

# Time Series Analysis

## Model Diagnostics

Hossein Moradi Rekabdarkolaee

South Dakota State University

**email:** [hossein.moradirekabdarkolaee@sdstate.edu](mailto:hossein.moradirekabdarkolaee@sdstate.edu).

**Office:** CAME Building, *Room: 240.*

# Introduction

- So far, we have learned
  - Different components of time series
    - Trend,
    - Seasonality,
    - Error.
  - Different type of time series
    - Stationary (weak and strong)
    - Non-stationary.
  - Different models for time series
    - Autoregressive,
    - Moving Average,
    - ARMA,
    - ARIMA.

- Different way to find the order of time series
  - ACF,
  - PACF,
  - EACF,
  - Information Criteria-based method (AIC, BIC, AICc, etc.)
- Different approaches to estimate the parameters of the model we choose to fit the time series.
  - Method of Moment,
  - Least Squares,
  - Maximum Likelihood.

# Introduction

- In this Section, we are going to talk about
  - Model diagnostics or model criticism,
  - What does it mean
  - How to do it.
  - Do some examples.

# Diagnostics

- Model diagnostics, or model criticism, is concerned with testing the goodness of fit of a model and, if the fit is poor, suggesting appropriate modifications.
- The main objective is to **check the adequacy** of the selected model.
- After fitting an ARIMA model to data, to evaluate the fit, diagnostics checking of the **residuals** need to be performed.

# Diagnostics

- If fact all of the relevant information from the data should be extracted by the model.
- For instance, the part of the data unexplained by the model (**residuals**) should be small and no systematic patterns should be left in the residuals.
- **residuals = actual - predicted**
- **Standardization** allows us to see residuals of unusual size much more easily.

# Residual Analysis

- Diagnostic checking essentially involves the **statistical properties** of the residuals of the fitted model (normality assumptions, weak white noise assumptions, etc.)

- The **standardized residuals** can be calculated as follows

$$e_t = \frac{Y_t - \hat{Y}_t}{s}, \quad \text{where } s^2 = \frac{1}{n-1} \sum_{t=1}^n (Y_t - \hat{Y}_t)^2$$

- Then,  $e_t$  should follow standard Normal White Noise.
- So residuals should be plotted in order to check for
  - substantial residual autocorrelation (Sample ACF, PACF),
  - normality (Q-Q-plots, histogram, normality test),
  - constant variance.

# Residual Analysis

- For instance, consider in particular an **AR(2)** model with a constant term:

$$Y_t = \phi_1 Y_{t-1} + \phi_2 Y_{t-2} + \theta_0 + \epsilon_t,$$

then, estimated  $\phi_1$ ,  $\phi_2$  and  $\theta_0$ , the residuals are defined as

$$e_t = Y_t - \hat{\phi}_1 Y_{t-1} - \hat{\phi}_2 Y_{t-2} - \hat{\theta}_0.$$

- For **general invertible ARMA model**, when  $\theta_0 = 0$ , we have

$$Y_t = \pi_1 Y_{t-1} + \pi_2 Y_{t-2} + \pi_3 Y_{t-3} + \dots + \epsilon_t,$$

- Thus, the estimated residuals are defined as

$$e_t = Y_t - \hat{\pi}_1 Y_{t-1} - \hat{\pi}_2 Y_{t-2} - \hat{\pi}_3 Y_{t-3} - \dots$$



- To check against **changes in mean and/or variance level**, we need to plot the  $e_t$ .
- Let us look at some examples.
- Things that we want to check:
  - We want to make sure that the residuals do not have significant ACF and PACF.
  - We want to check if the residuals are normally distributed.
  - Is the transformation necessary?

# Autocorrelation of the Residuals

- To check the independence of the noise terms in the model, we examine the **sample ACF of the residuals**
- The sample ACF of the residuals are denoted **with  $\hat{r}_k$**
- We know that for **true white noise and large  $n$** , the sample ACF are approximately **uncorrelated and normally distributed** with zero means and variance  $\frac{1}{n}$ .
- Generally, the residuals are approximately normally distributed with zero means; however, for **small lags  $k$  and  $j$** , the variance of  $\hat{r}_k$  can be substantially **less than  $\frac{1}{n}$** ,
- and the estimates  $\hat{r}_k$  and  $\hat{r}_j$  can be **highly correlated**.

# Autocorrelation of the Residuals - AR(1)

- For instance for **AR(1) model** it can be shown that

$$\text{Var}(\hat{r}_1) \approx \frac{\phi^2}{n}$$

$$\text{Var}(\hat{r}_k) \approx \frac{1 - (1 - \phi^2)\phi^{2k-2}}{n} \quad \text{for } k > 1$$

$$\text{corr}(\hat{r}_1, \hat{r}_k) \approx -\text{sign}(\phi) \frac{(1 - \phi^2)\phi^{k-2}}{1 - (1 - \phi^2)\phi^{2k-2}} \quad \text{for } k > 1$$

where

$$\text{sign}(\phi) = \begin{cases} 1 & \text{if } \phi > 0 \\ 0 & \text{if } \phi = 0 \\ -1 & \text{if } \phi < 0 \end{cases}$$

- Simulation results are showing that  $\text{Var}(\hat{r}_1) \approx \frac{1}{n}$  is a reasonable approximation for  $k \geq 2$  over a wide range of  $\phi$ -values.

# Autocorrelation of the Residuals - AR(2)

- For instance for **AR(2) model** it can be shown that

$$\text{Var}(\hat{r}_1) \approx \frac{\phi_2^2}{n}, \quad \text{Var}(\hat{r}_2) \approx \frac{\phi_2^2 + \phi_1^2(1 + \phi_2)^2}{n}$$

- If the AR(2) parameters are not too close to the stationarity boundary, then

$$\text{Var}(\hat{r}_k) \approx \frac{1}{n} \quad \text{for } k \geq 3$$

- It can be shown that results **analogous to those for AR models hold for MA models**.
- In particular, **replacing  $\phi$  by  $\theta$**  gives the results for the MA(1) case.
- Similarly, results for the MA(2) case can be stated by **replacing  $\phi_1$  and  $\phi_2$  by  $\theta_1$  and  $\theta_2$** , respectively.

# The Ljung-Box Test

- In addition to looking at residual correlations at individual lags, it is useful to have a **test that of their magnitudes as a group**.
- A common method for testing against residual auto-correlation is **Ljung-Box test**:

$$\begin{cases} H_0 : & \text{The error terms are uncorrelated,} \\ H_a : & \text{the error terms are correlated.} \end{cases}$$

- Ljung-Box statistic:

$$Q = n(n+2) \sum_{k=1}^K \frac{\hat{r}_k^2}{n-k}$$

# The Ljung-Box Test

- If  $e_t \sim WN$  then  $Q \sim \chi_{(K-p-q)}$ , as  $n \rightarrow \infty$
- Here the maximum lag  $K$  is selected somewhat arbitrarily but large enough that the weights are negligible for  $j > K$ .
- The test would reject the  $ARMA(p, q)$  model if the observed value of  $Q$  exceeded an appropriate critical value in a chi-square distribution with  $K - p - q$  degrees of freedom.

# Overfitting and Parameter Redundancy

- After specifying and fitting what we believe to be an adequate model, we fit a slightly more general model; that is, a model “close by” that contains the original model as a special case.
- For example, if an AR(2) model seems appropriate, we might overfit with an AR(3) model
- then, the original AR(2) model would be confirmed if:
  - 1 the estimate of the additional parameter,  $\phi_3$ , is not significantly different from zero, and
  - 2 the estimates for the parameters in common,  $\phi_1$  and  $\phi_2$ , do not change significantly from their original estimates
  - 3 Notice that, using higher order (p,q) leads to suboptimal estimation (overfitting).

# Overfitting and Parameter Redundancy

- The implications for fitting and overfitting models are as follows:
  - 1 Specify the original model carefully. If a simple model seems at all promising, check it out before trying a more complicated model.
  - 2 When overfitting, do not increase the orders of both the AR and MA parts of the model simultaneously.
  - 3 Extend the model in directions suggested by the analysis of the residuals.
  - 4 For example, if after fitting an MA(1) model, substantial correlation remains at lag 2 in the residuals, try an MA(2), not an ARMA(1, 1).



# Summary

- The parsimony principle, Keep it simple!
- If two models performs equally well in terms of residual variance or likelihood, pick the one with fewer parameters
- Including too many parameters can lead to overfitting
- This applies to  $p$  and  $q$ , but also to  $d$ .
- Too high  $d$  leads overdifferencing