

Lecture 2b: Linear Regression Diagnostics (VER 14.8-10)

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- Learning objectives

- ★ Assess assumptions linear regression
- ★ Identity outliers and influential observations
- ★ Develop approach linear reg. diagnostics

- Exercises

- ★ work on linear regression exercise 3

- Quiz #2

Example: WPC model (daisy2red.dta)

● VER example 14.12

- ★ outcome: wpc (interval from waiting period to conception)
- ★ predictors: parity, twin, dyst, rp, vag_disch, herd_size and herd_size2, calving_date (in months)
- ➔ also interactions: i.rp##i.vag_disch

● Final (candidate) model

```
. reg wpc hs100_ctr hs100_ctr_sq parity1 i.aut_calv i.twin i.dyst i.rp##i.vag_disch
```

Source	SS	df	MS	Number of obs	=	1,574
Model	296062.694	9	32895.8549	F(9, 1564)	=	13.22
Residual	3892027.86	1,564	2488.50886	Prob > F	=	0.0000
				R-squared	=	0.0707
				Adj R-squared	=	0.0653
Total	4188090.56	1,573	2662.48605	Root MSE	=	49.885

wpc	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
hs100_ctr	19.85708	2.163397	9.18	0.000	15.61361	24.10054
hs100_ctr_sq	11.13827	3.111145	3.58	0.000	5.035817	17.24073
parity1	1.13721	.8583103	1.32	0.185	-.5463501	2.82077
aut_calv						
yes	-8.263839	2.537751	-3.26	0.001	-13.24159	-3.286086
twin						
yes	20.68314	9.845165	2.10	0.036	1.37203	39.99425
dyst						
yes	11.70041	5.462576	2.14	0.032	.985666	22.41516
rp						
yes	5.98687	4.811976	1.24	0.214	-3.451734	15.42547
vag_disch						
yes	1.228196	7.161395	0.17	0.864	-12.81875	15.27514
rp#vag_disch						
yes#yes	22.85194	12.51605	1.83	0.068	-1.698056	47.40194
_cons	64.33029	2.634114	24.42	0.000	59.16352	69.49705

Evaluating major assumptions

● Homoscedasticity

```
. hettest

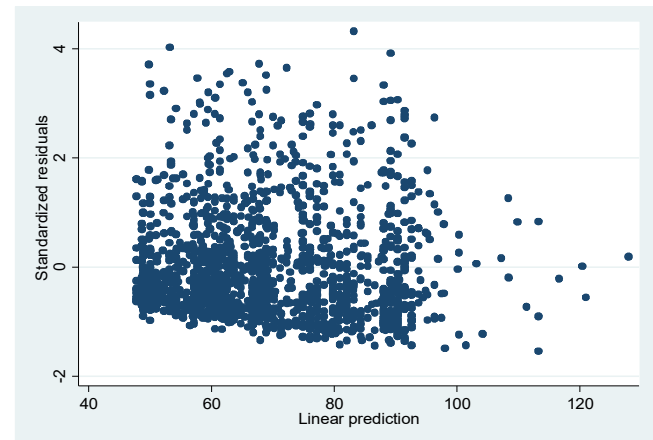
Breusch-Pagan / Cook-Weisberg test
for heteroskedasticity
    Ho: Constant variance
    Variables: fitted values of wpc

    chi2(1)      =    20.58
    Prob > chi2   =    0.0000

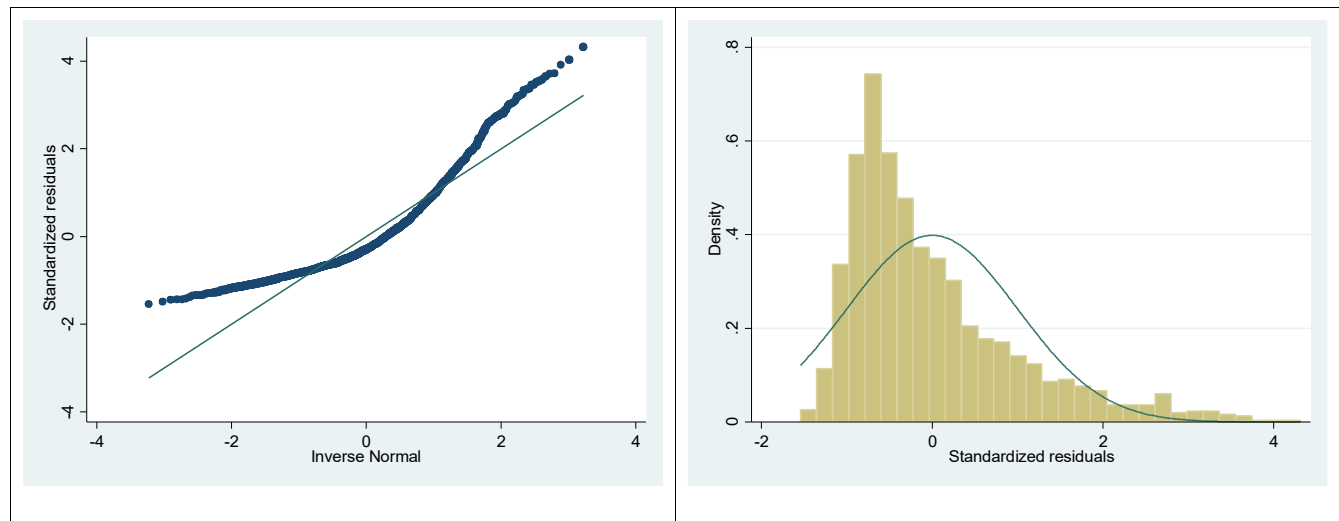
. imtest
```

Cameron & Trivedi's decomposition of IM-test

Source	chi2	df	p
Heteroskedasticity	74.11	44	0.0030
Skewness	143.84	9	0.0000
Kurtosis	33.27	1	0.0000
Total	251.22	54	0.0000



● Normality



```
. swilk stdres

Shapiro-Wilk W test for normal data

Variable | Obs   W      V      z      Prob>z
-----+-----
stdres   | 1574  0.87871 115.660 11.977  0.00000
```

Transformation of Y (L1a)

- Box-cox
- Interpretation

Re-assessing assumptions

- Log-transformed model

```
xi:boxcox wpc hs100_ctr hs100_ctr_sq parity1 i.aut_calv i.twin i.dyst  
i.rp*i.vag_disch
```

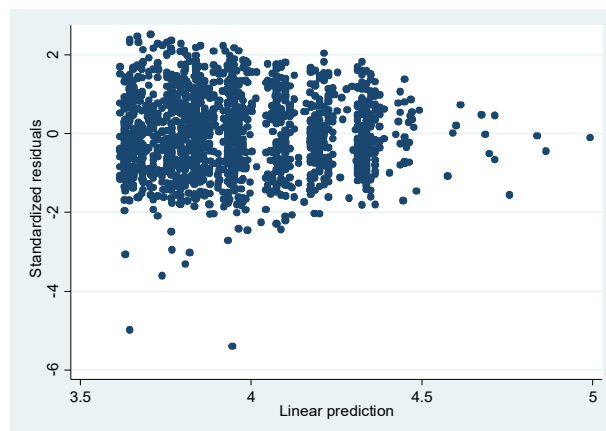
...output omitted...

```
Log likelihood = -7960.5368      Number of obs   =      1574  
                                LR chi2(8)           =      148.83  
                                Prob > chi2           =       0.000
```

	wpc	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
/theta		.1099104	.0271003	4.06	0.000	.0567947 .1630261

★ value of λ close to 0 $\rightarrow$ log transformation

- Homoscedasticity (L1a: p11)

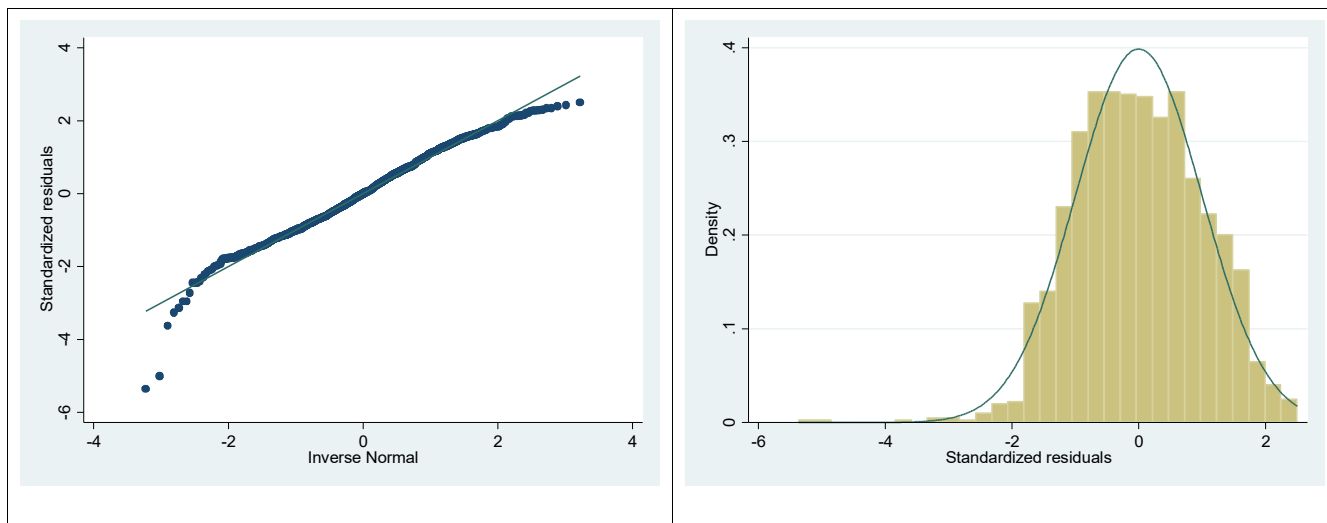


★ tests provide contradicting results

➡ imtest - constant variance

➡ however BPCW test highly significant

● Normality (L1a)



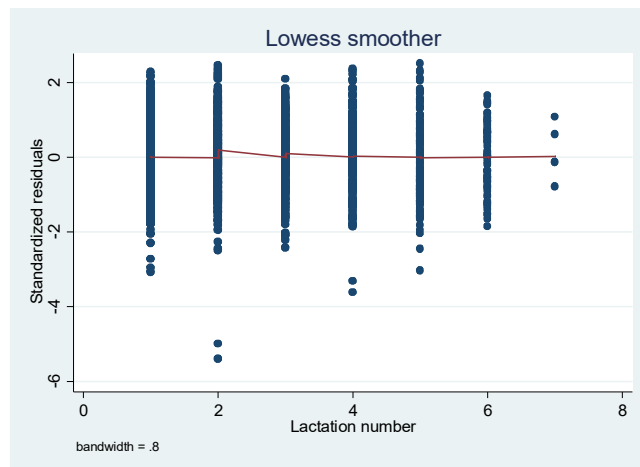
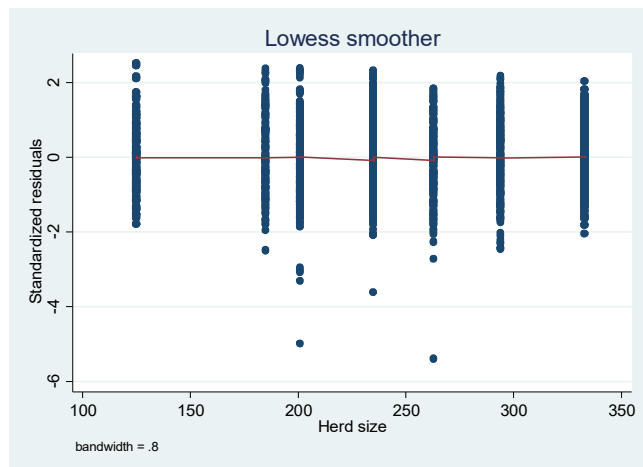
● Linearity (L1a)

★ continuous predictors only

★ standardized residuals vs predictor

Herd size

Parity



★ also pattern can be evaluated by looking at the residuals from a model without the predictor of interest

Evaluating individual observations

Outliers (lecture L1a)

- Large residuals
- Standardized residuals
 - ★ expect 5% > 2 and < -2
 - ★ expect 0.3% > 3 and < -3
- Deletion residuals
 - ★ t-test - $\text{prob} = 2 * n * t(\text{DFE}-1, di)$
- WPC log-transformed model - residuals
 - ★ standardized residuals
 - ➔ expected 5% = 79 (observed = 50)
 - ➔ expected 0.3% = 5 (observed = 6)
 - ➔ expected and observed values very similar
 - ★ deletion residuals
 - ➔ outlier cutoff value = 4.17
 - ➔ two observations $\geq$ this cutoff point
 - indication that these are outliers
 - cows with $\text{wpc} = 1$
 - ➔ delete and refit the model (later)
 - only for diagnostics purposes

Leverage (h)

- X-outliers
 - ★ depend on X values only
 - ➔ not affected by relationship with y
 - ★ expresses whether x_i is outlying in distribution of x's
 - so potentially influential
 - ★ less meaningful for categorical predictors
 - ➔ depends on the distribution of each category
- Cut-off values
 - ★ concerns if $h_i \geq 2(k+1)/n$
 - ➔ or $\geq 3(k+1)/n$
 - ➔ $k = \#$ of predictors, $n = \#$ obs.
 - ➔ should also look for unusual observations independently of these cutpoints
- Stata - predict varname, leverage
 - ★ eg. predict lev_lnwpc, lev
 - ★ lists/graphs to assess values

● Computed after full model (ln_wpc)

```
. summ lev
```

Variable	Obs	Mean	Std. Dev.	Min	Max
lev	1574	.0063532	.0083435	.0016636	.0642237

```
. display "leverage cutoff: " 2*nparam/nobs
```

leverage cutoff: .01270648

```
. display "conservative leverage cutoff: " 3*nparam/nobs
```

conservative leverage cutoff: .01905972

```
. count if lev>=.01905972 /* many (n=108) high leverage values */
```

```
. sort lev
```

```
. list wpc wpc_ln fit aut_calv herd_size parity1 twin dyst rp vag_disch stdres lev  
in -10/-1
```

	wpc	wpc_ln	fit	aut_calv	herd_size	parity1	twin	dyst	rp	vag_disch	stdres	lev
	40	3.689	4.407	yes	263	0	no	yes	yes	yes	-1.005	0.046
	23	3.135	4.121	yes	185	0	yes	yes	no	no	-1.381	0.050
	99	4.595	4.592	no	294	1	yes	yes	no	no	0.004	0.051
	32	3.466	4.262	yes	201	1	yes	yes	yes	no	-1.117	0.052
	110	4.700	4.162	yes	263	0	yes	no	no	yes	0.758	0.058
	53	3.970	4.227	yes	263	5	yes	no	no	yes	-0.362	0.059
	94	4.543	4.865	no	263	3	yes	no	yes	yes	-0.454	0.063
	137	4.920	4.994	no	294	2	yes	no	yes	yes	-0.105	0.064
	45	3.807	4.577	yes	201	4	yes	no	yes	yes	-1.089	0.064
	76	4.331	4.699	no	185	4	yes	no	yes	yes	-0.520	0.064

```
. table twin rp dyst
```

		Dystocia at calving and Retained placenta at calving			
		no		yes	
Twins		no	yes	no	yes
born	no	1332	124	76	15
	yes	15	9	2	1

★ large leverage values for less common combination of categorical predictors

Influence diagnostics: Cook's distance and DFITS

- Cook's distance and DFITS

$$\text{Cooks D} \quad D_i = \frac{r_{si}^2}{(k+1)} * \frac{h_i}{(1-h_i)}$$

$$\text{DFITS} \quad DFITS_i = r_{ii} \sqrt{\frac{h_i}{(1-h_i)}}$$

- ★ measure influence of an observation and have two interpretations:

- effect of deleting observation "i" on the predictions
- effect of outlier information about x's and y's

- ★ D_i is on the squared residual scale and based on standardized residuals

- ★ $DFITS_i$ is on absolute (signed) residual scale and are based on deletion residuals

- cut-off values

- ★ concern if either is >1 (rare) or if...

- $D_i \geq 1$ or $\geq 4/n$

- $DFITS_i$ outside $\pm 2 * \sqrt{\frac{k+1}{n}}$ (if $n \geq 120$)

- $n < 120$ only values beyond ± 1

- Stata: `predict varname, cooksdfits`

- ★ lists/graphs to assess values

● Model: ln_wpc

★ Cook's

```
. summ cook
```

Variable	Obs	Mean	Std. Dev.	Min	Max
cook	1574	.0006351	.0013679	5.57e-10	.0174304

```
. display "Cook's D cutoff: " 4/nobs  
Cook's D cutoff: .0025413
```

```
. count if cook>=.0025413 /* many (n=95) high Cook's D values */  
95
```

★ DFITS

```
. summ dfit
```

Variable	Obs	Mean	Std. Dev.	Min	Max
dfit	1574	.0003931	.0797594	-.4179343	.3664328

```
. display "DFITS cutoff: " 2*sqrt(nparam/nobs)*(nobs>=120)+1*(nobs<120)  
DFITS cutoff: .15941443
```

```
. count if abs(dfit)>=.15941443 /* many (n=95) high DFITS values */  
95
```

★ largest Cook's values

➡ not clear pattern – cows with at least one disease

★ largest negative DFIT values

➡ cows with twins and at least one of the diseases were the most influential

★ largest positive DFIT values

➡ cows without twins but with at least one of the diseases were the most influential

Delta-beta (or DFBETA)

★ influence of an observation on a specific regression coefficient (β)

★ for obs. "i" and predictor x_j

$$\Rightarrow \text{DFBETA}_{j(i)} = \frac{(\beta_j - \beta_{j(i)})}{SE_j}$$

→ where $\beta_{j(i)}$ is the estimate of β_j without obs. "i"

★ measures change in β_j in terms of SE's

→ how many SE's does β_j increase(decrease) if "i" is dropped?

● cut-off values

★ $\text{DFBETA}_{j(i)} > 0$ (< 0) ~ obs. "i" pulls β_j up (down)

→ extreme values of $\text{DFBETA}_{j(i)}$ are notable

→ most meaningful for continuous predictors

★ concern if outside $\pm \frac{2}{\sqrt{n}}$ (for $n \geq 120$)

→ $n < 120$ only values beyond ± 1

● Stata

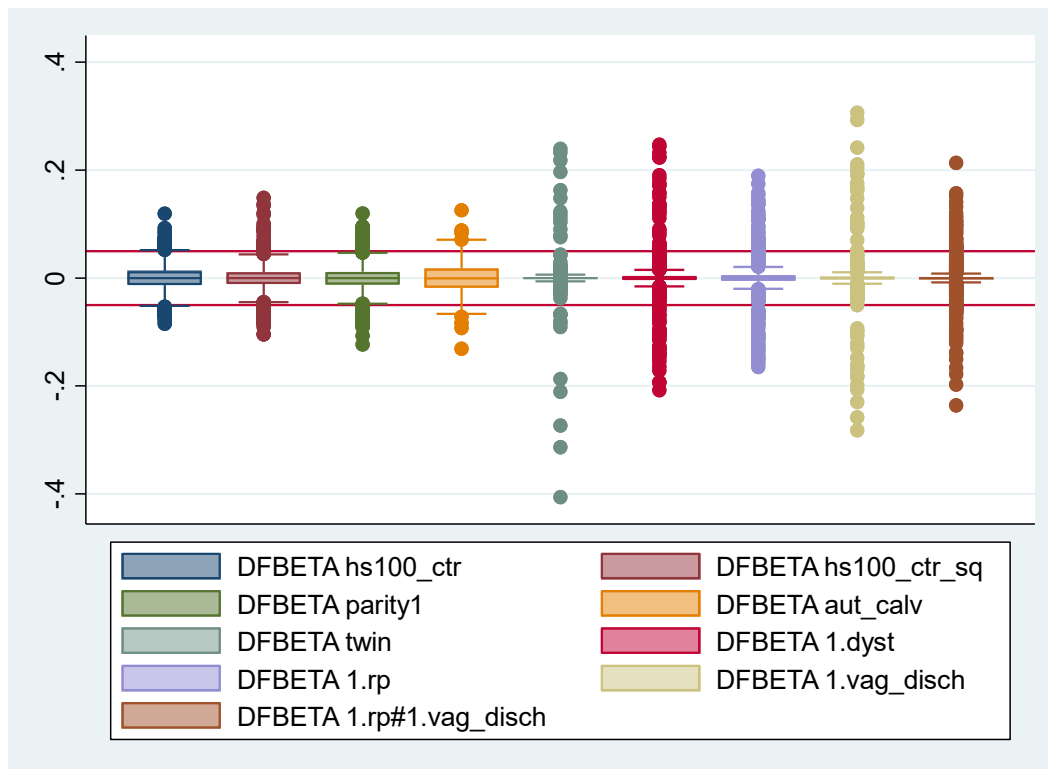
★ `predict varname, dfbeta(var in model)`

→ eg. `predict dfbdyst, dfbeta(dyst)`

★ `list`/`graphs` to assess values

● Model: ln_wpc

```
. display "DFBETA cutoff: " 2/sqrt(nobs)*(nobs>=120)+1*(nobs<120)
DFBETA cutoff: .05041127
```



● Explore values by browsing/listing

★ large values are cases of the correspondent predictor (eg. cows with diseases and twins)

★ linear effect of herd size similar to quadratic term

➡ need to be aware of collinearity

What to do with Outliers/Influential observations

- Verify that point(s) (observation(s)) are not errors (eg. data entry errors)
- May be very informative
 - ★ always examine observations and values of the predictors
- Determine if outliers are significant (L1a)
- Fit models with and without the outlier/influential observation(s)
- Keeping them in the analysis
 - ★ potential for biased estimates
 - ★ usually leads to larger SE and reduced power (is therefore conservative)
 - ★ always report them
- Eliminating them from the analysis
 - ★ should be always reported and justified
 - ★ narrows scope of inference from the study
 - ★ may create an unrealistically good model

Other transformations

- See “l2b_linear_reg_diag_other_transf.do”
- log-transformed model
 - ★ some indication of non-constant variance
 - ★ residuals still not normally-distributed

➡ two severe outliers

```
. l wpc wpc_ln fit herd_size parity dyst rp vag_disch if wpc==1, clean compress noobs
```

wpc	wpc~ln	fit	herd_size	parity	dyst	rp	vag~h
1	0	3.645548	201	2	no	no	no
1	0	3.945861	263	2	no	no	no

```
. l wpc wpc_ln delres lev dfit cook dfb_dyst dfb_rp dfb_vd dfb_int dfb_hs if wpc==1, clean compress noobs
```

wpc	wpc~ln	delres	lev	dfit	cook	df~st	dfb~p	dfb~d	dfb~nt	dfb_hs
1	0	-5.028	0.002	-0.243	0.006	0.051	0.037	0.028	-0.021	0.098
1	0	-5.449	0.002	-0.243	0.006	0.044	0.053	0.042	-0.026	-0.016

★ comparison without outliers

```
. estimate table ln ln_noout
```

Variable	ln	ln_noout
hs100_ctr	.34365799	.33916323
hs100_ctr_sq	.21187276	.20269649
parity1	.01298436	.0113333
aut_calv	-.13716214	-.13695147
twin	.39274261	.3894441
dyst	.1109405	.10338895
rp		
yes	.11231661	.10603408
vag_disch		
yes	-.02601346	-.03328151
rp#vag_disch		
yes#yes	.41369992	.42230092
_cons	3.8885867	3.9010244

- ★ repeat but without influential obs.
- ★ advantage log-transformed model
 - ➡ interpretation of estimates still possible

- Square root transformation

- ★ example VER – constant variance -> ok
- ★ residuals still not normally-distributed
- ★ deletion residuals show no indication of outlying observations
- ★ interpretation estimates not possible
 - ➔ obtain predicted values
 - ➔ parity=3 aut_calv=1 hsize=0 hsize2=0 twin=0

rp#vag_disch#dyst	Original scale			Sqrt transformation			log-transformation		
	est	lo_ci	up_ci	est	lo_ci	up_ci	est	lo_ci	up_ci
no#no#no	66.605	62.220	70.990	58.267	54.583	62.071	50.127	47.005	53.456
no#no#yes	78.305	67.338	89.272	66.840	57.187	77.244	56.008	47.688	65.780
no#yes#no	67.833	53.767	81.899	58.064	46.695	70.671	48.840	39.737	60.027
no#yes#yes	79.533	63.311	95.755	66.622	52.636	82.255	54.570	43.017	69.225
yes#no#no	72.592	63.048	82.135	64.364	56.085	73.213	56.085	48.761	64.510
yes#no#yes	84.292	70.918	97.666	73.359	61.107	86.731	62.666	51.506	76.243
yes#yes#no	96.672	78.396	114.947	90.259	71.883	110.723	82.645	63.217	108.045
yes#yes#yes	108.372	87.603	129.141	100.857	78.877	125.534	92.342	68.098	125.216

*ci estimated from t-distribution (Stata uses a different estimation procedure)

- ★ sqrt model – better than log-transformed
 - ➔ no outliers (eg. deletion res. range: -2.42; 3.28)
- ★ influential diagnostics
 - ➔ similar to log-transformed model
 - ➔ more in VER.