## Employment Analysis

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#### Organise analysis in preparation for constructing poster.

First load all required packages:

```
library(car)
library(tseries)
library(astsa)
```

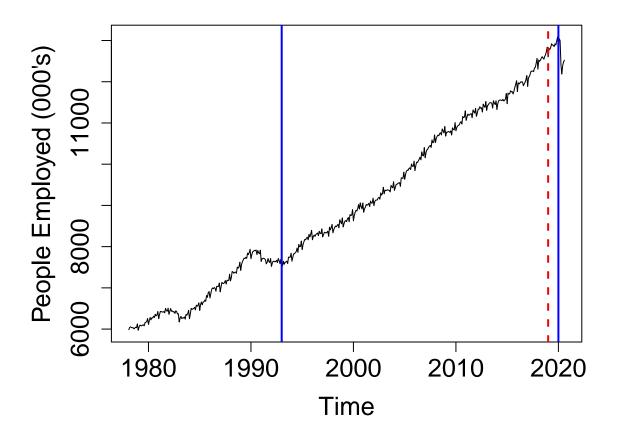
Load in the data:

```
dat <- read.csv("employment_data.csv", fileEncoding = 'UTF-8-BOM')
head(dat)</pre>
```

```
##
     Observation.times Time.series.values
## 1
                Feb-78
                                     5985.7
## 2
                Mar-78
                                     6040.6
                                     6054.2
## 3
                Apr-78
## 4
                May-78
                                     6038.3
## 5
                 Jun-78
                                     6031.3
                 Jul-78
                                     6036.1
```

Create a time series object from the data and plot. The blue lines are visually detected structural breakpoints - contextual reasoning is recession in the 90s and COVID-19 in 2020. The red line indicates the training/test split.

```
ts_dat_test <- ts(dat[, 2], start = c(1978, 2), end = c(2020, 8), frequency = 12)
#jpeg("plot_ts.jpg", width = 990, height = 400)
par(cex.lab = 1.6, mar = c(5, 7, 1, 1))
plot.ts(ts_dat_test, ylab="People Employed (000's)", axes=FALSE) #updated units
axis(1, cex.axis = 1.6)
axis(2, cex.axis = 1.6)
box()
abline(v = 1993, col = "blue", lwd = 2)
abline(v = 2020, col = "blue", lwd = 2)
abline(v = 2019, col = "red", lty = 2, lwd = 2)</pre>
```

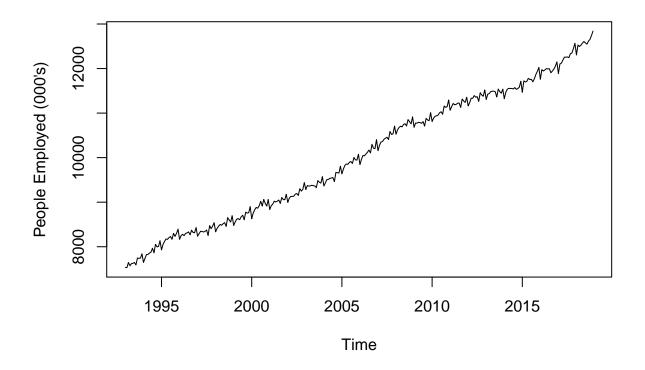


Instructed to truncate data from January 1993 to December 2019 (inclusive)

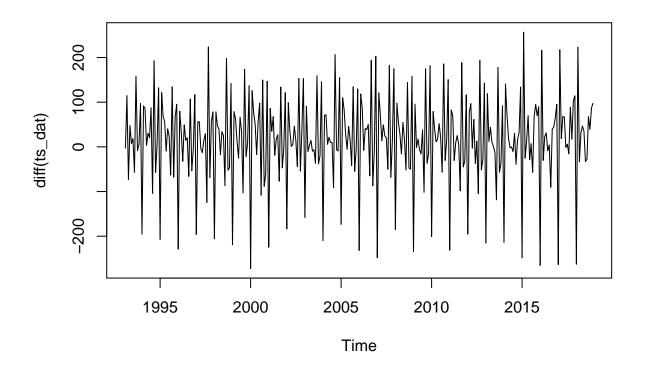
```
dat[dat$Observation.times == "Jan-93",]
##
       Observation.times Time.series.values
## 180
                  Jan-93
                                     7533.7
dat[dat$Observation.times == "Dec-19",]
##
       Observation.times Time.series.values
## 503
                  Dec-19
                                    13087.1
dat[dat$Observation.times == "Jan-19",]
##
       Observation.times Time.series.values
## 492
                  Jan-19
                                    12603.1
```

We only need rows 180-503 for the truncated dataset. Splitting into training and test sets where the test set is all of 2019, we have train [180:491] and test [492:503].

```
trunc_dat <- dat[180:503,] # all data after truncating
train_dat <- dat[180:491,] # training data
test_dat <- dat[492:503,] # test data
join_dat <- dat[491:492,] # the month between training data and predictions (for plotting)
# ts for model fitting
ts_dat <- ts(train_dat[, 2], start = c(1993, 1), end = c(2018, 12), frequency = 12)
# ts for model testing
test_ts <- ts(test_dat[, 2], start = c(2019, 1), end = c(2019, 12), frequency = 12)
# between train and test
ts_join <- ts(join_dat[, 2], start = c(2018, 12), end = c(2019, 1), frequency = 12)
plot.ts(ts_dat, ylab="People Employed (000's)")</pre>
```



plot.ts(diff(ts\_dat)) # We are not actually taking the difference yet!



# This 2nd plot is just to help observe trends in variance.

The trend in mean is readily observable. Difficult to determine a trend in variance - there appears to be frequent changes, which are easier to see after incorporating lags of 1. Check statistically for stationarity using the Kwiatkowski-Phillips-Schmidt-Shin (KPSS) test, which has the following hypotheses:

 $H_o$ : TS is stationary  $H_a$ : TS in not stationary

```
kpss.test(ts_dat)
```

```
## Warning in kpss.test(ts_dat): p-value smaller than printed p-value
##
## KPSS Test for Level Stationarity
##
## data: ts_dat
## KPSS Level = 5.3101, Truncation lag parameter = 5, p-value = 0.01
```

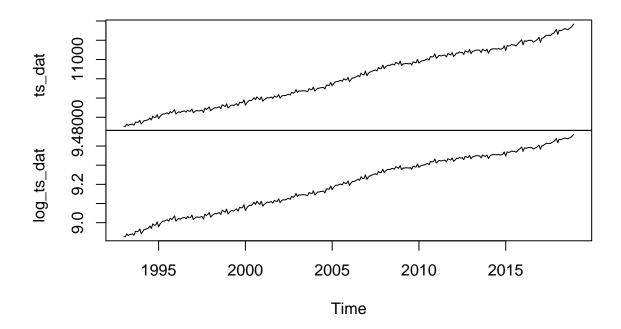
The small p-value indicates that we should reject the null and conclude that the ts is not stationary.

As a rough test of constant variance (Levene's isn't really valid because time series data isn't independent)

The small p-value of 0.04837 confirms that the data exhibits heteroscedasticity. Therefore we will perform a log transformation to attempt to reduce this:

```
log_ts_dat <- log(ts_dat)
plot.ts(cbind(ts_dat, log_ts_dat))</pre>
```

# cbind(ts\_dat, log\_ts\_dat)



```
leveneTest(log_ts_dat, Group)

## Warning in leveneTest.default(log_ts_dat, Group): Group coerced to factor.

## Levene's Test for Homogeneity of Variance (center = median)

## Df F value Pr(>F)

## group 3 0.4451 0.7209

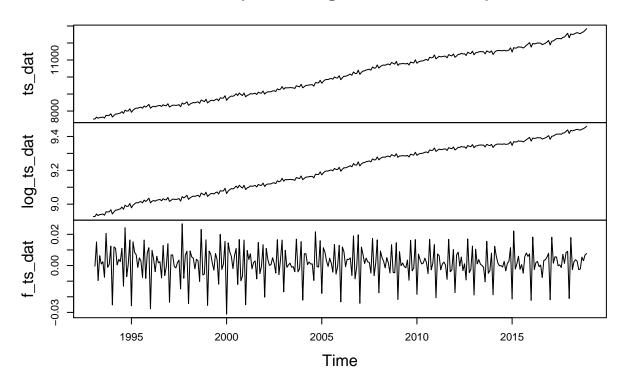
## 308
```

At a significance level of 5%, the p-value above of 0.7209 provides very weak evidence and we fail to reject the null hypothesis of equal variance among groups. Thus the heteroscedasticity has been reduced.

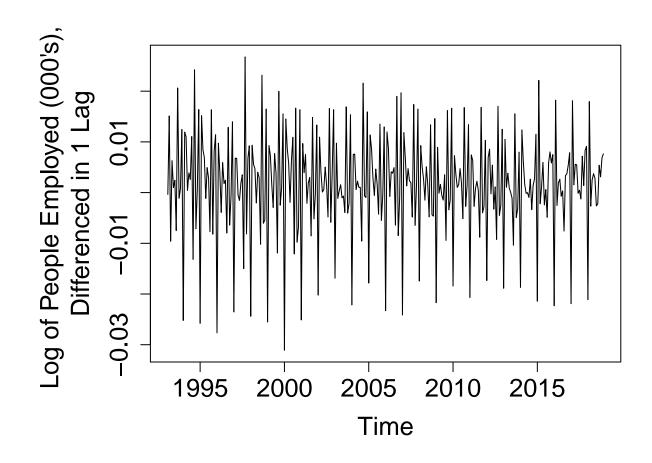
Next, to reduce the trend in mean, apply differencing of 1 lag to our TS with stabilised variance:

```
f_ts_dat <- diff(log_ts_dat, 1)
plot.ts(cbind(ts_dat, log_ts_dat, f_ts_dat))</pre>
```

### cbind(ts\_dat, log\_ts\_dat, f\_ts\_dat)



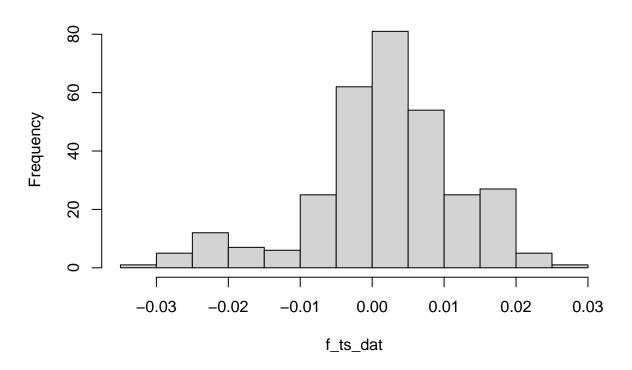
```
#jpeg("stationary_ts.jpg", width = 990, height = 400)
par(cex.lab = 1.6, mar = c(5, 7, 1, 1))
plot.ts(f_ts_dat, ylab="Log of People Employed (000's), \nDifferenced in 1 Lag", axes=FALSE)
axis(1, cex.axis = 1.6)
axis(2, cex.axis = 1.6)
box()
```



To confirm constant mean and variance and a Gaussian distribution for the time series, a Shapiro-Wilk normality test is performed:

hist(f\_ts\_dat)

# Histogram of f\_ts\_dat



### shapiro.test(f\_ts\_dat)

```
##
## Shapiro-Wilk normality test
##
## data: f_ts_dat
## W = 0.96193, p-value = 2.913e-07
```

The small p-value indicates likely non-normality, but this test isn't really valid for TS. Instead, check statistically for stationarity using the Kwiatkowski-Phillips-Schmidt-Shin (KPSS) test:

```
## Warning in kpss.test(log_ts_dat): p-value smaller than printed p-value

## ## KPSS Test for Level Stationarity

## data: log_ts_dat

## KPSS Level = 5.3037, Truncation lag parameter = 5, p-value = 0.01

kpss.test(f_ts_dat)

## Warning in kpss.test(f_ts_dat): p-value greater than printed p-value

## ## KPSS Test for Level Stationarity

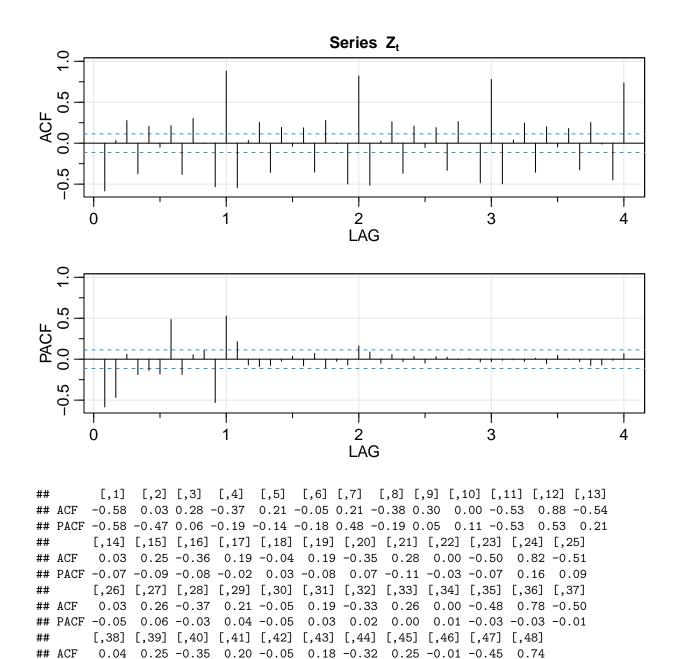
## data: f_ts_dat

## KPSS Level = 0.065128, Truncation lag parameter = 5, p-value = 0.1
```

The final ts has a high p-value of 0.1, which is statistically significant at a significance level of 5%. Therefore we fail to reject the null hypothesis, and have reasonable evidence that the final ts is stationary.

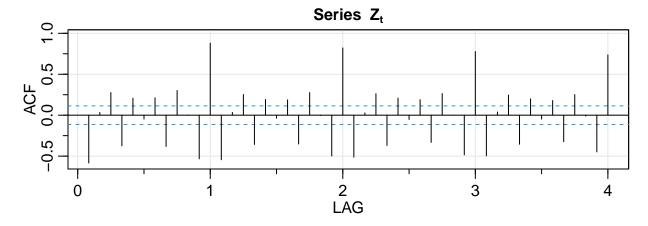
Next, the ACF and PACF of the differenced ts are plotted for analysis.

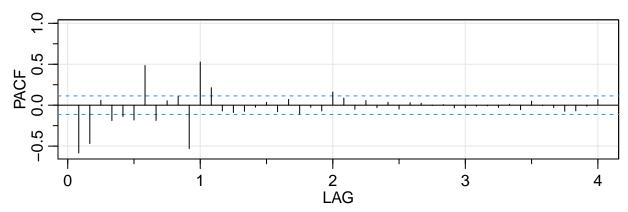
```
acf2(f_ts_dat, main=expression(bold("Series " ~ Z[t])))
```



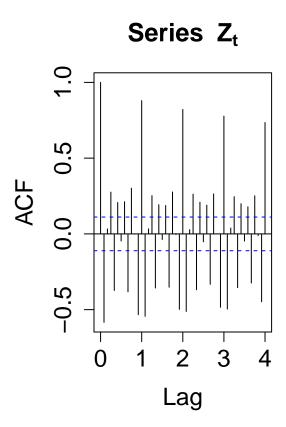
```
par(mfrow=c(1,2))
acf2(f_ts_dat, main=expression(bold("Series " ~ Z[t])))
```

## PACF -0.01 -0.03 0.01 -0.05 0.05 -0.01 -0.03 -0.07 -0.07 -0.01 0.07





```
#jpeg("acf_ts.jpg", width = 990, height = 300)
par(cex.lab = 1.5, cex.axis = 1.5, cex.main = 1.6, mar = c(5, 5, 4, 2))
acf(f_ts_dat,lag.max=48, main=expression(bold("Series " ~ Z[t])))
#dev.off()
```



Seasonal patterns in the ACF for TS\_name (Figure [above]) show a slow decay in the dominant lags. To mitigate this, the time series was differenced in 12 lags giving [differenced model] for which the ACF and PACF are plotted in Figure [below].

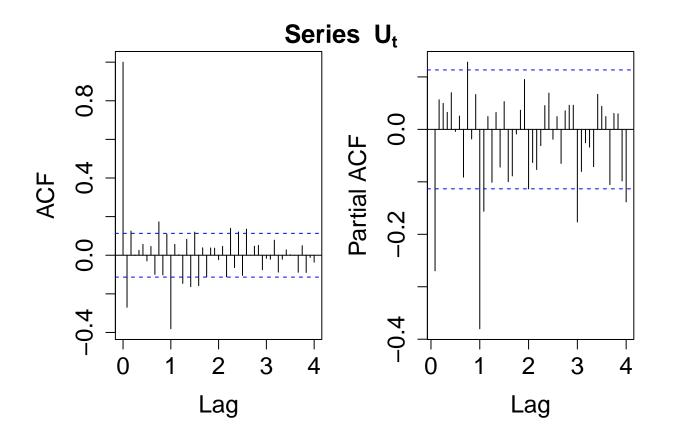
```
ts_dat_12 <- diff(f_ts_dat, 12)
kpss.test(ts_dat_12) #Big enough to call stationary

## Warning in kpss.test(ts_dat_12): p-value greater than printed p-value

##
## KPSS Test for Level Stationarity
##
## data: ts_dat_12
## KPSS Level = 0.026132, Truncation lag parameter = 5, p-value = 0.1

#acf2(ts_dat_12)

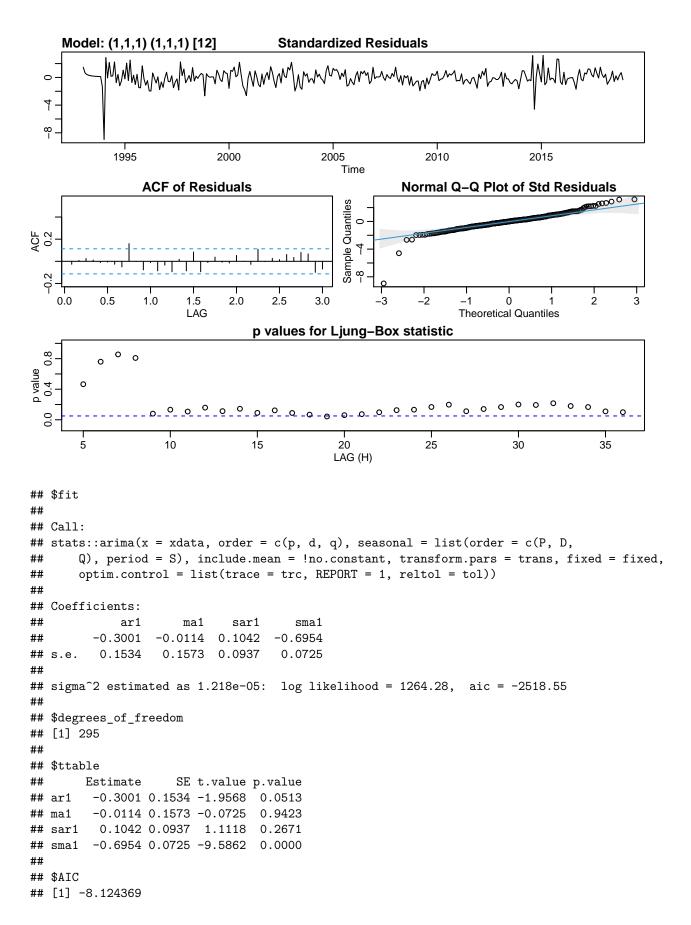
#jpeg("seasonal_acf.jpg", width = 990, height = 350)
par(cex.lab = 1.5, cex.main = 1.6, cex.axis = 1.5, mar = c(4.5, 4.5, 3, 1), mfrow=c(1,2))
acf(ts_dat_12, lag.max = 48, main = "")
pacf(ts_dat_12, lag.max = 48, main = "")
mtext(expression(bold("Series " ~ U[t])), line = -3, outer = TRUE, cex = 1.6)</pre>
```



At the seasonal level, these indicate a cutoff at 1 in the ACF and tailing off in PACF, possibly indicating P=0 and Q=1. Lags 1,2,...,11 suggest several choices, so estimates of  $0 \le p \le 1$  and  $0 \le q \le 1$  are made and explored.

```
sarima(log_ts_dat, p = 1, d = 1, q = 1, P = 1, D = 1, Q = 1, S = 12) #AICc -8.123946
```

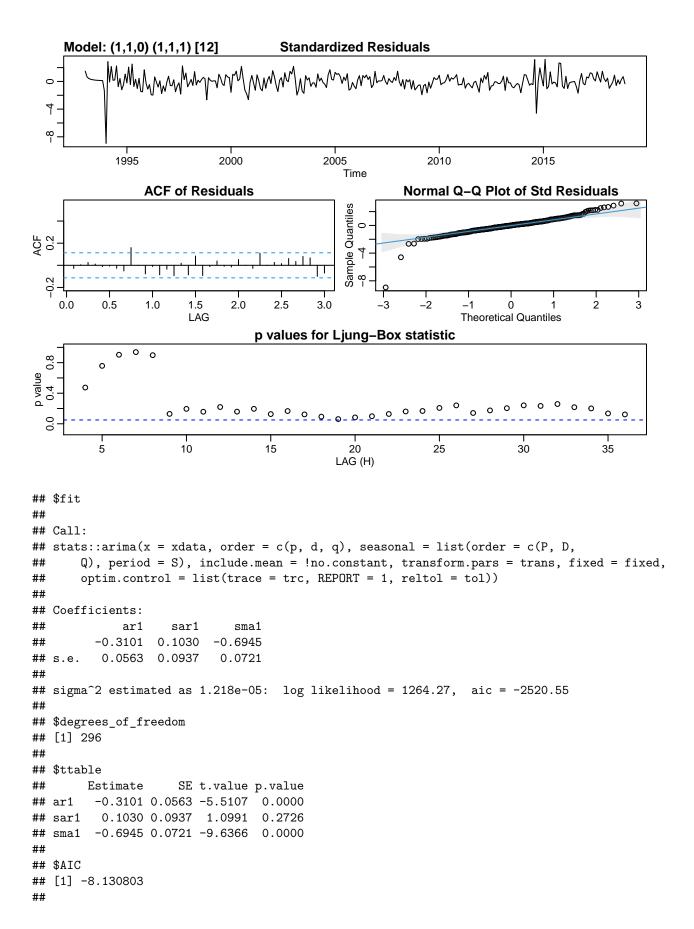
```
## initial value -5.492400
## iter
         2 value -5.587250
## iter
         3 value -5.645844
## iter
         4 value -5.649114
## iter
         5 value -5.653614
         6 value -5.658430
## iter
         7 value -5.659765
## iter
## iter
         8 value -5.660295
## iter
         9 value -5.660350
       10 value -5.660404
## iter
        11 value -5.660436
## iter
## iter
        12 value -5.660548
## iter
        13 value -5.660569
## iter
        14 value -5.660599
## iter
        15 value -5.660613
## iter
        16 value -5.660616
## iter
        16 value -5.660616
## iter 16 value -5.660616
## final value -5.660616
## converged
## initial value -5.643060
## iter
         2 value -5.644716
## iter
         3 value -5.646227
## iter
         4 value -5.647175
## iter
         5 value -5.647271
         6 value -5.647276
## iter
## iter
         7 value -5.647279
## iter
          8 value -5.647283
## iter
         9 value -5.647288
## iter 10 value -5.647290
## iter 10 value -5.647290
## final value -5.647290
## converged
```



```
##
## $AICc
## [1] -8.123946
##
## $BIC
## [1] -8.064684
\# ttable says mal coeff has highest p-value. removing this (model trimming):
sarima(log_ts_dat, p = 1, d = 1, q = 0, P = 1, D = 1, Q = 1, S = 12) #AICc -8.13055
## initial value -5.492400
## iter 2 value -5.615705
## iter 3 value -5.649834
## iter 4 value -5.652797
## iter 5 value -5.660080
## iter 6 value -5.660508
## iter 7 value -5.660561
## iter 8 value -5.660570
## iter 8 value -5.660571
## final value -5.660571
## converged
## initial value -5.643116
## iter 2 value -5.645066
## iter 3 value -5.646830
## iter 4 value -5.647172
## iter 5 value -5.647260
       6 value -5.647281
## iter
## iter
       7 value -5.647281
## iter 7 value -5.647281
## iter 7 value -5.647281
```

## final value -5.647281

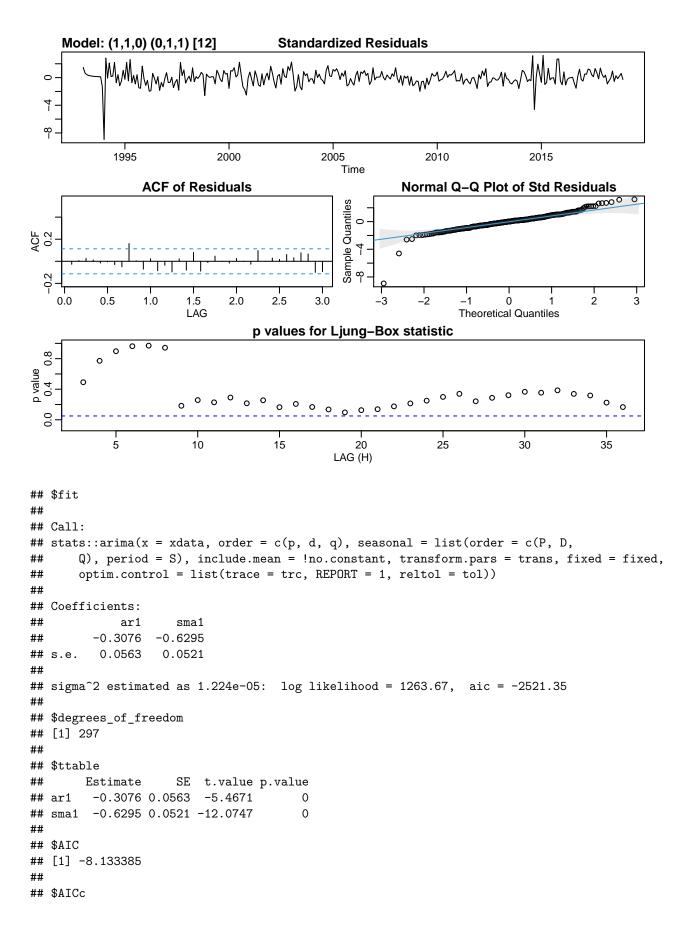
## converged



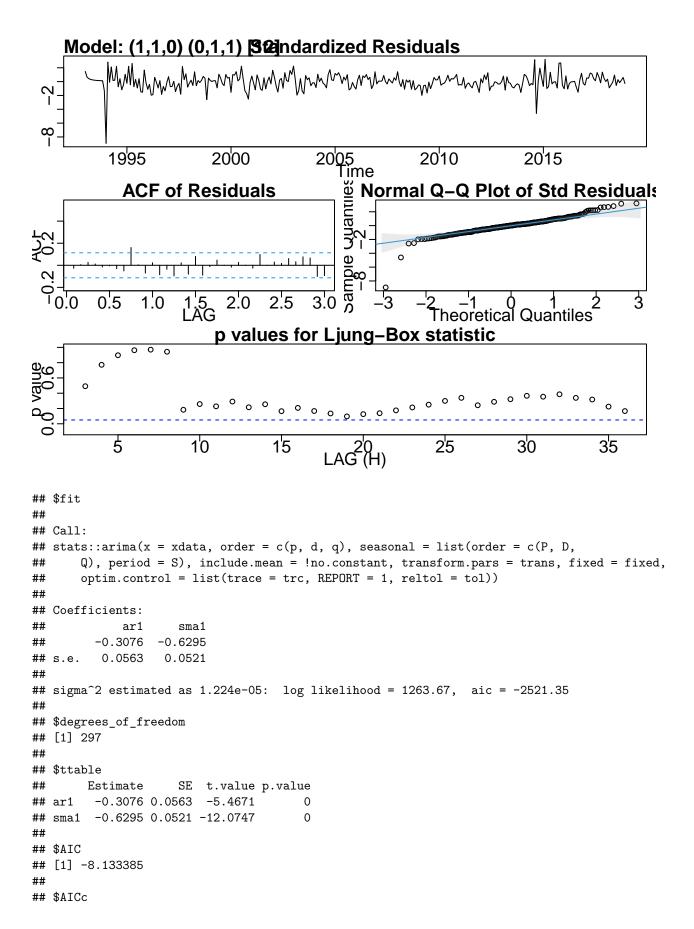
```
## $AICc
## [1] -8.13055
##
## $BIC
## [1] -8.083055
# ttable says sar1 coeff has highest p-value. removing this:
sarima(log_ts_dat, p = 1, d = 1, q = 0, P = 0, D = 1, Q = 1, S = 12) #AICc -8.133259
## initial value -5.482214
## iter 2 value -5.637939
## iter 3 value -5.652411
## iter 4 value -5.656510
## iter 5 value -5.657397
## iter 6 value -5.657437
## iter 7 value -5.657437
## iter 7 value -5.657437
## iter 7 value -5.657437
## final value -5.657437
## converged
## initial value -5.644718
## iter 2 value -5.645257
## iter 3 value -5.645275
## iter 4 value -5.645275
## iter 4 value -5.645275
## iter 4 value -5.645275
```

## final value -5.645275

## converged



```
## [1] -8.133259
##
## $BIC
## [1] -8.097574
\#jpeg("diagnostic\_plots.jpg", width = 1150, height = 700)
par(cex.lab = 1.6, cex.main = 1.8, cex.axis = 1.6)
sarima(log_ts_dat, p = 1, d = 1, q = 0, P = 0, D = 1, Q = 1, S = 12)
## initial value -5.482214
## iter 2 value -5.637939
## iter 3 value -5.652411
## iter 4 value -5.656510
## iter 5 value -5.657397
## iter 6 value -5.657437
## iter 7 value -5.657437
## iter 7 value -5.657437
## iter 7 value -5.657437
## final value -5.657437
## converged
## initial value -5.644718
## iter 2 value -5.645257
## iter 3 value -5.645275
## iter 4 value -5.645275
## iter 4 value -5.645275
## iter 4 value -5.645275
## final value -5.645275
## converged
```



```
## [1] -8.133259
##
## $BIC
## [1] -8.097574
```

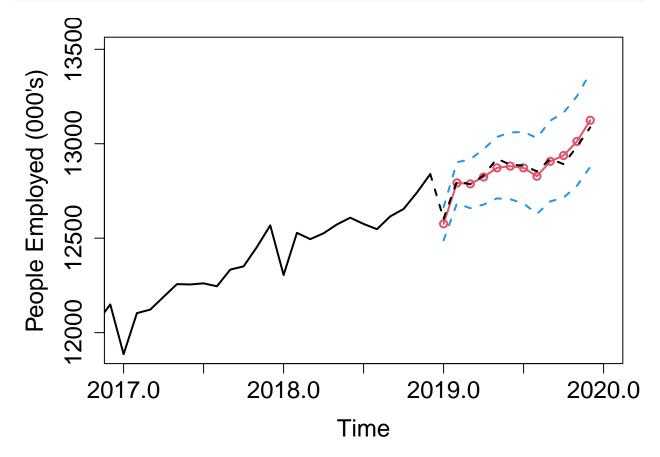
For the first model, the Ljung-Box statistic is not satisfactory at lag 20, and unnessed coefficients were present in the model. By trimming coefficients and finding the minimum bias-corrected AIC, the model selected was  $SARIMA(1,1,0)(0,1,1)_{12}$ .

We see a couple of outliers in the standardised residuals and Q-Q plot. The ACF of the residuals is as expected for normal data. The Ljung-Box statistic has a large enough p-value at lag 20 large enough to not reject the null hypothesis of the model exhibiting lack of fit.

Fit the model from above with lowest AIC

```
fit <- arima(log_ts_dat, c(1,1,0), seasonal = list(order = c(0,1,1), period = 12))
fore <- predict(fit, n.ahead = 12)</pre>
```

#### Display predictions:



### Assess the predictions

```
MAPE = 0
for (i in 1:12){
    MAPE = MAPE + abs((test_ts[i] - exp(fore$pred[i])) / test_ts[i])
}
MAPE = MAPE/12*100
MAPE

## [1] 0.1713158

Acc = 1-MAPE
Acc
```

## [1] 0.8286842

MAPE can be converted to accuracy by 1-MAPE. So we have a MAPE or 0.17% or an accuracy of prediction, or 99.837% accuracy for the 12 months' forcecast.

### Considering the possibility of adding a secondary time series

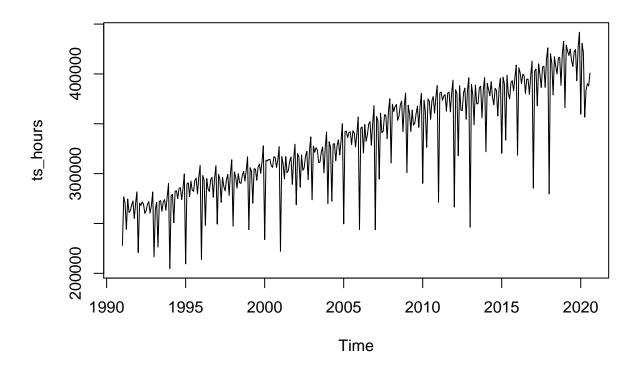
Load in the additional data:

```
dat2 <- read.csv("hours_data.csv", fileEncoding = 'UTF-8-BOM')
head(dat2)</pre>
```

```
##
     Observation.times Time.series.values
## 1
                Jan-91
                                  228022.4
## 2
                Feb-91
                                  276727.5
                Mar-91
## 3
                                  270555.0
## 4
                Apr-91
                                  244135.2
## 5
                May-91
                                  274548.9
## 6
                Jun-91
                                  261039.1
```

Create a time series object from the data and plot.

```
ts_hours <- ts(dat2[, 2], start = c(1991, 1), end = c(2020, 8), frequency = 12)
plot.ts(ts_hours)
```

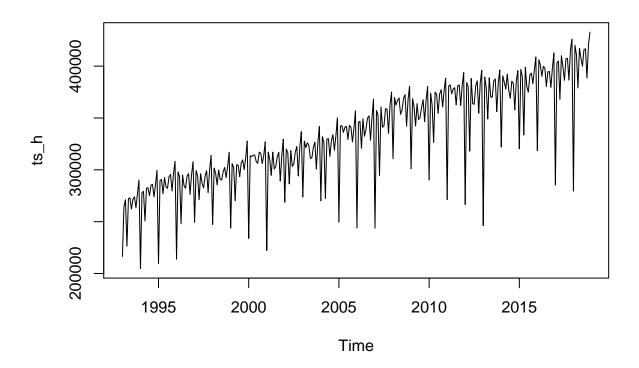


Cut the ts down to subset in appropriate time:

```
train_hours <- dat2[25:336,] # for training data
head(train_hours)</pre>
```

```
##
      Observation.times Time.series.values
## 25
                  Jan-93
                                    216445.1
## 26
                  Feb-93
                                    264350.3
  27
                  Mar-93
                                    271123.6
                  Apr-93
                                    226351.2
##
   28
## 29
                  May-93
                                    272067.0
## 30
                  Jun-93
                                    272984.9
```

```
ts_h \leftarrow ts(train_hours[, 2], start = c(1993, 1), end = c(2018, 12), frequency = 12) # ts compare to mod plot.ts(ts_h)
```



Plot a scatter plot comparing employed data and hours for those months.

pairs(cbind(Workers=ts\_dat, Hours=ts\_h))

