## Time Series: Issues with R

Andrius Buteikis, andrius.buteikis@mif.vu.lt http://web.vu.lt/mif/a.buteikis/ There are various (sometimes little, sometimes more serious) issues and function specifics which may do always seem to make sense. A number of them are mentioned at the homepage of Time Series, 4th Edition

# Issue No. 1: Functions with the same names override one another

There are two quite popular packages are plyr and dplyr. First, look at what happens when you load 'dplyr:

library(dplyr)

## Warning: package 'dplyr' was built under R version 3.5.2

## ## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##

## filter, lag

## The following objects are masked from 'package:base':
##
intersect, setdiff, setequal, union

As you can see, functions like filter and lag are overloaded with the new ones from the package.

### Now assume that we want to load plyr:

```
library(plyr)
## -----
## You have loaded plvr after dplvr - this is likely to cause problems.
## If you need functions from both plyr and dplyr, please load plyr first, then dplyr:
## library(plyr); library(dplyr)
## -----
##
## Attaching package: 'plyr'
## The following objects are masked from 'package:dplyr':
##
##
     arrange, count, desc, failwith, id, mutate, rename, summarise,
##
     summarize
```

In addition to function overloading we now get an additional warning specifically for this package.

- If we want to have both plyr and dplyr, we need to firstly load plyr and THEN dplyr;
- We need to be careful in regards to function names we might override functions from some packages with newer ones from different packages.
- You can load specific functions by referencing the package, like plyr::some\_function(...) and dplyr::another\_function() this prevents errors, though for more advanced functions and/or expressions, additional functions from the package may need to be loaded, see import package.
- plyr and dplyr may have additional conflicts with some Multivariate AR (VAR) model packages.

## Issue No. 2: dplyr and the lag() function

Assume that our data  $Y_t$  starts at time t = 1

x <- ts(1:5) x

## Time Series:
## Start = 1
## End = 5
## Frequency = 1
## [1] 1 2 3 4 5

Assume that we want to automatically create  $Y_{t-1}$ . By default the lag() function is a **forward** shift (i.e. into the future):

```
cbind(x, lag(x), lag(x, -1))
```

```
## Time Series:
## Start = 0
## End = 6
## Frequency = 1
     x lag(x) lag(x, -1)
##
## O NA
           1
                    NA
## 1 1
           2
                    NA
## 2 2
        3
                    1
## 3 3
        4
                     2
## 4 4
          5
                     3
## 5 5
          NA
                     4
## 6 NA
          NΔ
                     5
```

lag(x) - in this case, instead of lagging the values back, we append the series at time t = 0. At time t = 1, we have Y<sub>1</sub> = 1 and we would assume that Y<sub>t-1</sub> = Y<sub>0</sub> = NA. However, by default we would get the future value, i.e. Y<sub>t+1</sub> = Y<sub>2</sub> = 2.

lag(x, -1) - in this case at t = 1 we get an NA value - a BACKWARD shift - as we would expect.

Firstly, if we wanted to load dplyr and run the same function, we would get an **error**:

suppressPackageStartupHessages({# supress the package warnings if we are AESOLUTELY sure about what we are doing library(dplyr) }) lag(x)

## Error: `x` must be a vector, not a ts object, do you want `stats::lag()`?

# dplyr::lag(x) # alternatively, without loading

This does not work for ts(...) variables - it results in an error. If we use the value as a vector:

lag(c(x))

## [1] NA 1 2 3 4

We get what would be the equivalent of stats::lag(x, -1) - aBACKWARD shift

## Issue No. 3 Overreliance on auto.arima

```
suppressPackageStartupMessages({
    library(astsa)
    library(forecast)
})
tmp_mdl <- auto.arima(UnempRate)</pre>
```

tmp\_mdl

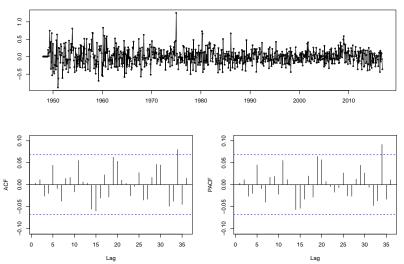
```
## Series: UnempRate
## ARIMA(3,0,1)(2,1,2)[12]
##
## Coefficients:
```

```
## Warning in sqrt(diag(x$var.coef)): NaNs produced
```

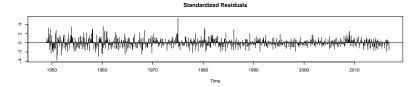
## ar1 ar2 ar3 ma1 sar1 sar2 sma1 sma2 ## 1.6852 -0.5763 -0.1204 -0.6106 -0.2849 0.0376 -0.4645 -0.2408 ## s.e. 0.0379 0.0657 0.0394 0.0461 0.0464 0.0510 0.0392 NaN ## ## sigma^2 estimated as 0.05465: log likelihood=25.52 ## ATC=-33 03 ATCc=-32 81 BTC=9 3

We even get a warning - the standard errors for the last coefficient are not calculated.

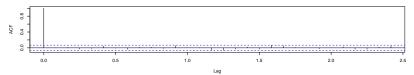
## forecast::tsdisplay(tmp\_mdl\$residuals)



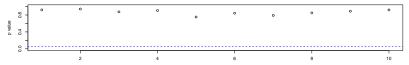
tmp\_mdl\$residuals











```
We can manually specify a model:
```

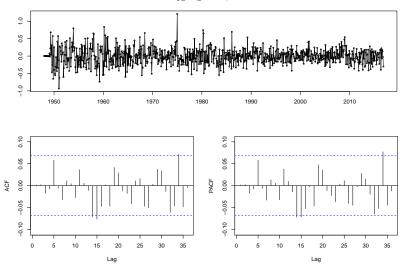
```
tmp_mdl_manual <- forecast::Arima(UnempRate, order = c(2, 1, 1), seasonal = c(0, 1, 1))
tmp_mdl_manual</pre>
```

```
## Scries: UnempRate
## ARIMA(2,1,1)(0,1,1)[12]
##
## Coefficients:
## ar1 ar2 ma1 sma1
## 0.5897 0.1342 -0.4831 -0.7676
## s.e. 0.1105 0.0465 0.1090 0.0254
##
## sigma^2 estimated as 0.05587: log likelihood=15.69
## AIC=-21.3 AICC=-21.3 BIC=-2.13
```

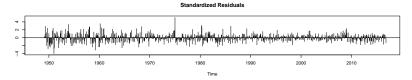
#### Note:

- The BIC is smaller for the manually specified model the penalty for including more variables in the auto.arima model is harsher.
- ▶ The AIC is smaller for the auto.arima model.

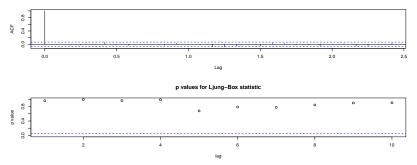
### forecast::tsdisplay(tmp\_mdl\_manual\$residuals)



tmp mdl manual\$residuals



ACF of Residuals



The residuals are not that much different in terms of their similarity to a WN process.

## Issue # 4: intercept vs mean

Remember that:

- for MA(q) models:  $Y_t = \alpha + \theta(L)\epsilon_t$ , the mean is the intercept:  $\mathbb{E}(Y_t) = \alpha$ .
- ► for stationary AR(p) or ARMA(p,q) models  $\phi(L)Y_t = \alpha + \theta(L)\epsilon_t$ , the mean is  $\mu = \alpha/(1 - \phi_1 - ... - \phi_p)$ , or  $\alpha = \mu \cdot (1 - \phi_1 - ... - \phi_p)$ .

In other words when there is an AR term in the model, the intercept is  $\ensuremath{\text{NOT}}$  the mean.

```
set.seed(1)
x = arima.sim(list(order = c(1,0,0), ar= 0.4), n = 1000) + 10
```

The true mean of the process is 50, which translates to  $\alpha = 10 \cdot (1 - 0.4) = 6.$ 

mean(x)

## [1] 9.978789

Verify this via manual simulation:

```
set.seed(1)
epsilon <- rnorm(1000, mean = 0, sd = 1)
x <- NULL
x[1] <- 6 + epsilon[1]
for(j in 2:length(epsilon)){
    x[j] <- 6 + 0.4 * x[j - 1] + epsilon[j]
}</pre>
```

mean(x)

## [1] 9.974864

If we estimate using stats::arima:

mdl\_1 <- arima(x, order = c(1, 0, 0))
mdl\_1</pre>

##
## Call:
## arima(x = x, order = c(1, 0, 0))
##
## Coefficients:
## ar1 intercept
## 0.3541 9.9717
## s.e. 0.0299 0.0510
##
## sigma^2 estimated as 1.085: log likelihood = -1459.74, aic = 2925.48

The naming is **incorrect** - the intercept is actually the mean.

On the other hand, if we use forecast::Arima:

```
mdl_2 \leftarrow forecast::Arima(x, order = c(1, 0, 0)) mdl_2
```

## Series: x
## ARIMA(1,0,0) with non-zero mean
## ARIMA(1,0,0) with non-zero mean
## Coefficients:
## 0.3541 9.9717
## s.e. 0.0299 0.0510
## sigma^2 estimated as 1.087: log likelihood=-1459.74
## alc=2925.48 AICc=2925.51 BIC=2940.21

The naming appears correct ... unless we extract the coefficients:

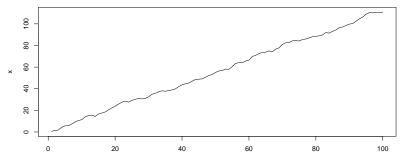
coef(mdl\_2)

## ar1 intercept
## 0.3541427 9.9716752

Then the naming is again **incorrect**.

Issue # 5.1: stats::arima vs forecast::Arima for  $Y_t \sim I(d)$ 

```
set.seed(1)
v = rnorm(100, mean = 1, sd = 1)
x = ts(cumsum(v))
plot.ts(x)
```



Time

The stats package is one of the default packages, which come with the installation of R.

```
mdl_stats <- arima(x, order = c(1, 1, 0), include.mean = TRUE)
mdl_stats</pre>
```

```
##
## Call:
## arima(x = x, order = c(1, 1, 0), include.mean = TRUE)
##
## Coefficients:
## ari
## ari
## o.6031
## s.e. 0.0793
##
## sigma^2 estimated as 1.294: log likelihood = -153.46, aic = 310.91
```

```
mdl_forcast <- forecast::Arima(x, order = c(1, 1, 0), include.drift = TRUE)
mdl_forcast</pre>
```

```
## Series: x
## ARIMA(1,1,0) with drift
##
## Coefficients:
## -0.0031 1.1163
## s.e. 0.1002 0.0897
##
## sigma^2 estimated as 0.8178: log likelihood=-129.51
## ALC=265.01 ALC==265.26 BLC==272.8
```

Note that when we are differencing the series:

- stats::arima() fits  $\Delta Y_t = \phi \Delta Y_{t-1} + \epsilon_t$  (no constant);
- forecast::Arima() fits  $\Delta Y_t = \alpha + \phi \Delta Y_{t-1} + \epsilon_t$  (constant).

Consequently, if we want to fit a I(d) series with a drift using stats::arima(), there are two ways to go about this:

```
fit the differenced series, diff(x), with a constant:
```

```
arima(diff(x), order = c(1, 0, 0), include.mean = TRUE)
```

```
##
## Call:
## call:
## arima(x = diff(x), order = c(1, 0, 0), include.mean = TRUE)
##
## Coefficients:
## ar1 intercept
## -0.0031 1.1163
## s.e. 0.1002 0.0897
##
# sigma^2 estimated as 0.8012: log likelihood = -129.51, aic = 265.01
```

In this case, for the AR model, the intercept is actually the mean of diff(x).

#### specify the constant as an exogeneous variable:

```
arima(x, order = c(1, 1, 0), xreg = 1:length(x))
```

```
##
## Call:
## call:
## arima(x = x, order = c(1, 1, 0), xreg = 1:length(x))
##
## Coefficients:
## ari 1:length(x)
## ari 1:length(x)
## o.0.0031 1.1163
## s.e. 0.1002 0.0897
##
## sigma^2 estimated as 0.8012: log likelihood = -129.51, aic = 265.01
```

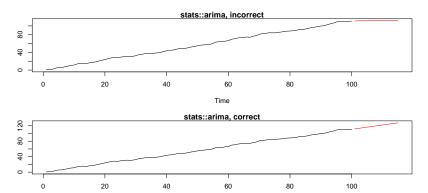
In this case, the exogeneous variable is the intercept of diff(x).

alternatively, use forecast::Arima() (where it is named drift).

## Issue # 5.2: Forecasting with $Y_t \sim I(d)$

Assume that we want to forecast via the **default** predict() function and we *ignore* the fact that we do not include a drift in our model:

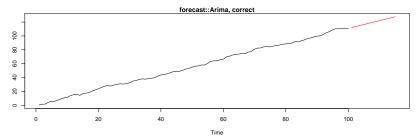
```
par(mfrow = c(2, 1), mai = c(1, 0.5, 0.2, 0.2))
ts.plot(x, forc_1$pred, col = 1:2, main = "stats::arima, incorrect")
ts.plot(x, forc_2$pred, col = 1:2, main = "stats::arima, correct")
```

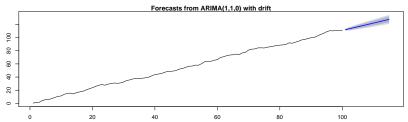


Time

```
mdl_forcast <- forecast::Arima(x, order = c(1, 1, 0), include.drift = TRUE)
#
forc_3 <- forecast(mdl_forcast, 15)</pre>
```

```
par(mfrow = c(2, 1), mai = c(1, 0.5, 0.2, 0.2))
ts.plot(x, forc_3$mean, col = 1:2, main = "forcast::Arima, correct")
plot(forc_3, cmain = "stats::arima, correct")
```





## Issue # 6: Tests for Residuals autocorrelation of an $\operatorname{ARMA}(p,q)$ model

The null hypothesis:

$$H_0: \rho(1) = ... = \rho(k) = 0$$
  
 $H_1: \exists j: \rho(j) \neq 0$ 

The Ljung-Box test:

$$Q(k) = T(T+2)\sum_{m=1}^{k} \frac{\widehat{\rho}^2(m)}{T-m}$$

Box-Pierce test:

$$Q_{
m BP} = T \sum_{m=1}^{k} \widehat{
ho}^2(m)$$

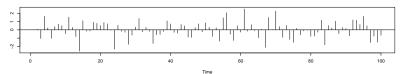
If we are testing whether Y<sub>t</sub> exhibits significant autocorrelation, then we reject the null hypothesis if

$$Q(k) > \chi^2_{1-lpha, k}$$

where k are the degrees of freedom.

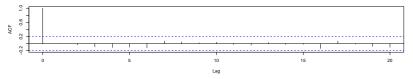
► If we are testing the residuals of an estimated ARIMA(p, q) model (without constant), the degrees of freedom need to be adjusted to reflect the parameter estimation. In such cases the degrees of freedom should be set to k - p - q > 0.

The same applies to  $Q_{\mathrm{BP}}$ 

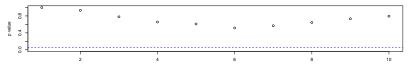


Standardized Residuals









##		INCORRECT	CORRECT
##	Lag_3	0.7796620	
##	Lag_4	0.6572870	0.2967712
##	Lag_5	0.6105057	
##	Lag_6	0.5171249	
##	Lag_7	0.5677989	0.3300712
##	Lag_8	0.6452965	0.4212575
##	Lag_9	0.7323974	0.5310294
##	Lag_10	0.7948932	0.6205997