ARMA-GARCH

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1 Data

1.1 Data Source

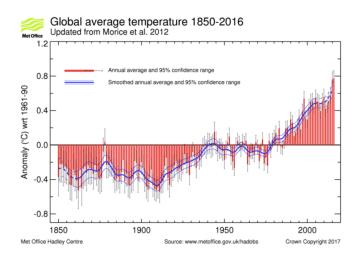
HadCRUT4 is a gridded dataset of global historical surface temperature anomalies relative to a 1961-1990 reference period. Data are available for each month since January 1850, on a 5 degree grid. The dataset is a collaborative product of the Met Office Hadley Centre and the Climatic Research Unit at the University of East Anglia.

 $url = \texttt{http://www.metoffice.gov.uk/hadobs/hadcrut4/data/current/time_series/HadCRUT.} \\ 4.5.0.0.monthly_ns_avg.txt.$

1.2 Brief Data Description

The gridded data are a blend of the CRUTEM4 land-surface air temperature dataset and the HadSST3 sea-surface temperature (SST) dataset. The dataset is presented as an ensemble of 100 dataset realisations that sample the distribution of uncertainty in the global temperature record given current understanding of non-climatic factors affecting near-surface temperature observations. This ensemble approach allows characterisation of spatially and temporally correlated uncertainty structure in the gridded data, for example arising from uncertainties in methods used to account for changes in SST measurement practices, homogenisation of land station records and the potential impacts of urbanisation. The HadCRUT4 data are neither interpolated nor variance adjusted.

Below is a graph provided by www.metoffice.gov.uk:



1.3 Loading the Data into R

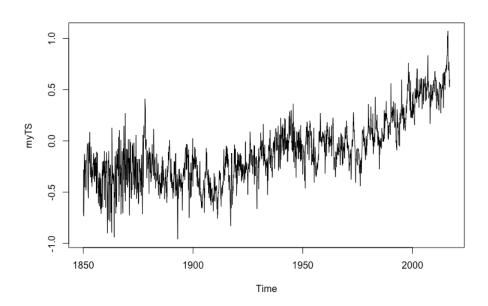
For our purpose, we are not interested in the spatial distribution, so we only use the *Global Mean*.

Loading the data into R:

```
library(curl)
tmpf <- tempfile()
curl_download(url, tmpf)
gtemp <- read.table(tmpf)[, 1:2]
temp = gtemp$V2[1:2004]</pre>
```

We only use the first two columns: time and monthly global mean. We pick the first 2004 observations, which is monthly data from 1850 \sim 2016, so that the data has complete periods.

The time series of temp is shown below:



2 Dealing with the time series

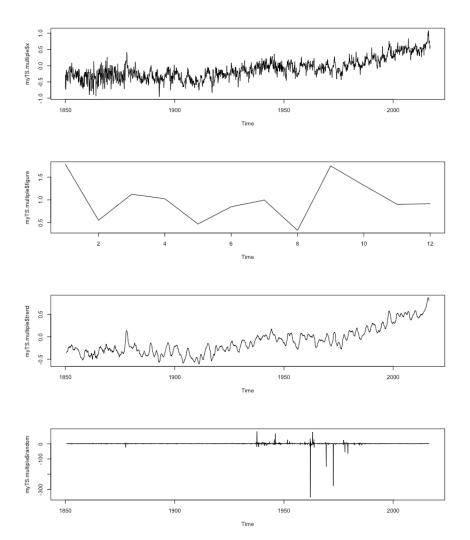
2.1 Decompose

The first thing we might want to do is to wipe out the seasonal component. There're many approaches to do it in R. We choose to use the decompose() provided by $\{stats\}$.

Looking at the time series, we think a additive model should be appropriate.

We can quickly checked that multiplicative model is not appropriate:





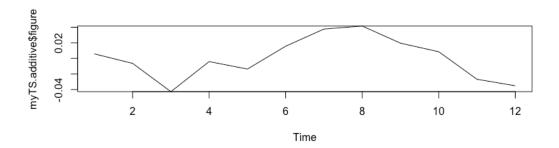
Obviously the random term is not easy to be modelled.

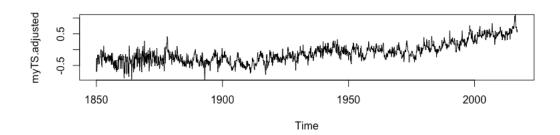
Therefore, decompose the time series with additive method in the following code:

```
library(TSA)
myTS = ts(as.numeric(temp), start = c(1850, 1), frequency = 12)
myTS.additive = decompose(myTS)
myTS.adjusted = myTS.additive$x - myTS.additive$seasonal
```

The myTS.adjusted now is free of seasonal components. The following graph illustrate the components.

Figure 2: additive model: seasonal component, adjusted series





Obviously, the time seires is not stationary. For simplicity, use diff().

dtemp = diff(myTS.adjusted)

> adf.test(dtemp)

Augmented Dickey-Fuller Test

data: dtemp

Dickey-Fuller = -16.175, Lag order = 12, p-value = 0.01

alternative hypothesis: stationary

So we can convince ourself that the time series dtemp is stationary.

2.2 Build a ARMA(p, q) for dtemp

For simplicity, we use auto.arima() from $\{forecast\}$ to judge the order.

library(forecast)

> auto.arima(dtemp> auto.arima(dtemp)

Series: dtemp

ARIMA(2,0,4)(2,0,1)[12] with non-zero mean

Coefficients:

ar1 ar2 ma1 ma2 ma3 ma4 sar1 sar2 sma1 mean

```
0.5041 0.3585 -1.0458 -0.1600 0.1362 0.0780 0.7619 0.0779 -0.7884 5e-04
s.e. 0.2431 0.2142
                       0.2423
                                0.3481 0.1082 0.0257 0.0926 0.0248
sigma^2 estimated as 0.01428: log likelihood=1417.03
AIC=-2812.05
              AICc=-2811.92
                               BIC=-2750.43
# first find the possible orders might be arma(2, 1) or arma(2, 4)
arma21.dtemp = arima(dtemp, c(2, 0, 1))
arma24.dtemp = arima(dtemp, c(2, 0, 4))
# next examine the two models' residuals
> auto.arima(arma21.dtemp$residuals)
Series: arma21.dtemp$residuals
ARIMA(2,0,3)(2,0,1)[12] with zero mean
Coefficients:
         ar1
                  ar2
                           ma1
                                   ma2
                                           ma3
                                                  sar1
                                                          sar2
                                                                   sma1
      1.5859 -0.6762 -1.6089
                                0.6706 0.0506 0.7454 0.0748 -0.7785
s.e. 0.1156
              0.1065
                        0.1169 0.1264
                                        0.0297
                                                0.1209 0.0258
                                                                 0.1181
sigma<sup>2</sup> estimated as 0.01429:
                               log likelihood=1415.92
AIC=-2813.83
               AICc=-2813.74
                               BIC=-2763.41
> auto.arima(arma24.dtemp$residuals)
Series: arma24.dtemp$residuals
ARIMA(0,0,0) with zero mean
sigma^2 estimated as 0.01435: log likelihood=1408.31
AIC=-2814.63
               AICc=-2814.62
                               BIC=-2809.02
Therefore it should be appropriate to choose arma(2, 4) for series dtemp.
my.arma = arma24.dtemp
> my.arma
Call:
arima(x = dtemp, order = c(2, 0, 4))
Coefficients:
                                           ma3
                                                        intercept
        ar1
                 ar2
                          ma1
                                   ma2
                                                   ma4
                                                            5e-04
      0.5204 0.3247
                      -1.0536
                               -0.1269
                                        0.1168 0.0755
s.e. 0.2733 0.2356
                       0.2727
                                0.3828
                                        0.1117
                                                0.0265
                                                            2e-04
sigma^2 estimated as 0.01435: log likelihood = 1407.67, aic = -2801.34
```

0.0910

2e-04

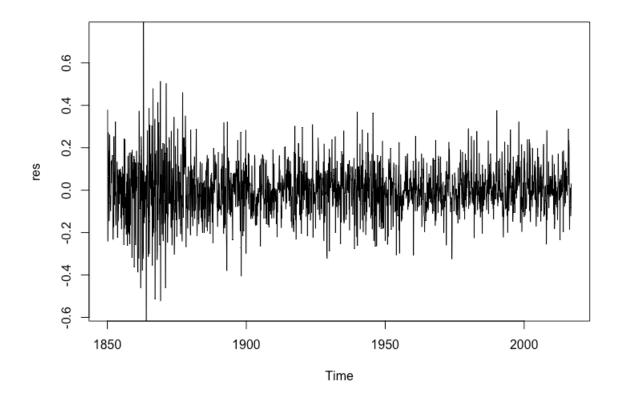
```
res = my.arma$residuals
> Box.test(res, type = 'Ljung-Box')
Box-Ljung test
data: res
X-squared = 0.0058853, df = 1, p-value = 0.9388
```

The ARMA(2,4) success in making the arma residual independent.

2.3 GARCH

One can be pretty satisfied with the results right now. Yet Looking at the plot of arma residual:

Figure 3: arma residual



It makes us wondering if it has ARCH effect. Using arch.test from $\{aTSA\}$ to perform the LM-Test:

library(aTSA)

> arch.test(my.arma)

ARCH heteroscedasticity test for residuals

0

0

alternative: heteroscedastic

Portmanteau-Q test:

	order	PQ	p.value		
[1,]	4	99.4	0		
[2,]	8	116.9	0		
[3,]	12	411.5	0		
[4,]	16	475.6	0		
[5,]	20	495.9	0		
[6,]	24	755.8	0		
${\tt Lagrange-Multiplier\ test:}$					
	order	LM p	o.value		
[1,]	4	1418	0		
[2,]	8	697	0		
[3,]	12	418	0		
[4,]	16	212	0		

20

24

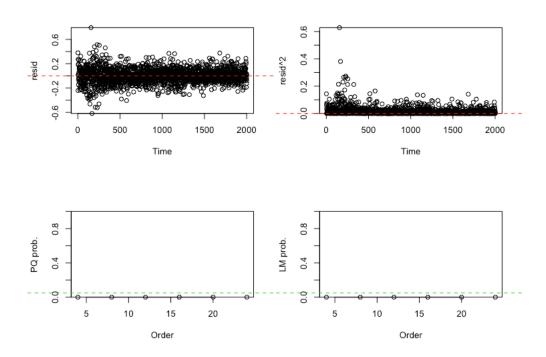
168

137

[5,]

[6,]

Figure 4: arma residual



It supports our doubt. Therefore we have the motive to build a GARCH model. We decide to use the package rugarch.

A standard GARCH model has the following variance equation:

$$\sigma_t^2 = (w + \sum_{j=1}^m \zeta_j v_{jt}) + \sum_{j=1}^q \alpha_j \varepsilon_{t-j}^2 + \sum_{j=1}^p \beta_j \sigma_{t-j}^2$$

So we wrote a function:

```
library(rugarch)
my_sGARCH_test <- function(p, q, m, n, ts.data = res)
{
# I use include.mean = FALSE after trying TRUE
# to find out insignificance
    myspec=ugarchspec(variance.model = list(model = "sGARCH", garchOrder = c(p, q)),
    mean.model = list(armaOrder = c(m, n), include.mean = FALSE),
    distribution.model = "norm")
    myfit=ugarchfit(myspec,data=ts.data, solver="solnp")
    return(myfit)
}</pre>
```

After trying a few times from (1,0), (0,0,0) to (5,5), (4,0,4), GARCH(1,1), ARIMA(2,0,3) is the most satisfying model. Here we realize that using GARCH model, the order of ARIMA might changes.

```
fit1 = my_sGARCH_test(1, 1, 2, 3, dtemp)
```

> fit1

```
*----*

* GARCH Model Fit *

*----*
```

Conditional Variance Dynamics

GARCH Model : sGARCH(1,1)
Mean Model : ARFIMA(2,0,3)

Distribution : norm

Optimal Parameters

Estimate Std. Error t value Pr(>|t|)
ar1 -0.090935 0.015226 -5.9724 0.000000
ar2 0.754333 0.015069 50.0583 0.000000
ma1 -0.391906 0.007313 -53.5917 0.000000
ma2 -0.863257 0.000130 -6647.5909 0.000000

ma3 0.295634 0.007800 37.9010 0.000000 omega 0.000082 0.000034 2.3956 0.016595 alpha1 0.023026 0.003614 6.3706 0.000000

beta1 0.970594 0.004705 206.2925 0.000000

Robust Standard Errors:

	Estimate	Std. Error	t value	Pr(> t)
ar1	-0.090935	0.017031	-5.3393	0.000000
ar2	0.754333	0.017312	43.5726	0.000000
ma1	-0.391906	0.002647	-148.0378	0.000000
ma2	-0.863257	0.000144	-6000.9074	0.000000
ma3	0.295634	0.002903	101.8443	0.000000
omega	0.000082	0.000036	2.2838	0.022382
alpha1	0.023026	0.003636	6.3329	0.000000
beta1	0.970594	0.003734	259.9413	0.000000

LogLikelihood: 1512.336

Information Criteria

Akaike -1.5021 Bayes -1.4797 Shibata -1.5021 Hannan-Quinn -1.4939

Weighted Ljung-Box Test on Standardized Residuals

statistic p-value
Lag[1] 0.3742 0.5407
Lag[2*(p+q)+(p+q)-1][14] 4.5219 1.0000
Lag[4*(p+q)+(p+q)-1][24] 13.5547 0.3236
d.o.f=5

HO : No serial correlation

Weighted Ljung-Box Test on Standardized Squared Residuals

statistic p-value
Lag[1] 31.06 2.506e-08
Lag[2*(p+q)+(p+q)-1][5] 37.84 1.904e-10
Lag[4*(p+q)+(p+q)-1][9] 51.28 4.433e-13
d.o.f=2

Weighted ARCH LM Tests

ARCH Lag[3] 0.004507 0.500 2.000 9.465e-01 ARCH Lag[5] 11.085543 1.440 1.667 3.565e-03 ARCH Lag[7] 20.592184 2.315 1.543 4.509e-05

Nyblom stability test

Joint Statistic: 2.6294 Individual Statistics:

0.25019 ar1

ar2 0.61592

ma1 0.19513

ma20.24162

ma3 0.07259

omega 0.10095

alpha1 0.42035

beta1 0.20019

Asymptotic Critical Values (10% 5% 1%) Joint Statistic: 1.89 2.11 2.59

Individual Statistic: 0.35 0.47 0.75

Sign Bias Test

t-value prob sig

0.3363 7.367e-01 Sign Bias

Negative Sign Bias 3.6815 2.380e-04 *** Positive Sign Bias 4.4421 9.396e-06 ***

Joint Effect 33.2983 2.786e-07 ***

Adjusted Pearson Goodness-of-Fit Test:

group statistic p-value(g-1) 1 20 57.32 1.019e-05 2 30 63.59 2.167e-04 85.01 2.883e-05 102.77 1.103e-05 85.01 3 40

Elapsed time : 0.365526

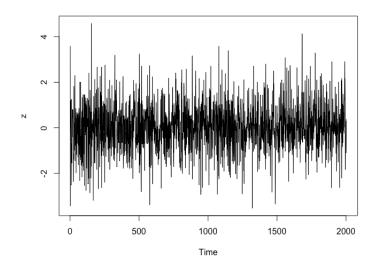
50

4

residuals(fit)Substract the standardized (w.r.t. the variance model) residuals z, which is z =sigma(fit)

z = residuals(fit1) / sigma(fit1) plot.ts(z)

Figure 5: z

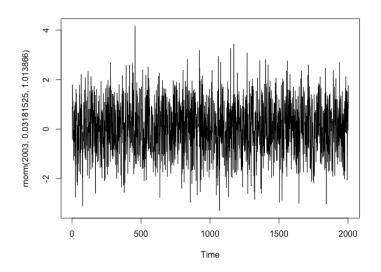


We can compute the mean and variance of z:

```
> mean(z)
[1] 0.03181525
> var(z)
[1] 1.013866
> length(z)
[1] 2003
> plot.ts(rnorm(2003, 0.03181525, 1.013866))
```

And then, just for fun, plot a normal sample series with the same parameters:

Figure 6: simulation of using rnorm



At least a human can't distinguish between them anymore. But what we most care about is whether standardized squared residuals can pass the LM-Test (substracting from the above long result):

Weighted ARCH LM Tests

ARCH Lag[3] 0.004507 0.500 2.000 9.465e-01 ARCH Lag[5] 11.085543 1.440 1.667 3.565e-03 ARCH Lag[7] 20.592184 2.315 1.543 4.509e-05

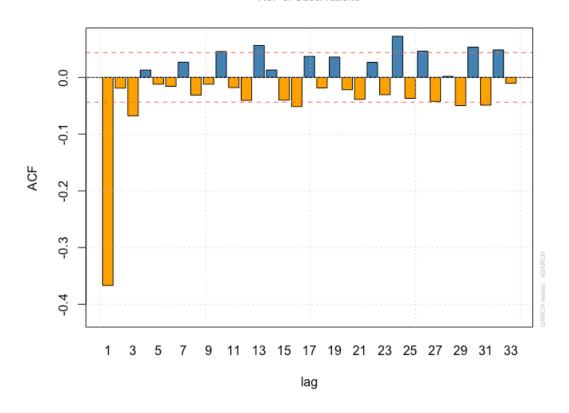
We can see our model successfully wipes out ARCH effect at small lags, but fails at larger lags. This is the best 'sGARCH' can give with respect to our data. Maybe we should try long memory models instead of diff() at first.

Graphical Diagnostics:

The original series' acf is shown below(the difference of temperature):

Figure 7: acf(dtemp)

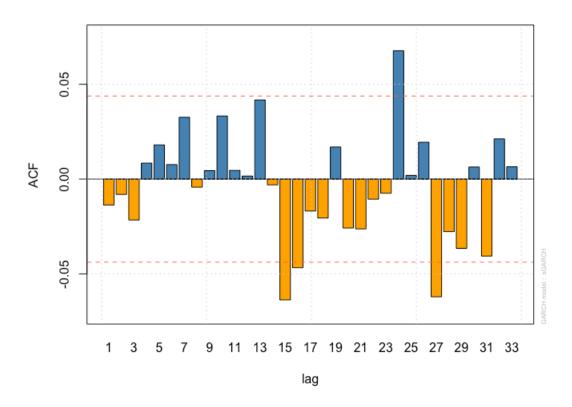
ACF of Observations



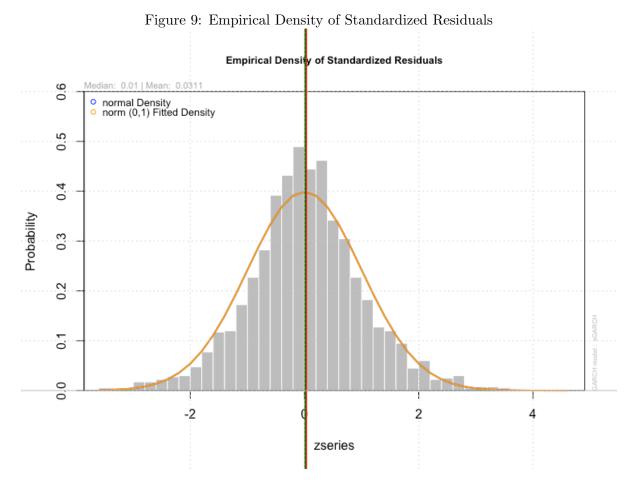
The acf of standardized residuals:

Figure 8: acf(z)

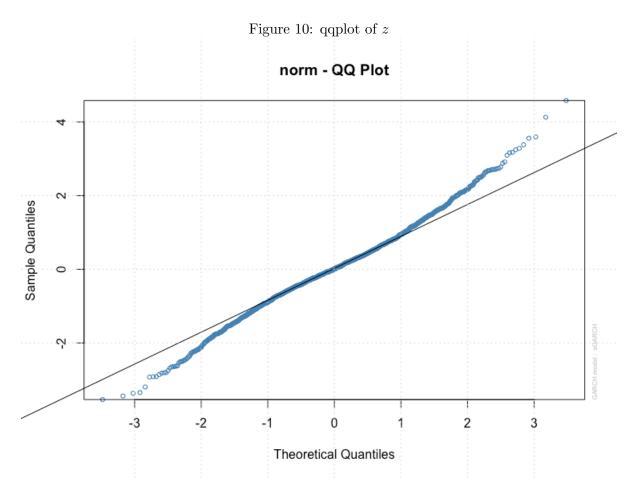
ACF of Standardized Residuals



This is a prove that we should have used a long memory model. The $\it Empirical\ Density\ of\ Standardized\ Residuals\ compared\ to\ normal\ distribution:$

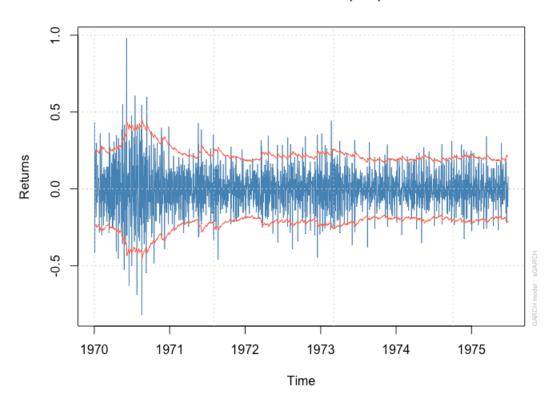


qqplot:



We can see that z is skewed, which might be correlated with long lags' heterodasticity. Series with 2 Conditional SD Superimposed:

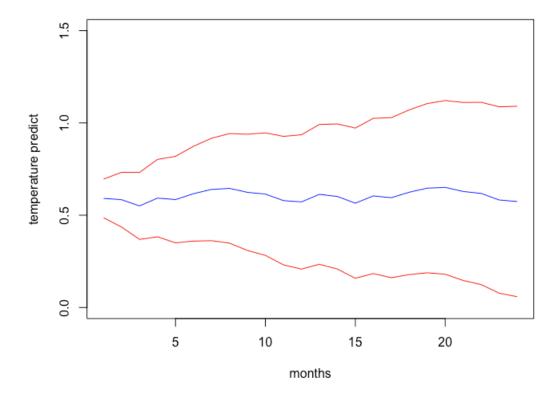
Series with 2 Conditional SD Superimposed



In this plot, we can find the reason that GARCH(1, 1) can't completely wipe out heterodasticity, because there is a huge anomaly that can't be explained by an arma on variance.

Forecast

Figure 11: forecast



Conclusion: 'sGARCH' have improved the model in the intuitive sense and to some extent fix the heterodasticity.