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Eg. food expenditure is known to vary much more at higher levels of income than at lower levels of income, the level of profits tends to vary more across large firms than across small firms)

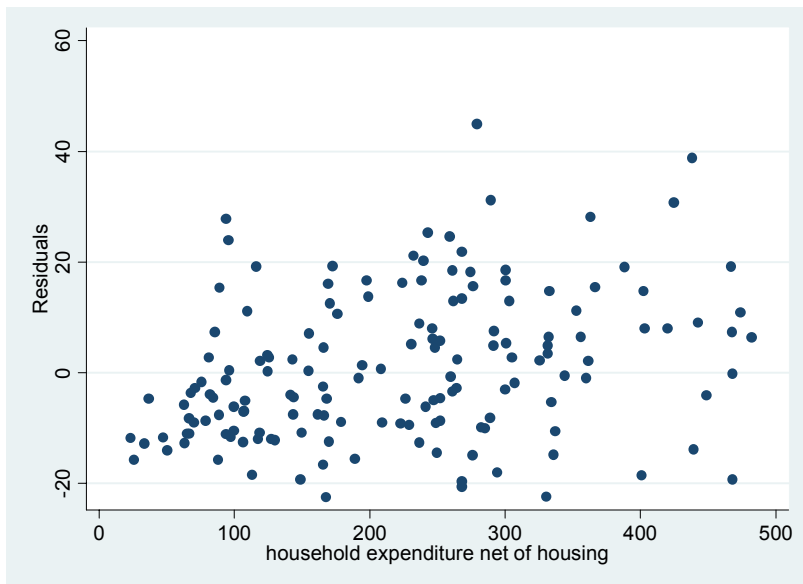
Example: the data set food.dta contains information on food expenditure and income. A graph of the *residuals* from a regression of food spending on total household expenditure clearly that the residuals tend to be more spread out at higher levels of income – this is typical pattern associated with heteroskedasticity.

```
. reg food expnethsum
```

Source	SS	df	MS	Number of obs = 200		
Model	22490.0823	1	22490.0823	F(1, 198)	=	107.19
Residual	41544.8096	198	209.822271	Prob > F	=	0.0000
-----				R-squared	=	0.3512
-----				Adj R-squared	=	0.3479
Total	64034.8918	199	321.783376	Root MSE	=	14.485

food	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
expnethsum	.0355189	.0034308	10.35	0.000	.0287534	.0422844
_cons	28.55002	1.56964	18.19	0.000	25.45466	31.64537

```
. predict res, resid
. two (scatter res expnet if expnet<500)
```



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(intuitively, if all observations are distributed unevenly about the regression line then OLS is unable to distinguish the “quality” of the observations - observations further away from the regression line should be given less weight in the calculation of the standard errors (since they are more unreliable) but OLS can't do this, so the standard errors are biased).

Testing for Heteroskedasticity

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In absence of Heteroskedasticity there should be no obvious pattern to the **spread** of the residuals, so useful to plot the residuals against the X variable thought to be causing the problem,

- assuming you know which X variable it is (often difficult)

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Again assuming know which variable is causing the problem then can test formally whether the residual spread varies with values of the suspect X variable.

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- i) Order the data by the size of the X variable and split the data into 2 equal sub-groups (one high variance the other low variance)
- ii) Drop the middle “c” observations where c is approximately 30% of your sample

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- iv) Compute

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- vii) If estimated $F > F_{critical}$, **reject** null of no heteroskedasticity (intuitively the residuals from the high variance sub-sample are much larger than the residuals from the high variance sub-sample)

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A test that is valid asymptotically (ie in large samples) that does not rely on knowing which variable is causing the problem is the Breusch-Pagan test

Given

$$Y_i = a + b_1X_1 + b_2X_2 + u_i \quad (1)$$

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or

compute $N * R^2_{auxillary} \sim \chi^2(k-1)$

If F or $N * R^2_{auxillary} >$ respective critical values reject null of no heterosked.

Example: Breusch-Pagan Test of Heteroskedasticity

The data set smoke.dta contains information on the smoking habits, wages age and gender of a cross-section of individuals

```

. u smoke.dta                /* read in data */

. reg lhw age age2 female smoke

-----+-----
Source |      SS      df      MS                Number of obs =      7970
-----+-----+-----+-----                F( 4, 7965) =     284.04
Model |  304.964893      4   76.2412233            Prob > F      =    0.0000
Residual | 2137.94187   7965   .268417059            R-squared      =    0.1248
-----+-----+-----+-----            Adj R-squared =    0.1244
Total | 2442.90677   7969   .306551232            Root MSE     =    .51809

-----+-----
      lhw |      Coef.   Std. Err.      t    P>|t|     [95% Conf. Interval]
-----+-----+-----+-----+-----+-----
      age |   .0728466   .0031712    22.97   0.000   .0666301   .0790631
     age2 |  -.000847   .0000382   -22.17   0.000  -.0009219  -.0007721
    female |  -.2583456   .0116394   -22.20   0.000  -.2811618  -.2355294
     smokes | -.1501679   .0128866   -11.65   0.000  -.1754291  -.1249068
      _cons |   .8732505   .062907    13.88   0.000   .7499363   .9965646
-----+-----

/* save residuals */
. predict reshat, resid

. g reshat2=reshat^2                /* square them */

/* regress square of residuals on all original rhs variables */

. reg reshat2 age age2 female smoke

-----+-----
Source |      SS      df      MS                Number of obs =      7970
-----+-----+-----+-----                F( 4, 7965) =      6.59
Model |  13.2179958      4   3.30449895            Prob > F      =    0.0000
Residual | 3996.90523   7965   .501808566            R-squared      =    0.0033
-----+-----+-----+-----            Adj R-squared =    0.0028
Total | 4010.12323   7969   .503215363            Root MSE     =    .70838

```

reshat2	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]
age	.0012546	.004336	0.29	0.772	-.0072452 .0097544
age2	.0000252	.0000522	0.48	0.630	-.0000772 .0001276
female	.0022702	.0159145	0.14	0.887	-.0289264 .0334668
smokes	-.0174587	.0176199	-0.99	0.322	-.0519983 .0170808
_cons	.1766929	.0860128	2.05	0.040	.0080854 .3453004

Breusch-Pagan test is $N \cdot R^2$

```
. di 7970*.0033
26.301
```

which is chi-squared $k-1$ degrees of freedom (4 in this case) and the critical value is 9.48. So estimated value exceeds critical value

Similarly the F test for goodness of fit in stata output in the top right corner is test for joint significance of all the rhs variables in this model (excluding the constant)

From F tables, $F_{critical}^{5\% \text{ level}}(4, 7970) = 2.37$

So estimated $F = 6.59 > F_{critical}$, so **reject** null of **no** heteroskedasticity
Or could use Stata's version of the Breusch-Pagan test

What to do if heteroskedasticity present?

1. Try different functional form

Sometimes taking logs of dependent or explanatory variable can reduce the problem

```

. reg food expnethsum if exp<1000

```

Source	SS	df	MS			
Model	21179.4196	1	21179.4196	Number of obs =	192	
Residual	36583.9436	190	192.547072	F(1, 190) =	110.00	
Total	57763.3632	191	302.425986	Prob > F =	0.0000	
				R-squared =	0.3667	
				Adj R-squared =	0.3633	
				Root MSE =	13.876	

	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
expnethsum	.0504532	.0048106	10.49	0.000	.0409641	.0599423
_cons	24.60655	1.770093	13.90	0.000	21.11499	28.0981

```

. bpagan expn

```

Breusch-Pagan LM statistic: 7.54351 Chi-sq(1) P-value = .006

The Breusch-Pagan test indicates the presence of heteroskedasticity (estimated chi-squared value > critical value). This means the standard errors, t statistics etc are biased

If use the log of the dependent variable rather than in levels

```

. g lfood=log(food)
. reg lfood expnethsum

```

Source	SS	df	MS			
Model	14.6377436	1	14.6377436	Number of obs =	200	
Residual	31.1642937	198	.157395423	F(1, 198) =	93.00	
Total	45.8020374	199	.230160992	Prob > F =	0.0000	
				R-squared =	0.3196	
				Adj R-squared =	0.3162	
				Root MSE =	.39673	

	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
expnethsum	.0009062	.000094	9.64	0.000	.0007209	.0010915
_cons	3.290222	.0429903	76.53	0.000	3.205444	3.374999

```

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```

Breusch-Pagan LM statistic: .4280017 Chi-sq(1) P-value = .513

2. Drop “Outliers”

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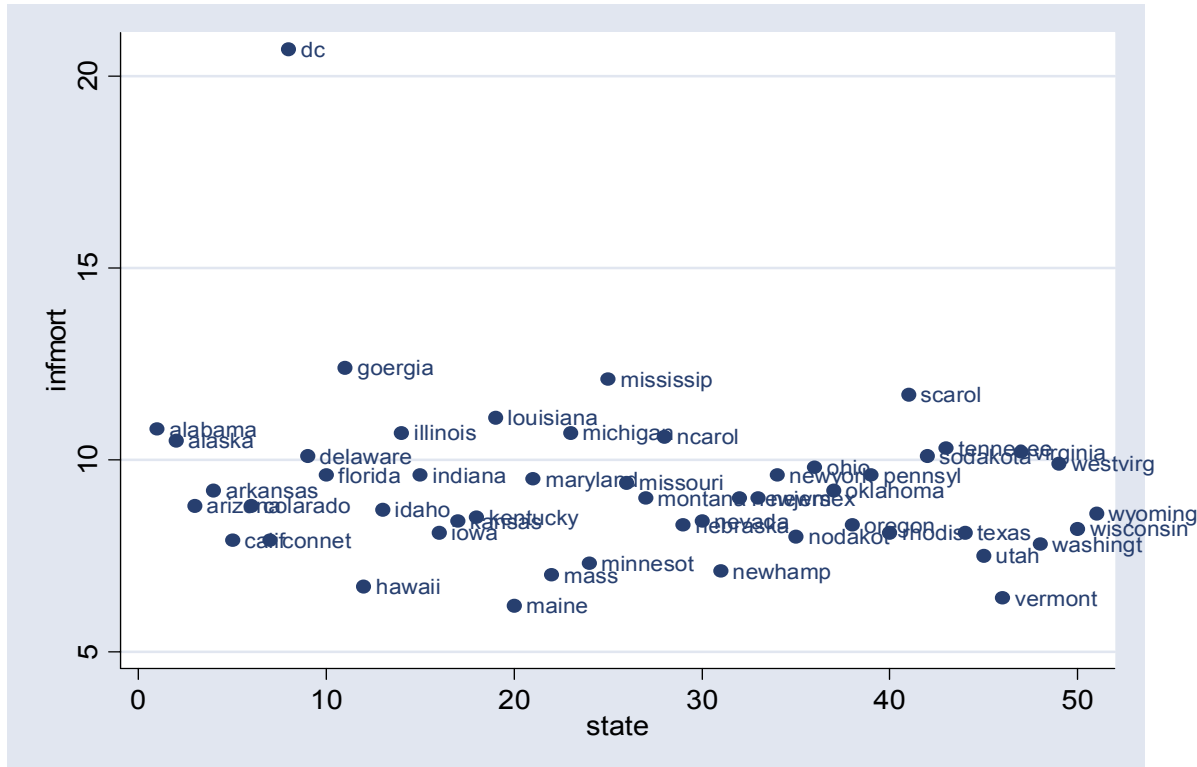
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Often these observations may be genuine – in which case you should not drop them – but sometimes they may be the result of measurement error or miscoding in which case you may have a case for dropping them.

Example

The data `infmort.dta` gives infant mortality for 51 U.S. states along with the number of doctors per capita in each state. A graph of infant mortality against number of doctors clearly shows that Washington D.C. is something of an outlier (it has lots of doctors but also a very high infant mortality rate)

```
. twoway (scatter infmort state, mlabel(state)), ytitle(infmort) ylabel(, labels) xtitle(state)
```



A regression of infant mortality on (the log of) doctor numbers for all 51 observations suffers from heteroskedasticity

```
. reg infmort ldocs
```

Source	SS	df	MS	Number of obs =	51
-----+-----				F(1, 49) =	4.08
Model	17.7855153	1	17.7855153	Prob > F =	0.0488

Residual		213.461954	49	4.3563664		R-squared	=	0.0769
Total		231.247469	50	4.62494938		Adj R-squared	=	0.0581
						Root MSE	=	2.0872

infmort		Coef.	Std. Err.	t	P> t	[95% Conf. Interval]		
ldocs		2.130049	1.054189	2.02	0.049	.0115765	4.248522	
_cons		-1.959674	5.572467	-0.35	0.727	-13.15797	9.238617	

. bpagan ldocs

Breusch-Pagan LM statistic: 67.14974 Chi-sq(1) P-value = 2.5e-16

However if the outlier is excluded then

. reg infmort ldocs if dc==0

Source		SS	df	MS		Number of obs	=	50
Model		9.49879378	1	9.49879378		F(1, 48)	=	5.13
Residual		88.8244081	48	1.8505085		Prob > F	=	0.0280
Total		98.3232019	49	2.00659596		R-squared	=	0.0966
						Adj R-squared	=	0.0778
						Root MSE	=	1.3603

infmort		Coef.	Std. Err.	t	P> t	[95% Conf. Interval]		
ldocs		-1.915912	.8456428	-2.27	0.028	-3.616191	-.2156336	
_cons		19.12582	4.448765	4.30	0.000	10.18098	28.07066	

. bpagan ldocs

Breusch-Pagan LM statistic: .0825086 Chi-sq(1) P-value = .7739

Can see that the problem of heteroskedasticity disappears – though the D.C. observation is genuine so you need to think carefully about the benefits of dropping it against the costs.

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Consider the term $\text{Var}(u_i/X) = 1/X_i^2 \text{Var}(u_i) = 1/X_i^2 * \sigma^2 X_i^2 = \sigma^2$

So the variance of this **is** constant for all observations in the data set

This means if we divide all the observations by $1/X_i$

$$Y_i = b_0 + b_1 X_i + u_i \quad (1)$$

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So the variance of this **is** constant for all observations in the data set

This means if we divide all the observations by $1/X_i$

$$Y_i = b_0 + b_1 X_i + u_i \quad (1)$$

becomes

$$Y_i / X_i = b_0 / X_i + b_1 X_i / X_i + u_i / X_i \quad (2)$$

2. Feasible GLS

If (and this is a big if) you think you know the exact functional form of the heteroskedasticity

eg you know that $\text{var}(u_i) = \sigma^2 X_1^2$ (and not say $\sigma^2 X_2^3$)

so that there is a common component to the variance, σ^2 , and a part that rises with the square of the level of the variable X_1

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and the estimates of b_0 and b_1 in (2) will not be affected by heterosked.

This is called a Feasible Generalised Least Squares Estimator (FGLS) and will be more efficient than OLS

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If not the “solution” may be much worse than OLS

Example

```
. reg hourpay age
```

Source	SS	df	MS			
Model	5207.03058	1	5207.03058	Number of obs =	12098	
Residual	473292.608	12096	39.1280264	F(1, 12096) =	133.08	
Total	478499.638	12097	39.5552317	Prob > F =	0.0000	
				R-squared =	0.0109	
				Adj R-squared =	0.0108	
				Root MSE =	6.2552	

hourpay	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
age	.0586134	.005081	11.54	0.000	.0486539	.0685729
_cons	6.168383	.2066433	29.85	0.000	5.763329	6.573437

```
. bpagan age
```

```
Breusch-Pagan LM statistic: 17.27396 Chi-sq( 1) P-value = 3.2e-05
```

Test suggests heteroskedasticity present

Suppose you decide that heteroskedasticity is given by $\text{var}(u_i) = \sigma^2 \text{Age}_i$

So transform variables by dividing by SQUARE ROOT of Age
(including the constant)

```
. g ha=hourpay/sqrt(age)
. g aa=age/sqrt(age)
. g ac=1/sqrt(age)          /* this is new constant term */
. reg ha aa ac, nocon
```

Source	SS	df	MS			
Model	22854.251	2	11427.1255	Number of obs =	12098	
Residual	12576.8073	12096	1.03974928	F(2, 12096) =	10990.27	
				Prob > F =	0.0000	
				R-squared =	0.6450	
				Adj R-squared =	0.6450	

Total		35431.0584	12098	2.92867072		Root MSE	=	1.0197
ha		Coef.	Std. Err.	t	P> t	[95% Conf. Interval]		
aa		.0932672	.0049446	18.86	0.000	.0835749	.1029594	
ac		4.813435	.184437	26.10	0.000	4.451908	5.174961	

If heteroskedastic assumption is correct these are the GLS estimates and should be preferred to OLS. If assumption is not correct they will be misleading.

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Again should only really do this in **large** samples

Example

```
. reg food expnethsum
```

Source	SS	df	MS			
Model	22490.0823	1	22490.0823	Number of obs =	200	
Residual	41544.8096	198	209.822271	F(1, 198) =	107.19	
Total	64034.8918	199	321.783376	Prob > F =	0.0000	
				R-squared =	0.3512	
				Adj R-squared =	0.3479	
				Root MSE =	14.485	

food	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
expnethsum	.0355189	.0034308	10.35	0.000	.0287534	.0422844
_cons	28.55002	1.56964	18.19	0.000	25.45466	31.64537

```
. reg food expnethsum, robust
```

Linear regression

```
Number of obs = 200
F( 1, 198) = 80.37
Prob > F = 0.0000
R-squared = 0.3512
Root MSE = 14.485
```

food	Coef.	Robust Std. Err.	t	P> t	[95% Conf. Interval]	
expnethsum	.0355189	.0039619	8.97	0.000	.0277059	.0433319
_cons	28.55002	1.549629	18.42	0.000	25.49412	31.60591

The Breusch-Pagan test indicates the presence of heteroskedasticity (estimated chi-squared value > critical value). This means the standard errors, t statistics etc are biased, so decide to fix up the standard errors using the white correction

Note that the OLS coefficients are unchanged, only the standard errors and t values change

Testing for Heteroskedasticity in Time Series Data (ARCH)

Sometimes what appears to be autocorrelation in time series data can be caused by heteroskedasticity

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Model (1) is called an Autoregressive Conditional Heteroskedasticity (ARCH) model

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i) Estimate the model $y_t = b_0 + b_1 X_t + u_t$ by OLS

ii) Save the estimated residuals, \hat{u}_t

iii) Square them, \hat{u}_t^2

iv) Regress the squared OLS residuals on their value lagged by 1 period and a constant

$$\hat{u}_t^2 = g_0 + g_1 \hat{u}_{t-1}^2 + v_t$$

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vi) If so adjust the standard errors using the white robust correction

Example: This example uses Wooldridge's (2000) data, (*stocks.dta*) on New York Stock Exchange price movements to test the efficient markets hypothesis.

EMH suggests that information on returns (ie the percentage change in the share price) in the week before should not predict the percentage change in this week's share price. A simple way to test this is to regress current returns on lagged returns.

```
. u shares
```

```
. reg return return1
```

Source	SS	df	MS			
Model	10.6866237	1	10.6866237	Number of obs =	689	
Residual	3059.73813	687	4.4537673	F(1, 687) =	2.40	
Total	3070.42476	688	4.46282668	Prob > F =	0.1218	
				R-squared =	0.0035	
				Adj R-squared =	0.0020	
				Root MSE =	2.1104	

return	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
return1	.0588984	.0380231	1.55	0.122	-.0157569	.1335538
_cons	.179634	.0807419	2.22	0.026	.0211034	.3381646

In this example it would appear that lagged returns have little power in predicting current price changes.

However it may be that the t value is influenced by heteroskedasticity in the variance of the residuals.

```
. predict reshat, resid
. g reshat2=reshat^2 /* ols residuals squared */
. g reshat2l=reshat[_n-1] /* lagged one period */
```

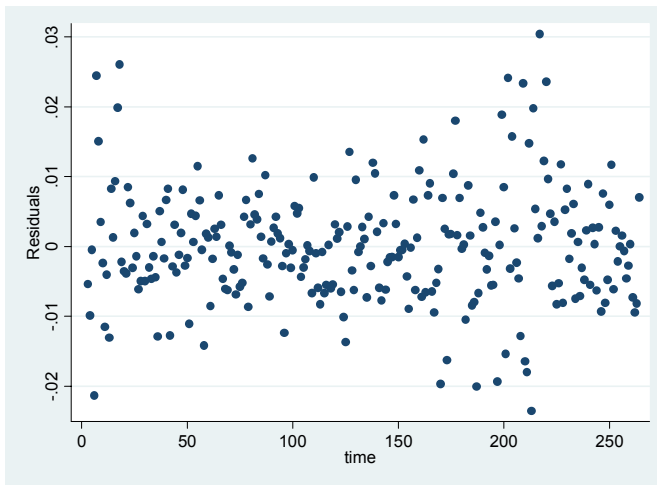
The graph of residuals over time, suggests heteroskedasticity may exist

```
Two (scatter reshat time , yline(0))
```

As does the Breusch-Pagan test

```
. bpagan return1
```

Breusch-Pagan LM statistic: 95.21722 Chi-sq(1) P-value = 1.7e-22



The ARCH test is found by a regression of the squared ols residuals on lagged values (in this case 1)

```
. reg return return1
. predict reshat, resid          /* save residuals */

. g reshat2=reshat^2            /* square residuals */
. g reshat21=reshat2[_n-1]     /* lag by 1 period */

. reg reshat2 reshat21
```

Source	SS	df	MS			
Model	10177.7088	1	10177.7088	Number of obs =	688	
Residual	79409.7826	686	115.757701	F(1, 686) =	87.92	
Total	89587.4914	687	130.403918	Prob > F =	0.0000	
				R-squared =	0.1136	
				Adj R-squared =	0.1123	
				Root MSE =	10.759	

reshat2	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
reshat21	.3370622	.0359468	9.38	0.000	.2664833	.4076411
_cons	2.947434	.4402343	6.70	0.000	2.083065	3.811802

Since estimated t value on lagged dependent variable is highly significant reject null of homoskedasticity.

Need to fix up the standard errors in the original regression.