9: AR(2) AIC and AICc

For AR(1),

```
> X.model1 <- sarima(X, 1,0,0, no.constant = TRUE, details = TRUE)
initial value 0.276624
iter
      2 value 0.180533
iter
      3 value 0.180527
      4 value 0.180527
iter
      4 value 0.180527
iter
final value 0.180527
converged
initial value 0.176837
     2 value 0.176828
iter
iter
      3 value 0.176828
      3 value 0.176828
iter
      3 value 0.176828
iter
final value 0.176828
converged
Coefficients:
Estimate SE t.value p.value ar1 0.4158 0.0906 4.5916 0
sigma^2 estimated as 1.421564 on 99 degrees of freedom
AIC = 3.231533 AICc = 3.231941 BIC = 3.283636
```

Figure 10: AR(1) AIC and AICc

```
> X.model3 <- sarima(X, 3,0,0, no.constant = TRUE, details = TRUE)
initial value 0.279170
iter
      2 value -0.002298
iter
      3 value -0.064675
      4 value -0.088947
iter
      5 value -0.095630
iter
      6 value -0.096191
iter
      7 value -0.096505
iter
iter
      8 value -0.096506
iter
      8 value -0.096506
final value -0.096506
converged
initial value -0.092284
iter
      2 value -0.092354
      3 value -0.092391
iter
      4 value -0.092400
iter
      5 value -0.092400
iter
      5 value -0.092400
iter
      5 value -0.092400
final value -0.092400
converged
Coefficients:
   Estimate
                SE t.value p.value
ar1 0.5661 0.0994 5.6949 0.0000
ar2 -0.5185 0.1014 -5.1138 0.0000
ar3 -0.1705 0.0998 -1.7082 0.0908
sigma^2 estimated as 0.8205279 on 97 degrees of freedom
AIC = 2.733077 AICC = 2.735577 BIC = 2.837284
```

Figure 11: AR(3) AIC and AICc

Since AIC and AICc choose the model with the smallest value, and AR(3) gives the smallest AIC with 2.733077 and AICc with 2.735577, AR(3) is a better fit.

3) I am using The AR(2) process from the first assignment:

$$X_t - 0.8X_{t-1} + 0.2X_{t-2} = W_t,$$

$$\mathrm{AIC}(\beta) = -2\log L\left(\beta, \frac{1}{n}S(\beta)\right) + 2(p+q+1).$$

The penalty term in the formula will increase to account for the extra parameter.

$$k=3 \quad \text{(2 AR coefficients + variance)}.$$

$$\text{AIC}_{\text{no mean}} = -2 \cdot (-200) + 2 \cdot 3 = 406$$

With Mean:

$$k = 4$$
 (2 AR coefficients + mean + variance).
AIC_{with mean} = $-2 \cdot (-200) + 2 \cdot 4 = 408$

The difference is 2 which makes sense since the additional parameter of mean is added.

I calculated log likelihood with sarima

```
set.seed(123)
n <- 200
ar_params <- c(0.8, -0.2)
x <- arima.sim(model = list(ar = ar_params), n = n)
#AR(2) model without mean
fitzmean <- sarima(x, 2, 0, 0, no.constant = TRUE,
details = TRUE)
#AR(2) model with mean
fitmean <- sarima(x, 2, 0, 0, no.constant = FALSE,
details = TRUE)
#log likelihoods
loglikzmean <- fitzmean$fit$loglik # -271.9306
loglikmean <- fitmean$fit$loglik # -271.9306</pre>
```

Since including a mean adds an extra parameter to the model, it can lead to a penalty in the AIC and AICc. If the true process has a non-zero mean, including a mean in the model can improve the fit to the data, potentially leading to a decrease in the AIC and AICc despite the added parameter. So it depends on the trade-off between the penalty for the extra parameter and the improvement in the model fit. In terms of dataset, it is a small dataset, the penalty for adding a mean is greater, making it less likely for AICc to decrease when including a mean. Considering the AICc formula;

$$AICc(\beta) = -2\log L\left(\beta, \frac{1}{n}S(\beta)\right) + 2(p+q+1) \cdot \frac{n}{n-p-q-2}$$

Adding more parameters will cause more severe penalty and may outweigh the improvement in model fit.

