

SOCY7708: Hierarchical Linear Modeling
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Class notes: HLM Diagnostics

Like OLS, HLM models rely on certain assumptions that have to be satisfied in order for regression coefficients to be unbiased and efficient estimates of the parameters of interest. Therefore, it is important to watch out for possible assumption violations and to take steps to prevent them. We will address the issues of model specification, collinearity, homoscedasticity, normality of level 1 and level 2 residuals, and linearity.

1. Model specification.

In HLM models, the issue of model specification concerns two main questions: (1) Did we include the right fixed effects? (2) Did we include the right random components? As we discussed, when specifying your model, you should rely heavily on your theory as well as utilize hypothesis testing. But there are some additional steps you can take to prevent model misspecification:

- Consider including aggregates of level 1 variables. It is always possible that what appears to be an effect of a level 1 variable is, in reality, an effect of its level 2 aggregate. The only way to test is to introduce such an aggregate. So far, we discussed aggregates to the mean, but sometimes, it is also possible to use group-level standard deviations. For example, you can use MEANSES to indicate the average level of SES in the school and DEVSES (within-school standard deviation) to indicate how diverse each school is in terms of SES. Such diversity may have an impact above and beyond the impact of the average level.

```
. bysort id: egen devses=sd(ses)

. mixed mathach c.ses##c.meanses c.ses##i.sector i.female##c.meanses
i.female##i.sector devses || id: female, cov(unstr)

Mixed-effects ML regression                Number of obs    =      7,185
Group variable: id                        Number of groups =      160

                                           Obs per group:
                                           min =           14
                                           avg =           44.9
                                           max =            67

                                           Wald chi2(9)     =      861.54
                                           Prob > chi2      =      0.0000

Log likelihood = -23217.242
```

mathach	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
ses	2.852182	.147225	19.37	0.000	2.563626 3.140738
meanses	3.123813	.5007382	6.24	0.000	2.142384 4.105242
c.ses#					
c.meanses	.7762981	.2691847	2.88	0.004	.2487057 1.30389
ses	0	(omitted)			
1.sector	1.00463	.4108992	2.44	0.014	.1992828 1.809978
sector#c.ses					

1		-1.54621	.2220969	-6.96	0.000	-1.981512	-1.110908
1.female		-1.218688	.2383675	-5.11	0.000	-1.68588	-.7514966
meanses		0	(omitted)				
female#							
c.meanses							
1		-.0401053	.5063098	-0.08	0.937	-1.032454	.9522437
female#							
sector							
1 1		.0546438	.4182702	0.13	0.896	-.7651507	.8744384
devses		-2.666202	1.457351	-1.83	0.067	-5.522557	.1901533
_cons		14.57225	1.041436	13.99	0.000	12.53107	16.61343

Random-effects Parameters	Estimate	Std. Err.	[95% Conf. Interval]	
id: Unstructured				
var(female)	.9479572	.5671689	.2934333	3.062444
var(_cons)	2.934085	.5636358	2.013527	4.27551
cov(female,_cons)	-1.170931	.4832245	-2.118034	-.2238285
var(Residual)	36.3806	.6189185	35.18755	37.61411

LR test vs. linear model: $\chi^2(3) = 183.93$ Prob > $\chi^2 = 0.0000$

Note: LR test is conservative and provided only for reference.

- Consider including level 2 predictors of level 1 slopes (i.e., cross-level interactions) if you find significant variation in these slopes
- If the proportion of explained variance (R-squared – we discussed earlier how to calculate R-squared within each level and overall) is substantially reduced when you add a fixed effect, that can be a sign of misspecification.
- Sometimes a fixed effect misspecification (e.g., a nonlinearity) can lead to a misspecification of the random effects (excluded curvilinear effect may show up as a significant variance component for the slope). We will return to the issue of linearity below.

To prevent the misspecification problems in terms of random components:

- Always test whether each of your level 1 slopes varies across level 2 units (i.e., try to estimate each slope as random). However, you have to be careful not to “overtax” your data – if you have very few lower level cases within each upper level unit, you can’t do too many random components.
- If the model doesn’t converge or if it takes a long time to converge, that may mean that the model has too many random effects and the data are relatively sparse. In general, you should be cautious in specifying level-1 coefficients as random – as the number of random effects grows, the number of variances/covariances to be estimated increases even faster (for m random predictors, there are $1+m(m+1)/2$ variance covariance components). As the number of random effects grows, significantly more information is required to obtain reasonable estimates of variance/covariance components. The maximum depends on a number of factors: the magnitude of the variance components,

the degree of intercorrelation among the random effects, the magnitude of sigma squared, and other characteristics of the data.

- If there are high correlations among level-1 coefficients (i.e., slopes for different variables—correlations with the intercept are ok), the model must be simplified. There are a number of ways of dealing with it. You can, for example, use factor analysis to form scales and reduce the number of variables. You can also constrain one or more random effects to be zero (i.e. keep only the fixed effect for that variable), thus eliminating the correlation. This works well if that random effect is in fact negligible in size and/or non-significant.

2. Multicollinearity

Like regular OLS, HLM models can be affected by multicollinearity. There are no tools to check for it specifically for mixed command, but you can check basic correlations among your independent variables as well as variance inflation factors (VIFs) for the same model estimated with OLS:

```
. pwcorr mathach ses female meanses sector size
```

	mathach	ses	female	meanses	sector	size
mathach	1.0000					
ses	0.3608	1.0000				
female	-0.1231	-0.0679	1.0000			
meanses	0.3437	0.5306	-0.0589	1.0000		
sector	0.2040	0.1896	0.0065	0.3573	1.0000	
size	-0.0506	-0.0673	-0.0388	-0.1268	-0.4237	1.0000

```
. reg mathach ses female meanses sector size
```

Source	SS	df	MS	Number of obs =	7185
Model	61205.6611	5	12241.1322	F(5, 7179) =	315.35
Residual	278671.273	7179	38.8175614	Prob > F =	0.0000
				R-squared =	0.1801
				Adj R-squared =	0.1795
Total	339876.934	7184	47.3102637	Root MSE =	6.2304

mathach	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]
ses	2.148034	.1113801	19.29	0.000	1.929697 2.366372
female	-1.321295	.1478042	-8.94	0.000	-1.611034 -1.031555
meanses	2.889622	.2206451	13.10	0.000	2.457093 3.322151
sector	1.503238	.1724585	8.72	0.000	1.165169 1.841308
size	.0003457	.0001345	2.57	0.010	.0000821 .0006093
_cons	12.32108	.2222729	55.43	0.000	11.88536 12.7568

```
. vif
```

Variable	VIF	1/VIF
meanses	1.54	0.648947
ses	1.39	0.717093
sector	1.38	0.726733
size	1.22	0.818586
female	1.01	0.992362
Mean VIF	1.31	

Different researchers advocate for different cutoff points for VIF. Some say that if any one of VIF values is larger than 4, there are some multicollinearity problems associated with that variable. Others use cutoffs of 5 or even 10. It can also be useful to check level 2 separately using means of your dependent variable as an outcome:

```
. bysort id: egen mathachm=mean(mathach)
```

```
. reg mathachm meanses sector size if tagged==1
```

Source	SS	df	MS	Number of obs	=	160
Model	1016.16465	3	338.72155	F(3, 156)	=	99.84
Residual	529.275622	156	3.39279245	Prob > F	=	0.0000
				R-squared	=	0.6575
				Adj R-squared	=	0.6509
Total	1545.44027	159	9.71975014	Root MSE	=	1.842

mathachm	Coefficient	Std. err.	t	P> t	[95% conf. interval]
meanses	5.359654	.3777613	14.19	0.000	4.613467 6.105841
sector	1.524517	.349298	4.36	0.000	.8345531 2.214481
size	.0005152	.0002603	1.98	0.050	1.00e-06 .0010293
_cons	11.38922	.4055502	28.08	0.000	10.58814 12.1903

```
. vif
```

Variable	VIF	1/VIF
sector	1.42	0.706227
size	1.26	0.794707
meanses	1.15	0.872531
Mean VIF	1.27	

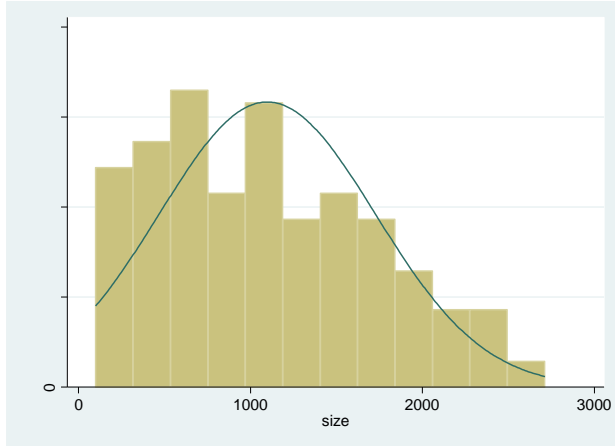
When running your mixed models, you can also watch out for potential signs of multicollinearity (e.g., large coefficients for two correlated variables going in opposite directions, high standard errors).

3. Normality

HLM models assume that the level-1 and level 2 error terms are normally distributed. To make sure this assumption will be met, it is important to do some preliminary data screening before running mixed models. It is especially important to ensure that your dependent variable distribution is as close to normal as possible, but you should check independent variables as well. If substantial deviations from normality are identified, consider fixing them with a transformation. Note that when examining normality of level 2 variables, you should either have a separate level 2 file or you should limit your analysis to one record per higher level unit.

```
. egen tagged=tag(id)
```

```
. histogram size if tagged==1  
(bin=12, start=100, width=217.75)
```

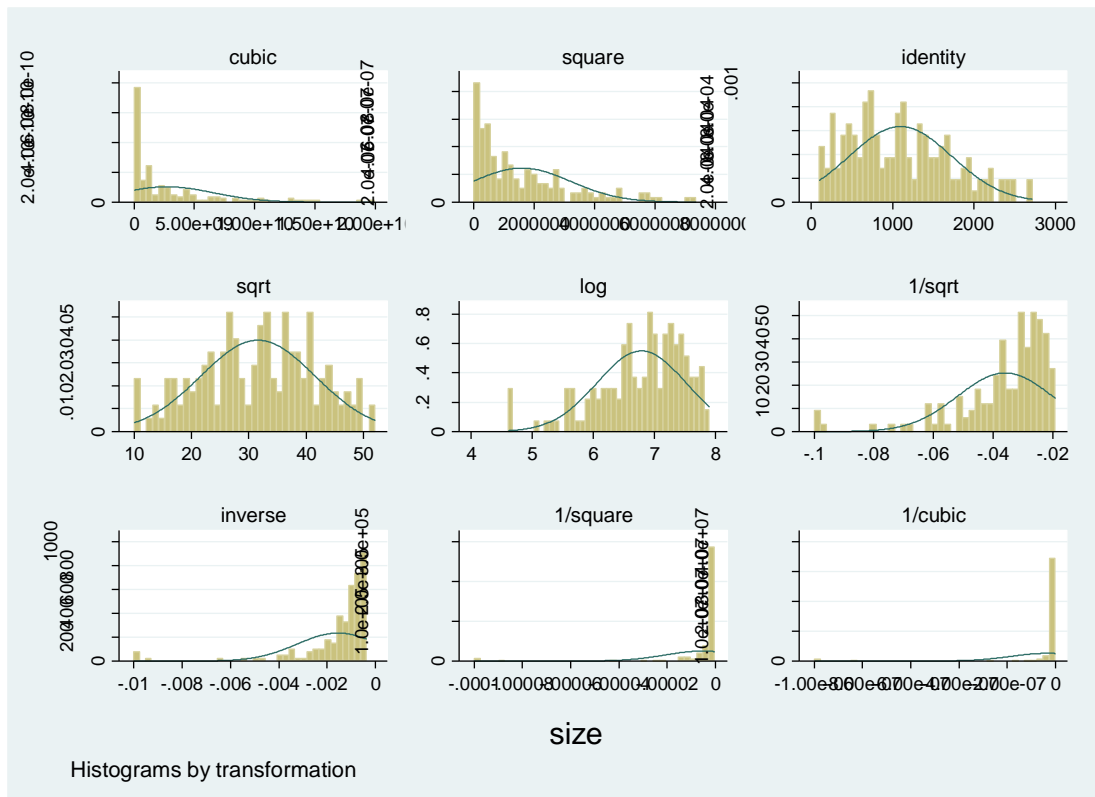


Looks like a right skew; to find a transformation:

```
. ladder size if tagged==1
```

Transformation	formula	chi2 (2)	P (chi2)
cubic	size^3	60.02	0.000
square	size^2	31.36	0.000
identity	size	8.37	0.015
square root	sqrt(size)	7.18	0.028
log	log(size)	16.55	0.000
1/(square root)	1/sqrt(size)	58.10	0.000
inverse	1/size	.	0.000
1/square	1/(size^2)	.	0.000
1/cubic	1/(size^3)	.	0.000

```
. gladder size if tagged==1
```



Square root looks the best, so we would generate it and then later on import that transformed variable into HLM:

```
. gen sizesqrt=sqrt(size)
```

If a variable contains zero or negative values, you need to add a constant to it before looking for transformations (such that all values of the variable become larger than zero). For example:

```
. ladder mathach
```

Transformation	Formula	chi2(2)	Prob > chi2
Cubic	mathach^3	758.08	0.000
Square	mathach^2	758.90	0.000
Identity	mathach	914.19	0.000
Square root	sqrt(mathach)	.	.
Log	log(mathach)	.	.
1/(Square root)	1/sqrt(mathach)	.	.
Inverse	1/mathach	.	.
1/Square	1/(mathach^2)	.	.
1/Cubic	1/(mathach^3)	.	.

```
. gladder mathach
```

```
. sum mathach
```

Variable	Obs	Mean	Std. dev.	Min	Max
mathach	7,185	12.74785	6.878246	-2.832	24.993

```
. gen mathach_c=mathach-r(min)+1
```

```
. sum mathach_c
```

Variable	Obs	Mean	Std. dev.	Min	Max
mathach_c	7,185	16.57985	6.878246	1	28.825

```
. ladder mathach_c
```

Transformation	Formula	chi2(2)	Prob > chi2
Cubic	mathac~c^3	595.26	0.000
Square	mathac~c^2	1029.42	0.000
Identity	mathac~c	914.19	0.000
Square root	sqrt(mathac~c)	380.40	0.000
Log	log(mathac~c)	.	.
1/(Square root)	1/sqrt(mathac~c)	.	.
Inverse	1/mathac~c	.	.
1/Square	1/(mathac~c^2)	.	.
1/Cubic	1/(mathac~c^3)	.	.

```
. gladder mathach_c
```

If your sample size is large, everything will be significantly different from normal, so you should either rely on graphical examination (gladder) or randomly select a subsample of your dataset and do this type of analysis for that subsample.

If a variable is negatively skewed, you might have an easier time finding a transformation for it after reversing it. To reverse the variable and yet keep all the values positive, you can subtract it from its maximum value + 1; for example:

```
. sum mathach
```

Variable	Obs	Mean	Std. Dev.	Min	Max
mathach	7185	12.74785	6.878246	-2.832	24.993

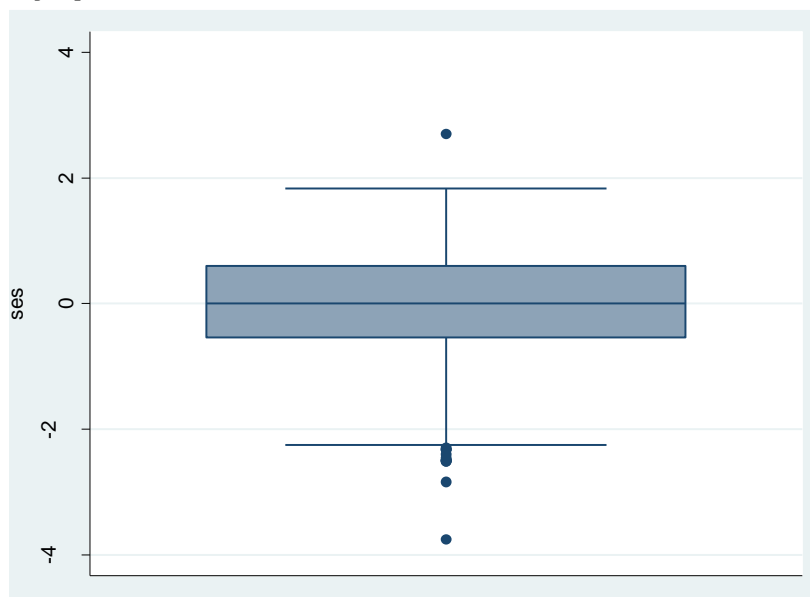
```
. gen mathachr=r(max)+1-mathach
```

```
. sum mathachr
```

Variable	Obs	Mean	Std. Dev.	Min	Max
mathachr	7185	13.24515	6.878246	.9999999	28.825

As you are examining normality, pay attention to outliers as well – sometimes, it is useful to top-code or bottom-code outliers in addition to or instead of transforming a variable.

```
. graph box ses
```



```
. sum ses, detail
```

Percentiles		Smallest		
1%	-1.848	-3.758		
5%	-1.318	-2.838		
10%	-1.038	-2.508	Obs	7185
25%	-.538	-2.508	Sum of Wgt.	7185
50%	.002		Mean	.0001434
		Largest	Std. Dev.	.7793552
75%	.602	1.732		
90%	1.022	1.762	Variance	.6073945
95%	1.212	1.832	Skewness	-.2281447
99%	1.512	2.692	Kurtosis	2.620279

```
. gen ses1=clip(ses, -2.9, 1.9)
```

Never top-code or bottom-code more than 5% of the distribution; better yet, do 1% or less. Sometimes transformation might be a better way to bring in outliers so consider both options or a combination of them.

If you do a good job dealing with normality problems and with outliers during preliminary screening, you should not run into problems with multivariate normality. Still, we need to check both level 1 and level 2 residuals for normality. Let's estimate a model, obtain residuals, and inspect them.

```
. mixed mathach c.ses#c.meanses c.ses##i.sector i.female#c.meanses
i.female##i.sector || id: female, cov(unstr)
```

```
Mixed-effects ML regression      Number of obs      =      7,185
Group variable: id              Number of groups   =      160

                                Obs per group:
                                min =      14
                                avg =      44.9
                                max =      67

                                Wald chi2(8)      =      849.53
                                Prob > chi2       =      0.0000

Log likelihood = -23218.854
```

mathach	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
ses	2.856887	.1473012	19.39	0.000	2.568182 3.145592
meanses	3.203604	.4968255	6.45	0.000	2.229844 4.177364
c.ses#					
c.meanses	.832274	.268213	3.10	0.002	.3065862 1.357962
ses	0	(omitted)			
1.sector	1.167355	.3994507	2.92	0.003	.3844458 1.950264
sector#c.ses					
1	-1.554133	.2223377	-6.99	0.000	-1.989907 -1.118359
1.female	-1.22104	.238342	-5.12	0.000	-1.688182 -.7538981
meanses	0	(omitted)			
female#					
c.meanses					
1	-.0074533	.5063542	-0.01	0.988	-.9998894 .9849827
female#					
sector					
1 1	.047171	.4186158	0.11	0.910	-.773301 .8676429
_cons	12.71993	.2439351	52.14	0.000	12.24183 13.19804

Random-effects Parameters	Estimate	Std. Err.	[95% Conf. Interval]
id: Unstructured			
var(female)	.9446425	.5677046	.290883 3.067726
var(_cons)	2.897432	.55434	1.991409 4.215665
cov(female,_cons)	-1.091921	.4762149	-2.025285 -.1585568


```
-----+-----
var(Residual) | 36.37821 .6188368 35.18531 37.61155
-----+-----
LR test vs. linear model: chi2(3) = 190.77          Prob > chi2 = 0.0000
```

Note: LR test is conservative and provided only for reference.

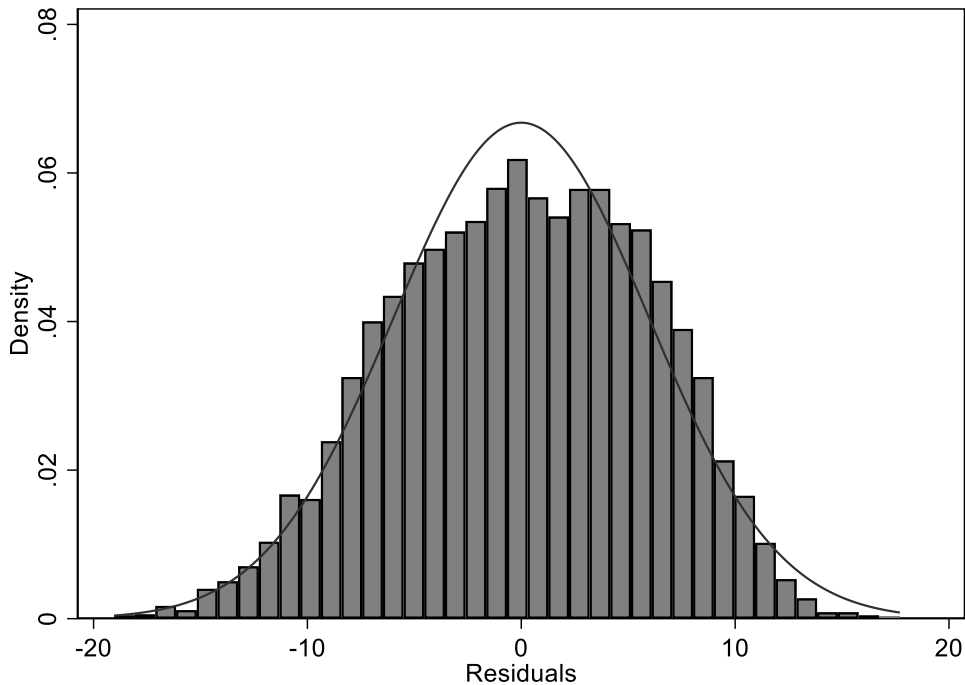
To check the distribution of error terms, we obtain level-1 residuals and level-2 residuals. Here is what we can obtain using predict command:

```
xb          xb, linear predictor for the fixed portion of the model
stdp       standard error of the fixed-portion linear prediction xb
fitted     fitted values, linear predictor of the fixed portion plus
           contributions based on predicted random effects
residuals  residuals, response minus fitted values
rstandard  standardized residuals
reffects   best linear unbiased predictions (BLUPs) of the
           random effects. By default, BLUPs for all random effects in the
           model are calculated. You must specify q new variables, where q is
           the number of random-effects terms in the model.
reses      standard errors of the best linear unbiased predictions (BLUPs) of
           the random effects. By default, standard errors for all BLUPs in
           the model are calculated. You must specify q new variables.
```

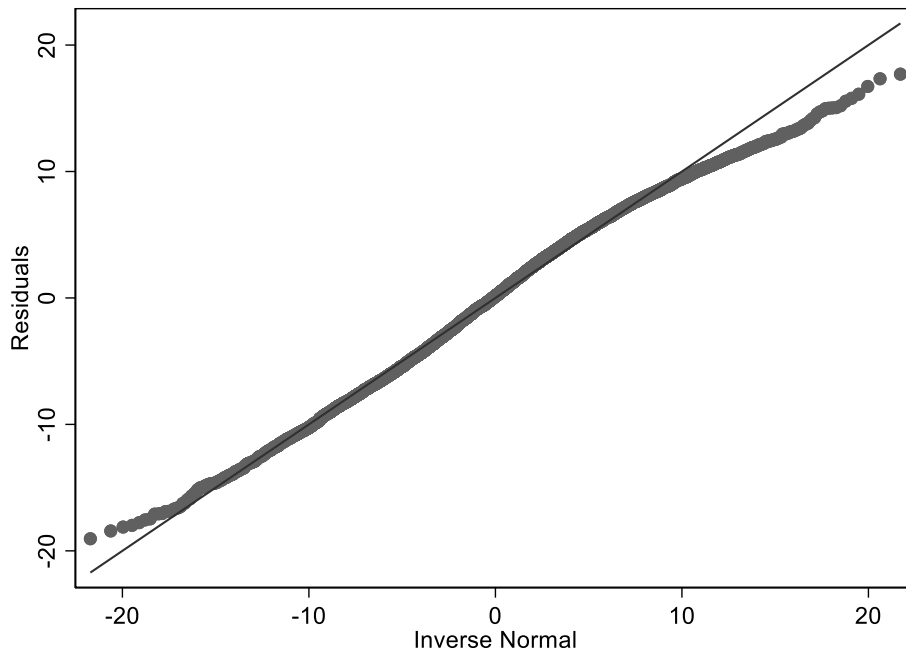
Thus, residuals will give you level 1 residuals, and reffects will give you level 2 residuals for each level 2 random component. You should examine both types of residuals to assess normality.

Level-1 residuals:

```
. predict llresid, resid
. histogram llresid, normal
(bin=38, start=-19.084782, width=.96868813)
```



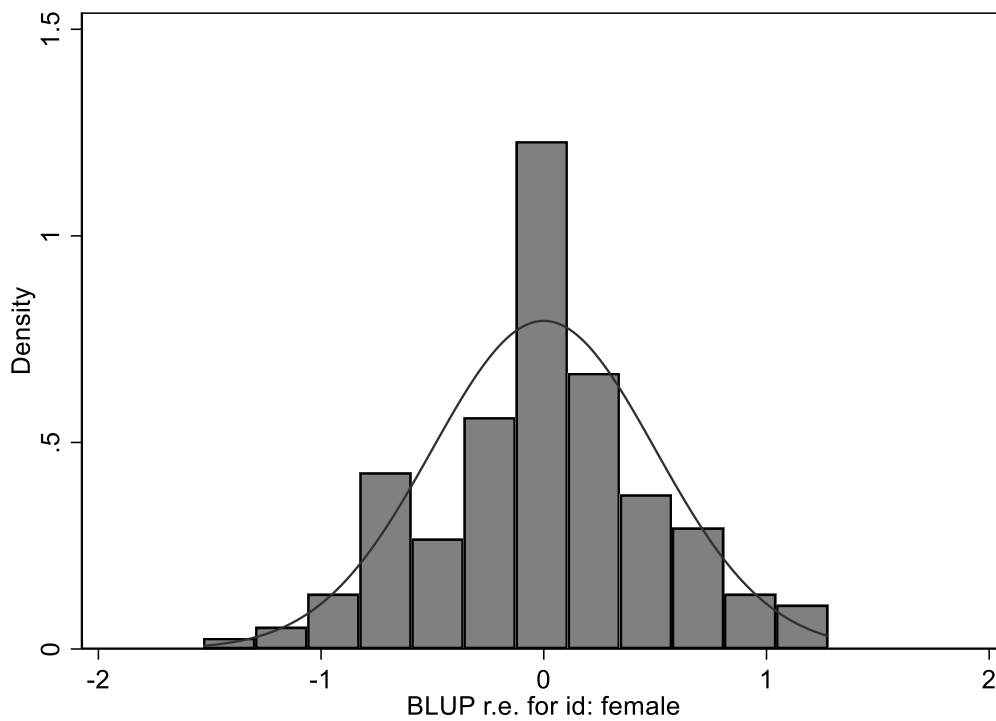
```
. qnorm llresid
```



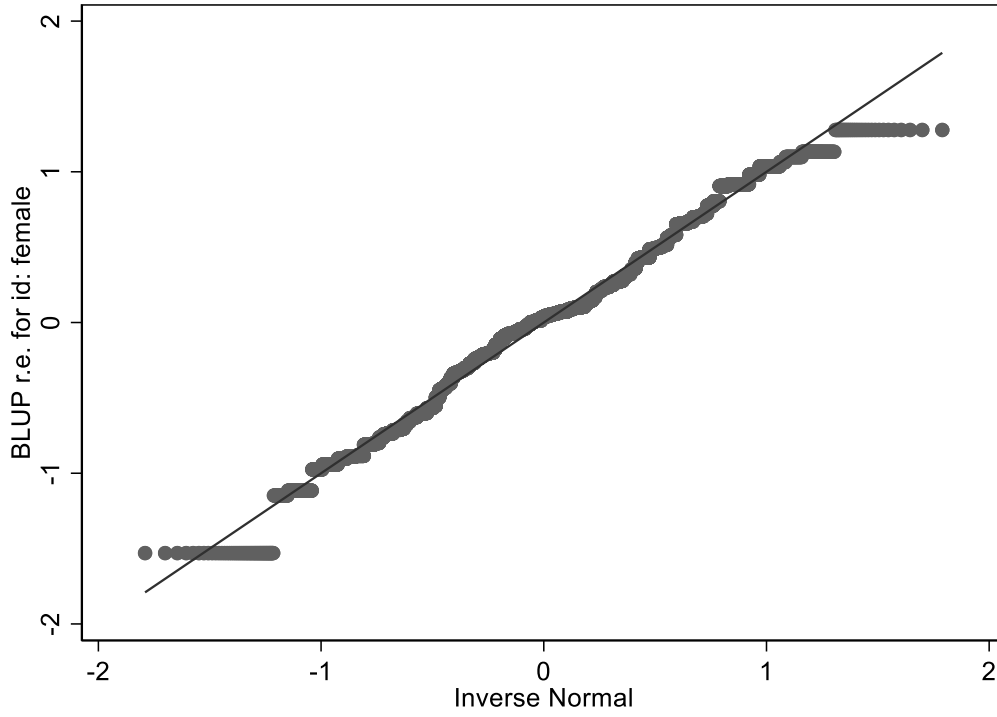
Level 2 residuals:

```
. predict l2resid*, reffects
```

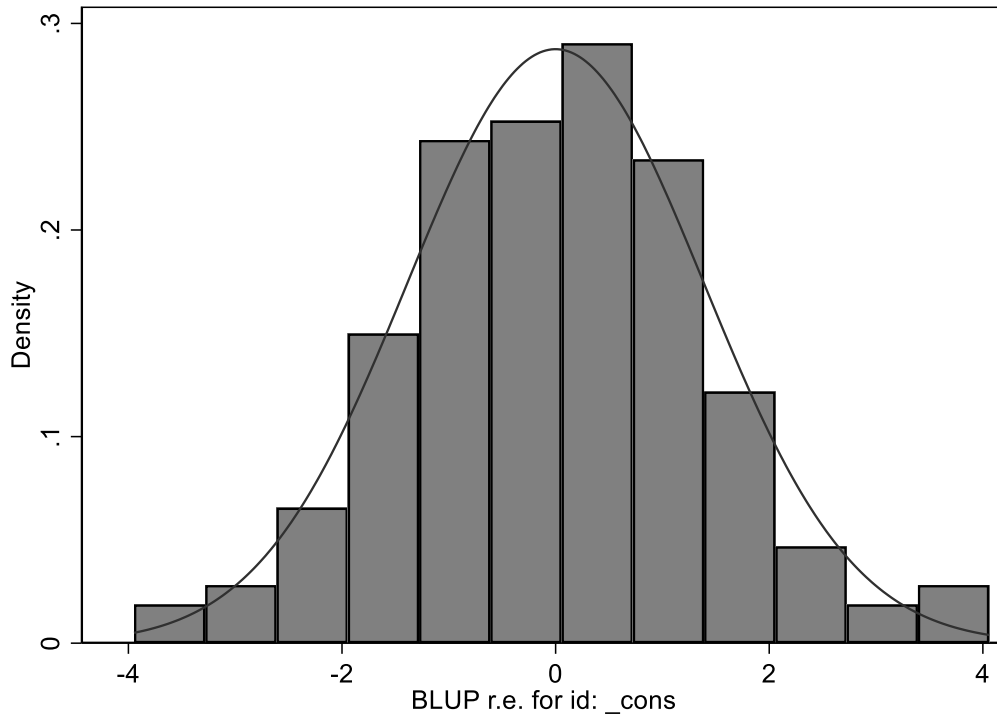
```
. histogram l2resid1 if tagged==1, normal  
(bin=12, start=-4.0914528, width=.6652911)
```



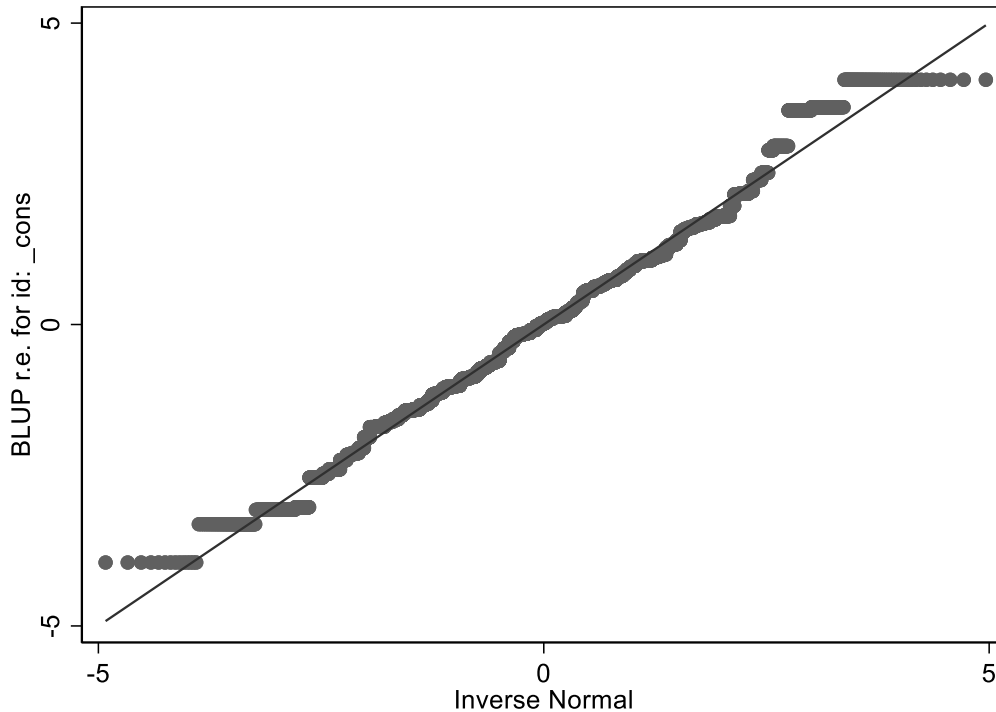
```
. qnorm l2resid1
```



```
. histogram l2resid2 if tagged==1, normal
(bin=12, start=-1.3546074, width=.2191729)
```



```
. qnorm l2resid2
```



If there are problems with normality of residuals but you can't fix them with simple transformations or top/bottomcoding, you can use robust option available with mixed for robust standard errors to obtain standard errors and significance tests that are less dependent on assumptions:

```
. mixed mathach c.ses##c.meanses c.ses##i.sector i.female##c.meanses i.female#
> #i.sector || id: female, cov(unstr) robust
note: ses omitted because of collinearity
note: meanses omitted because of collinearity
```

```
Mixed-effects regression      Number of obs   =      7,185
Group variable: id           Number of groups =      160

                               Obs per group:
                               min =      14
                               avg =     44.9
                               max =      67

                               Wald chi2(8)      =    1204.55
                               Prob > chi2       =      0.0000

Log pseudolikelihood = -23218.854
```

(Std. Err. adjusted for 160 clusters in id)

mathach	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ses	2.856887	.1408367	20.29	0.000	2.580852	3.132922
meanses	3.203604	.4716531	6.79	0.000	2.27918	4.128027
c.ses#						
c.meanses	.832274	.2933011	2.84	0.005	.2574144	1.407134
ses	0	(omitted)				

1.sector		1.167355	.3975957	2.94	0.003	.3880816	1.946628
sector#c.ses							
1		-1.554133	.2228935	-6.97	0.000	-1.990996	-1.11727
1.female		-1.22104	.2216856	-5.51	0.000	-1.655536	-.7865441
meanses		0	(omitted)				
female#							
c.meanses							
1		-.0074533	.4225481	-0.02	0.986	-.8356324	.8207257
female#							
sector							
1 1		.047171	.4142162	0.11	0.909	-.7646779	.8590199
_cons		12.71993	.2224148	57.19	0.000	12.28401	13.15586

Random-effects Parameters	Estimate	Robust Std. Err.	[95% Conf. Interval]	
id: Unstructured				
var(female)	.9446425	.5153373	.3242692	2.751879
var(_cons)	2.897432	.5871549	1.947691	4.310289
cov(female,_cons)	-1.091921	.452792	-1.979377	-.2044647
var(Residual)	36.37821	.7087571	35.01526	37.79421

You can also use bootstrapping, although it does take time to calculate:

```
. bootstrap, cluster(id): mixed mathach c.ses##c.meanses c.ses##i.sector
i.female##c.meanses i.female##i.sector || id: female, cov(unstr)
(running mixed on estimation sample)
```

Bootstrap replications (50)

```
----- 1 ----- 2 ----- 3 ----- 4 ----- 5
..... 50
```

```
Mixed-effects ML regression      Number of obs   =      7,185
Group variable: id              Number of groups =      160
```

```
Obs per group:
      min =      14
      avg =     44.9
      max =      67
```

```
Log likelihood = -23218.854      Wald chi2(8)    =     1548.18
                                Prob > chi2       =      0.0000
```

(Replications based on 160 clusters in id)

mathach	Observed Coef.	Bootstrap Std. Err.	z	P> z	Normal-based [95% Conf. Interval]	
ses	2.856887	.1349074	21.18	0.000	2.592474	3.121301
meanses	3.203604	.3618389	8.85	0.000	2.494412	3.912795
c.ses#						
c.meanses	.832274	.3020349	2.76	0.006	.2402965	1.424252
ses	0	(omitted)				

1.sector		1.167355	.3569849	3.27	0.001	.4676774	1.867032
sector#c.ses							
1		-1.554133	.2129321	-7.30	0.000	-1.971472	-1.136794
1.female		-1.22104	.1914944	-6.38	0.000	-1.596362	-.8457178
meanses		0	(omitted)				
female#							
c.meanses							
1		-.0074533	.3536247	-0.02	0.983	-.700545	.6856383
female#							
sector							
1 1		.047171	.3585003	0.13	0.895	-.6554766	.7498186
_cons		12.71993	.1796728	70.79	0.000	12.36778	13.07209

Random-effects Parameters		Observed Estimate	Bootstrap Std. Err.	Normal-based [95% Conf. Interval]	
id: Unstructured					
var(female)		.9446425	.2268802	.5899694	1.512535
var(_cons)		2.897432	.4663447	2.113545	3.972053
cov(female,_cons)		-1.091921	.2330362	-1.548663	-.6351782
var(Residual)		36.37821	.6943167	35.04251	37.76482

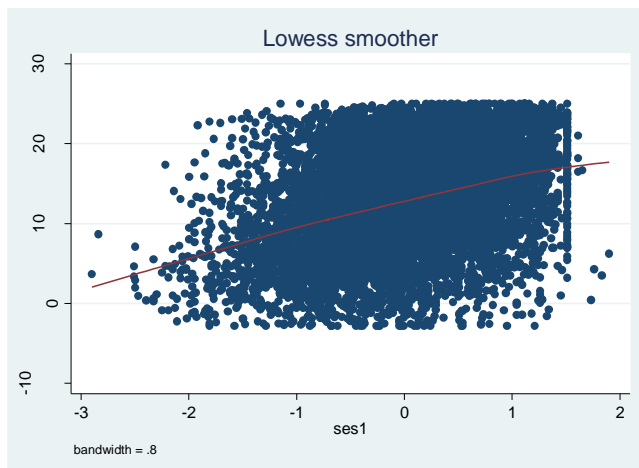
LR test vs. linear model: $\chi^2(3) = 190.77$ Prob > $\chi^2 = 0.0000$

Note: LR test is conservative and provided only for reference.

4. Linearity

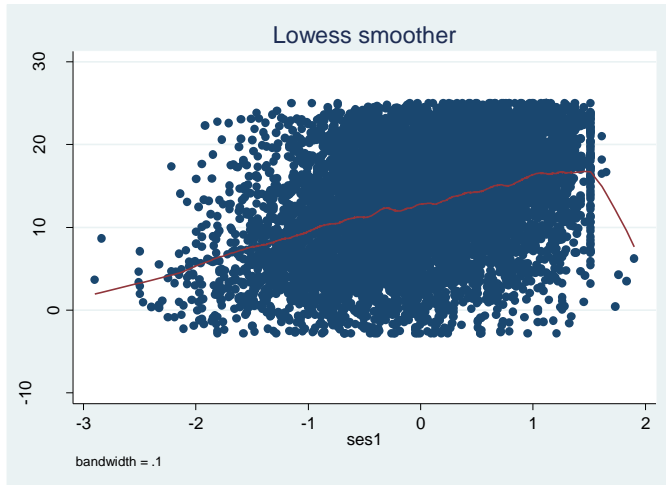
Before running mixed models, it's also a good idea to examine the relationship of each independent variable to the dependent to assess its linearity. A good tool for such an examination is a lowess plot – that is, a scatterplot with locally weighted regression line (based on means or medians) going through it:

```
. lowess mathach ses1
```



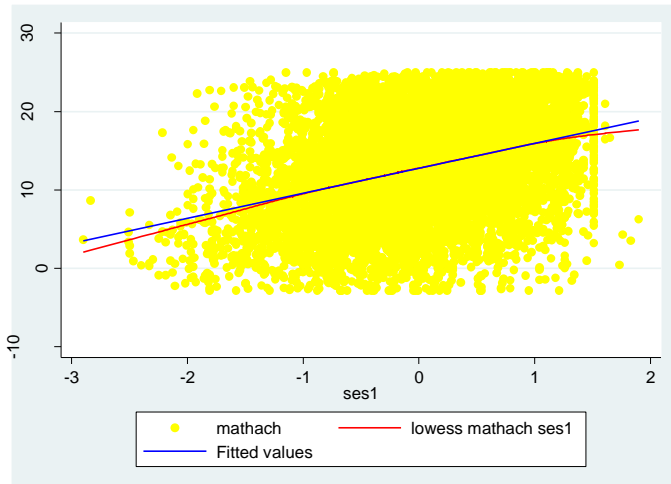
We can change bandwidth to make the curve less smooth (decrease the number) or smoother (increase the number):

```
. lowess mathach ses1, bwidth(.1)
```



We can also add a regression line to see the difference better:

```
. scatter mathach ses1, mcolor(yellow) || lowess mathach ses1, lcolor(red) ||
lfit mathach ses1, lcolor(blue)
```

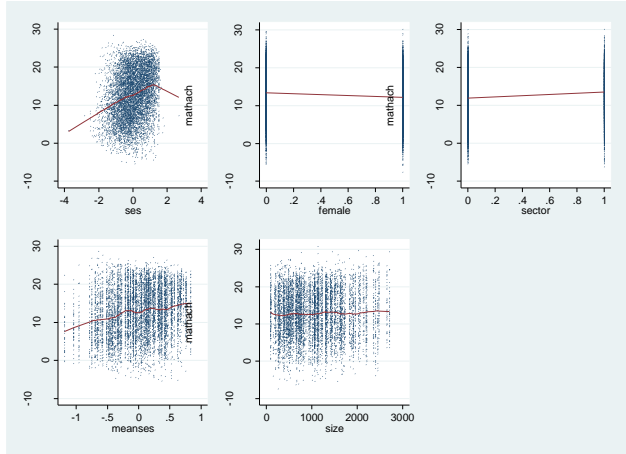


You can do an approximate test for multivariate linearity (based on OLS) with a user-written mrunning program:

```
. search mrunning
Keyword search
  Keywords:  mrunning
  Search:   (1) Official help files, FAQs, Examples, SJs, and STBs
Search of official help files, FAQs, Examples, SJs, and STBs
SJ-5-3  gr0017  . . . . . A multivariable scatterplot smoother
(help mrunning, running if installed) . . . . P. Royston and N. J. Cox
Q3/05   SJ 5(3):405--412
presents an extension to running for use in a
multivariable context
```

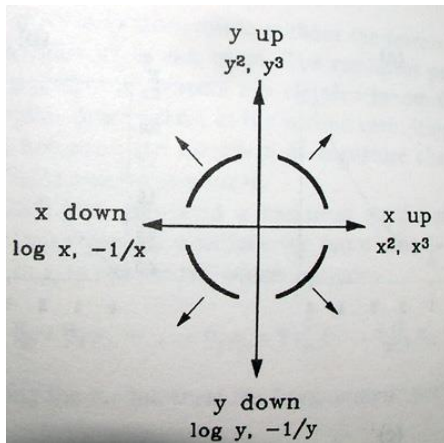
Click on gr0017 to install the program. Now we can use it:

```
. mrunning mathach ses female sector meanses size
```



If the relationship looks nonlinear on lowess plot, consider using transformations to fix it before importing data into HLM. (Note that if the relationship is too complex, sometimes we may choose to break up the corresponding independent variable into a series of dummies instead.)

Monotone nonlinear relationship: Power transformations can be used to linearize relationships if strong monotone nonlinearities are found. The following chart gives suggestions for transformations when the curve looks a certain way:



Nonmonotone relationship: For non-monotone relationships (e.g. parabola or cubic), use polynomial functions of the variable, e.g. ses and ses squared, etc. Note that when including variables that are generated using other variables already in the model (as in this case, or when we want to enter a product of two variables to model an interaction term), we should mean-center the variable outside of HLM (only if it is continuous; don't mean-center dichotomous variables!), and then square and/or cube the mean-centered variable. We will then include the mean-centered variable itself and its transformations into our HLM file and our models. For example, if we are dealing with a second level variable, we would get its mean across 160 level 2 cases by restricting the calculation to one case per level 2 unit:

```
. sum size if tagged==1
  Variable |      Obs      Mean      Std. Dev.      Min      Max
-----+-----
```



```

size |      160    1097.825    629.5064      100      2713
. gen size=size-r(mean)
. gen size2=size^2

```

Oftentimes, the same transformation that helps with normality also will improve linearity, but that it is not always the case. Overall, linearity is more important to enforce than normality for a given variable, so if you end up with incompatible transformations, opt for the one improving linearity.

Once we estimated our HLM model and obtained residuals, we can inspect them to further assess linearity. First, we can assess the overall pattern by plotting level 1 residuals against predicted values; there should be no discernable pattern:

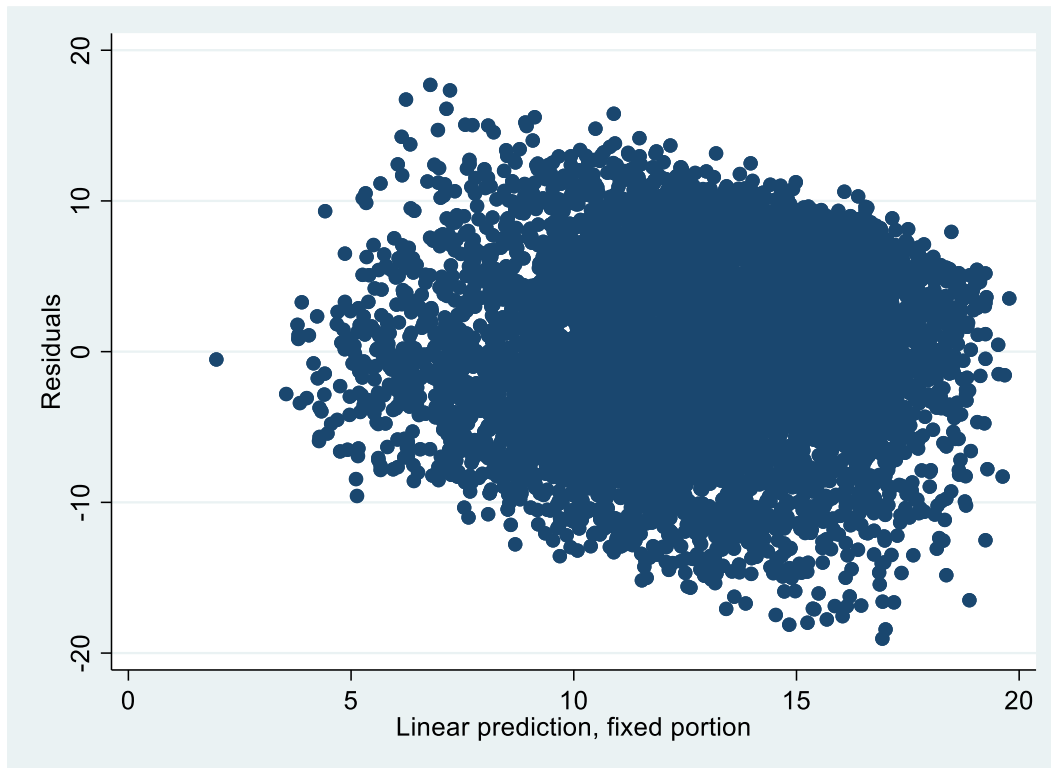
```

. qui mixed mathach c.ses##c.meanses c.ses##i.sector i.female##c.meanses
i.female##i.sector || id: female, cov(unstr)

. predict xb, xb

. scatter llresid xb

```



This does not look too good; indicates potential heteroscedasticity or nonlinearity problems.

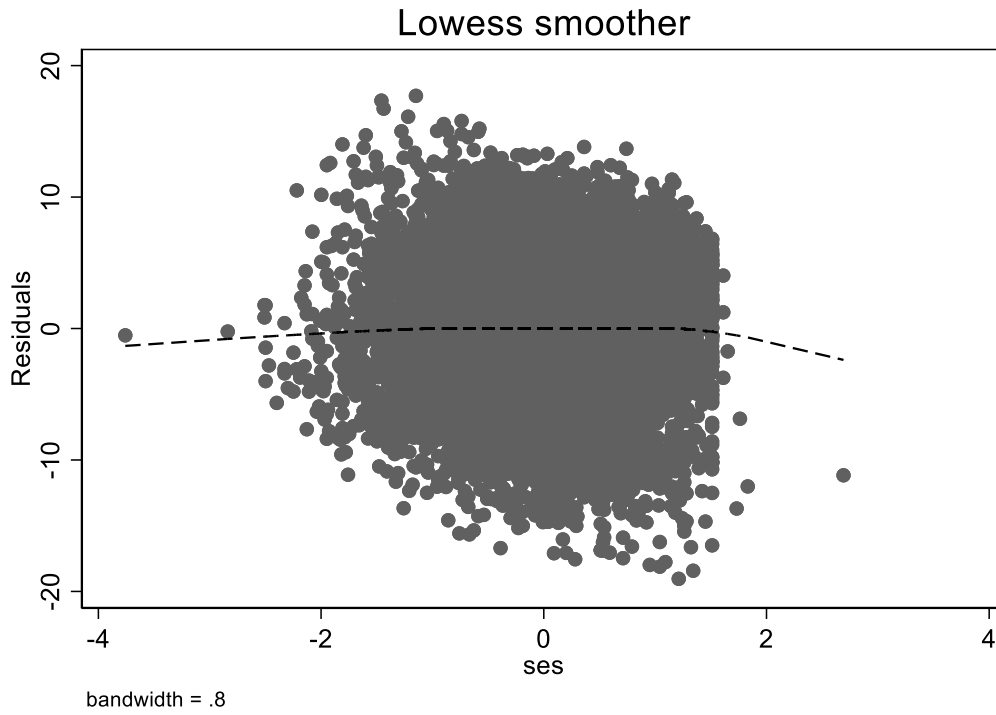
To test the linearity assumption for continuous predictors, it is useful to plot residuals against each of the continuous dependent variable. To improve our ability to detect a curvilinear relationship, we will include a smoother in our plot using lowess command.

In level 1 file:

```

. lowess llresid ses

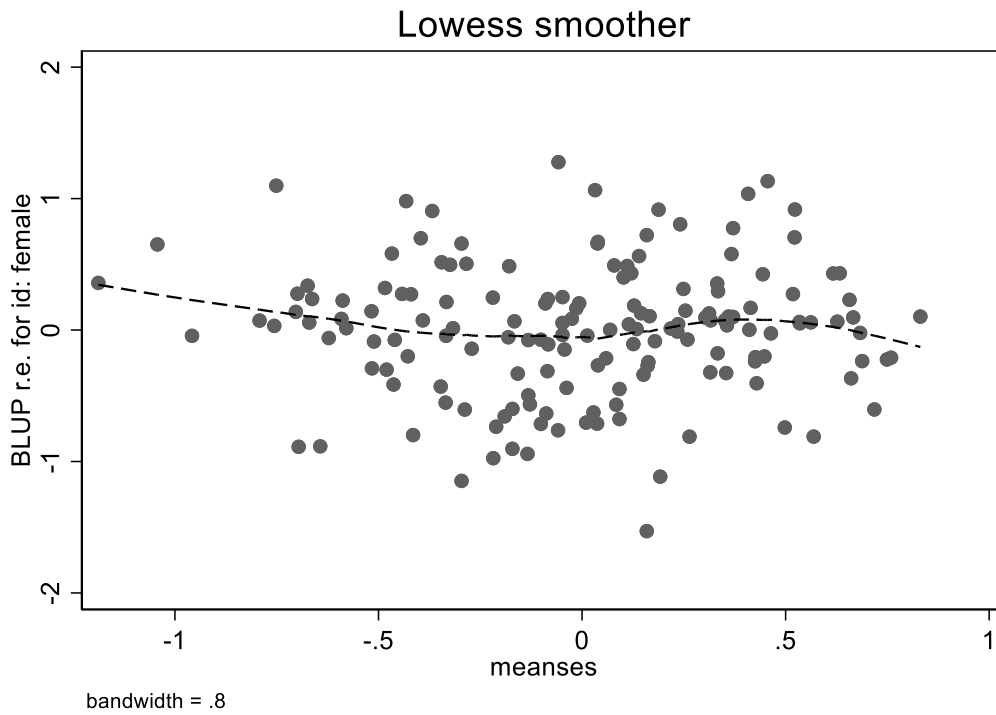
```



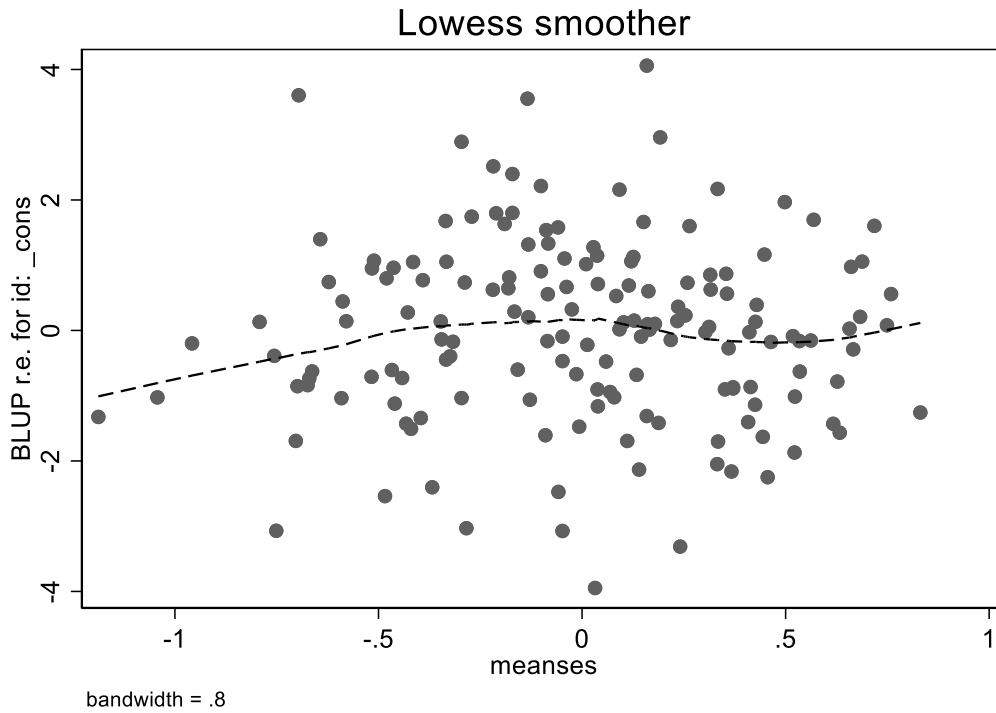
Looks more or less fine, but we do see some outliers on SES.

In level 2 file:

```
. lowess l2resid1 meansas if tagged==1
```



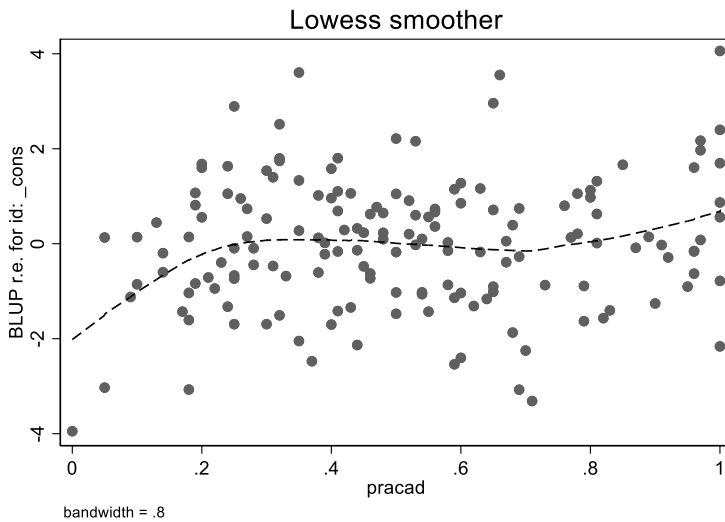
```
. lowess l2resid2 meansas if tagged==1
```



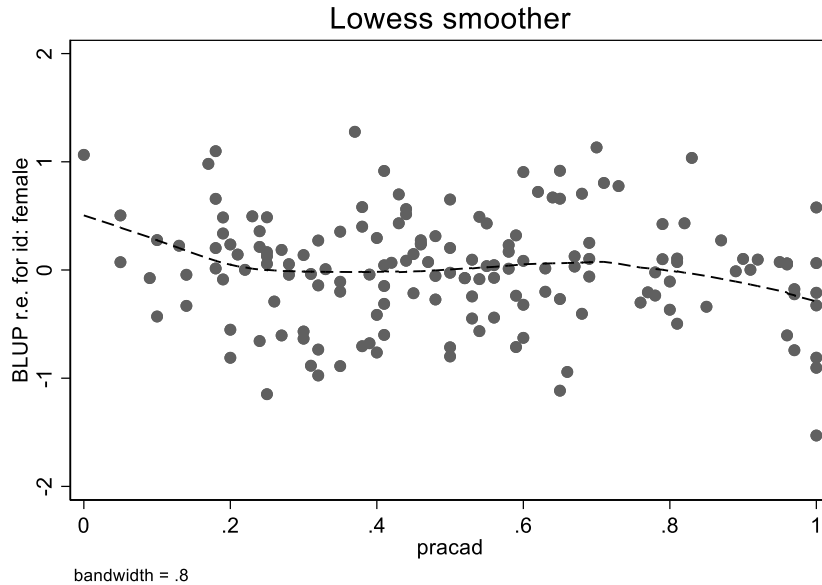
Based on these graphs, we could consider modeling nonlinear relationships with MEANSES (e.g. cubic).

We can also use such plots to search for potential other relationships and examine their shape, e.g. with PRACAD:

```
. lowess l2resid1 pracad if tagged==1
```



```
. lowess l2resid2 pracad if tagged==1
```



Based on graphs for MEANSES, looks like we need a quadratic and cubic term; let's try that. We run the following model:

Level-1 Model

$$MATHACH_{ij} = \beta_{0j} + \beta_{1j}*(FEMALE_{ij}) + \beta_{2j}*(SES_{ij}) + r_{ij}$$

Level-2 Model

$$\beta_{0j} = \gamma_{00} + \gamma_{01}*(SECTOR_j) + \gamma_{02}*(MEANSESM_j) + \gamma_{03}*(MEANSES2_j) + \gamma_{04}*(MEANSES3_j) + u_{0j}$$

$$\beta_{1j} = \gamma_{10} + \gamma_{11}*(SECTOR_j) + \gamma_{12}*(MEANSESM_j) + \gamma_{13}*(MEANSES2_j) + \gamma_{14}*(MEANSES3_j) + u_{1j}$$

$$\beta_{2j} = \gamma_{20} + \gamma_{21}*(SECTOR_j) + \gamma_{22}*(MEANSESM_j) + \gamma_{23}*(MEANSES2_j) + \gamma_{24}*(MEANSES3_j) + u_{2j}$$

Mixed Model

$$\begin{aligned}
 MATHACH_{ij} = & \gamma_{00} + \gamma_{01}*SECTOR_j + \gamma_{02}*MEANSESM_j + \gamma_{03}*MEANSES2_j \\
 & + \gamma_{04}*MEANSES3_j \\
 & + \gamma_{10}*FEMALE_{ij} + \gamma_{11}*SECTOR_j*FEMALE_{ij} + \gamma_{12}*MEANSESM_j*FEMALE_{ij} + \gamma_{13}*MEANSES2_j*FEMALE_{ij} \\
 & + \gamma_{14}*MEANSES3_j*FEMALE_{ij} \\
 & + \gamma_{20}*SES_{ij} + \gamma_{21}*SECTOR_j*SES_{ij} + \gamma_{22}*MEANSESM_j*SES_{ij} + \gamma_{23}*MEANSES2_j*SES_{ij} \\
 & + \gamma_{24}*MEANSES3_j*SES_{ij} \\
 & + u_{0j} + u_{1j}*FEMALE_{ij} + u_{2j}*SES_{ij} + r_{ij}
 \end{aligned}$$

```
. mixed mathach c.ses##c.meanses##c.meanses##c.meanses c.ses##i.sector
i.female##c.meanses##c.meanses##c.meanses i.female##i.sector || id: female, cov(unstr)
```

```
Mixed-effects ML regression
Group variable: id
```

```
Number of obs    =    7,185
Number of groups =    160
```

```
Obs per group:
    min =    14
    avg =   44.9
```

max = 67

Log likelihood = -23211.143 Wald chi2(14) = 876.38
 Prob > chi2 = 0.0000

mathach	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
ses	3.050629	.1656633	18.41	0.000	2.725935	3.375323
meanses	1.817196	.8053935	2.26	0.024	.2386537	3.395738
c.ses#						
c.meanses	.4610032	.4533027	1.02	0.309	-.4274538	1.34946
c.meanses#						
c.meanses	-.9934494	1.086095	-0.91	0.360	-3.122156	1.135258
c.ses#						
c.meanses#						
c.meanses	-1.44887	.6373161	-2.27	0.023	-2.697987	-.1997537
c.meanses#						
c.meanses#						
c.meanses	3.866993	1.659089	2.33	0.020	.6152377	7.118749
c.ses#						
c.meanses#						
c.meanses#						
c.meanses	.8161978	.9309449	0.88	0.381	-1.008421	2.640816
ses	0	(omitted)				
1.sector	1.283584	.3963461	3.24	0.001	.5067598	2.060408
sector#c.ses						
1	-1.4467	.2256309	-6.41	0.000	-1.888929	-1.004472
1.female	-1.230308	.2740098	-4.49	0.000	-1.767358	-.693259
meanses	0	(omitted)				
female#						
c.meanses	.7925874	.8138469	0.97	0.330	-.8025233	2.387698
1						
female#						
c.meanses#						
c.meanses#						
c.meanses	-.1274134	1.087919	-0.12	0.907	-2.259695	2.004869
1						
female#						
c.meanses#						
c.meanses#						
c.meanses	-1.805483	1.583418	-1.14	0.254	-4.908926	1.297959
1						
female#						
sector						
1 1	-.0036908	.4203907	-0.01	0.993	-.8276415	.8202598
_cons	12.8906	.2731983	47.18	0.000	12.35514	13.42606

Random-effects Parameters	Estimate	Std. Err.	[95% Conf. Interval]	
id: Unstructured				

```

      var(female) |      .91266   .5645054   .2715297   3.067614
      var(_cons) |      2.727112   .530815   1.862182   3.993778
cov(female,_cons) |     -1.011546   .4651488   -1.923221   -.0998711
-----+-----
      var(Residual) |      36.32734   .6179661   35.13612   37.55895
-----+-----
LR test vs. linear model: chi2(3) = 180.35          Prob > chi2 = 0.0000

```

Note: LR test is conservative and provided only for reference.

It looks like there is a bunch of non-significant coefficients that we could omit; let's omit and compare using BIC (we could also do joint hypothesis test for these coefficients first):

```
. estat ic
```

Akaike's information criterion and Bayesian information criterion

```

-----+-----
      Model |          N   ll(null)   ll(model)     df       AIC       BIC
-----+-----
      . |          7,185           .   -23211.14     19   46460.29   46591
-----+-----

```

Note: BIC uses N = number of observations. See [R] BIC note.

```
. mixed mathach c.ses##c.meanses##c.meanses##c.meanses c.ses##i.sector i.female || id:
female, cov(unstr)
```

```

Mixed-effects ML regression          Number of obs   =       7,185
Group variable: id                   Number of groups =       160

Obs per group:
      min =          14
      avg =          44.9
      max =           67

```

```

Log likelihood = -23212.015          Wald chi2(10)   =       873.56
                                      Prob > chi2     =       0.0000

```

```

-----+-----
      mathach |      Coef.   Std. Err.     z   P>|z|   [95% Conf. Interval]
-----+-----
      ses |      3.053147   .1655297   18.44   0.000   2.728715   3.377579
      meanses |      2.324254   .6047559    3.84   0.000   1.138954   3.509553
      c.ses# |
      c.meanses |      .4447929   .4528177    0.98   0.326   -.4427134   1.332299
      c.meanses# |
      c.meanses |     -1.088537   .8269356   -1.32   0.188   -2.709301   .5322275
      c.ses# |
      c.meanses# |
      c.meanses |     -1.458181   .6368045   -2.29   0.022   -2.706295   -.2100674
      c.meanses# |
      c.meanses# |
      c.meanses |      2.697434   1.288303    2.09   0.036   .1724068   5.222461
      c.ses# |
      c.meanses# |
      c.meanses# |
      c.meanses |      .8568271   .9304757    0.92   0.357   -.9668717   2.680526
-----+-----

```

```

      ses |          0 (omitted)
1.sector |  1.285378  .2859008    4.50  0.000    .7250229    1.845734
sector#c.ses
  1 | -1.447254  .2254075   -6.42  0.000   -1.889045   -1.005464
  1.female | -1.213093  .1813475   -6.69  0.000   -1.568528   -.8576586
    _cons |  12.87913  .2388608   53.92  0.000   12.41097   13.34729
-----

```

```

Random-effects Parameters |   Estimate  Std. Err.   [95% Conf. Interval]
-----+-----
id: Unstructured
      var(female) |   .9800848   .5711366    .3127755    3.071104
      var(_cons) |   2.742527   .5344897    1.871805    4.018289
      cov(female,_cons) | -1.047902   .4702886   -1.969651   -.1261537
-----+-----
      var(Residual) |   36.3267   .6179416   35.13552   37.55826
-----

```

```
LR test vs. linear model: chi2(3) = 182.23          Prob > chi2 = 0.0000
```

Note: LR test is conservative and provided only for reference.

```
. estat ic
```

Akaike's information criterion and Bayesian information criterion

```

-----+-----
Model |          N  ll(null)  ll(model)    df        AIC        BIC
-----+-----
. |          7,185          . -23212.02    15  46454.03  46557.23
-----

```

Note: BIC uses N = number of observations. See [R] BIC note.

BIC strongly prefers the more parsimonious model. We could also consider getting rid of cubed term for MEANSES, but let's keep for now and examine graphically using margins and marginsplot:

```
. sum ses
```

```

Variable |          Obs          Mean    Std. Dev.    Min    Max
-----+-----
ses |          7,185    .0001434    .7793552   -3.758    2.692

```

```
. global sesmin=r(min)
```

```
. global sesmax=r(max)
```

```
. global sesmean=r(mean)
```

```
. global plusd=r(mean)+r(sd)
```

```
. global minusd=r(mean)-r(sd)
```

```
. sum meanses if tagged==1
```

```

Variable |          Obs          Mean    Std. Dev.    Min    Max
-----+-----
meanses |          160   -.0001875    .4139731   -1.188    .831

```

```
. global meansesmin=r(min)
```

```

. global meansesmax=r(max)

. global meansesmean=r(mean)

. global meansesplussd=r(mean)+r(sd)

. global meansesminusd=r(mean)-r(sd)

. margins, at(ses= ($sesmin $minusd $sesmean $plussd $sesmax) meanses=($means
> esmin $meansesminusd $meansesmean $meansesplussd $meansesmax))

```

```

Predictive margins                                Number of obs    =        7,185

```

```

Expression   : Linear prediction, fixed portion, predict()

```

1._at	: ses	=	-3.758
	meanses	=	-1.188
2._at	: ses	=	-3.758
	meanses	=	-.4141606
3._at	: ses	=	-3.758
	meanses	=	-.0001875
4._at	: ses	=	-3.758
	meanses	=	.4137856
5._at	: ses	=	-3.758
	meanses	=	.831
6._at	: ses	=	-.7792118
	meanses	=	-1.188
7._at	: ses	=	-.7792118
	meanses	=	-.4141606
8._at	: ses	=	-.7792118
	meanses	=	-.0001875
9._at	: ses	=	-.7792118
	meanses	=	.4137856
10._at	: ses	=	-.7792118
	meanses	=	.831
11._at	: ses	=	.0001434
	meanses	=	-1.188
12._at	: ses	=	.0001434
	meanses	=	-.4141606
13._at	: ses	=	.0001434
	meanses	=	-.0001875
14._at	: ses	=	.0001434
	meanses	=	.4137856
15._at	: ses	=	.0001434
	meanses	=	.831
16._at	: ses	=	.7794985
	meanses	=	-1.188


```

17._at      : ses          =    .7794985
              meanses     =   -0.4141606

18._at      : ses          =    .7794985
              meanses     =   -0.0001875

19._at      : ses          =    .7794985
              meanses     =    .4137856

20._at      : ses          =    .7794985
              meanses     =    .831

21._at      : ses          =    2.692
              meanses     =   -1.188

22._at      : ses          =    2.692
              meanses     =   -0.4141606

23._at      : ses          =    2.692
              meanses     =   -0.0001875

24._at      : ses          =    2.692
              meanses     =    .4137856

25._at      : ses          =    2.692
              meanses     =    .831

```

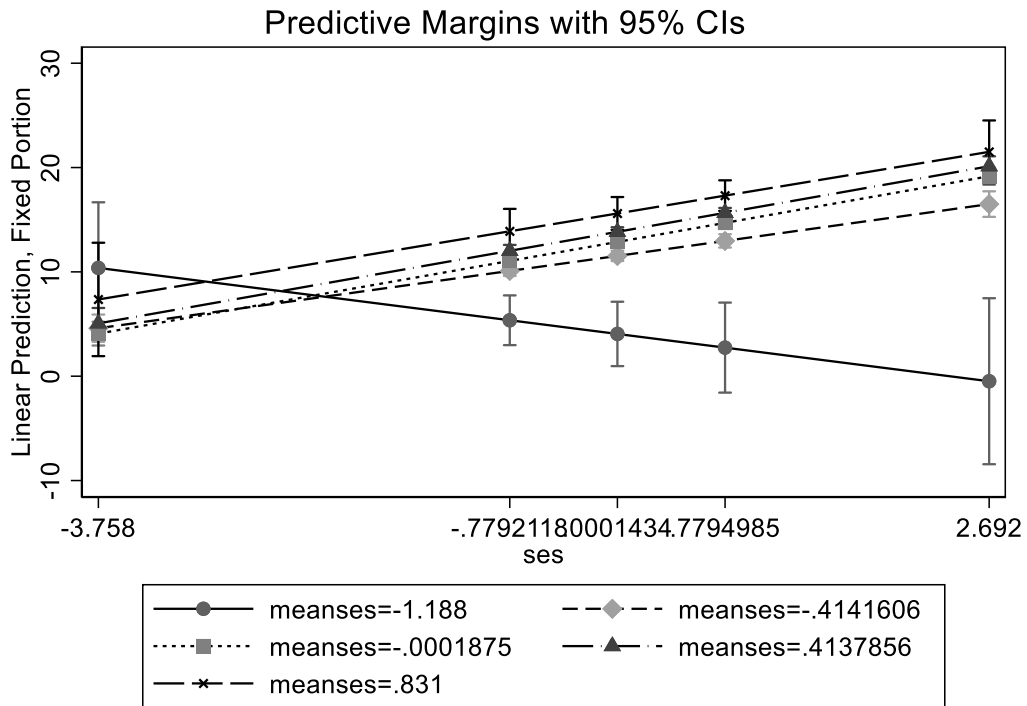
```

-----
|                               Delta-method
|                               Margin   Std. Err.   z   P>|z|   [95% Conf. Interval]
-----+-----
  _at |
  1 |   10.37874   3.209518   3.23  0.001    4.0882   16.66928
  2 |    4.600451  .6663532   6.90  0.000    3.294423   5.90648
  3 |    4.080304  .5811327   7.02  0.000    2.941305   5.219303
  4 |    5.065367  .751264   6.74  0.000    3.592916   6.537817
  5 |    7.355452  2.772092   2.65  0.008    1.922252   12.78865
  6 |    5.363829  1.216027   4.41  0.000    2.980459   7.747198
  7 |   10.09419  .2479459  40.71  0.000    9.60822   10.58015
  8 |   11.0489   .2189395  50.47  0.000   10.61979   11.47801
  9 |   12.01957  .2887416  41.63  0.000   11.45365   12.5855
 10 |   13.89045  1.095341  12.68  0.000   11.74362   16.03728
 11 |    4.051753  1.573768   2.57  0.010    .9672248   7.136281
 12 |   11.53154  .2505531  46.02  0.000   11.04046   12.02261
 13 |   12.87213  .18279   70.42  0.000   12.51387   13.23039
 14 |   13.83904  .2253969  61.40  0.000   13.39727   14.28081
 15 |   15.60024  .801719   19.46  0.000   14.0289   17.17158
 16 |    2.739676  2.201374   1.24  0.213   -1.574938   7.054291
 17 |   12.96889  .3257726  39.81  0.000   12.33039   13.60739
 18 |   14.69536  .2105043  69.81  0.000   14.28278   15.10794
 19 |   15.6585   .2385924  65.63  0.000   15.19087   16.12613
 20 |   17.31002  .7472577  23.16  0.000   15.84543   18.77462
 21 |   -0.4800974  4.058521  -0.12  0.906   -8.434652   7.474457
 22 |   16.49609  .625075   26.39  0.000   15.27096   17.72121
 23 |   19.16948  .4228441  45.33  0.000   18.34072   19.99824
 24 |   20.12338  .4857143  41.43  0.000   19.1714   21.07536
 25 |   21.50576  1.53443   14.02  0.000   18.49833   24.51318
-----

```

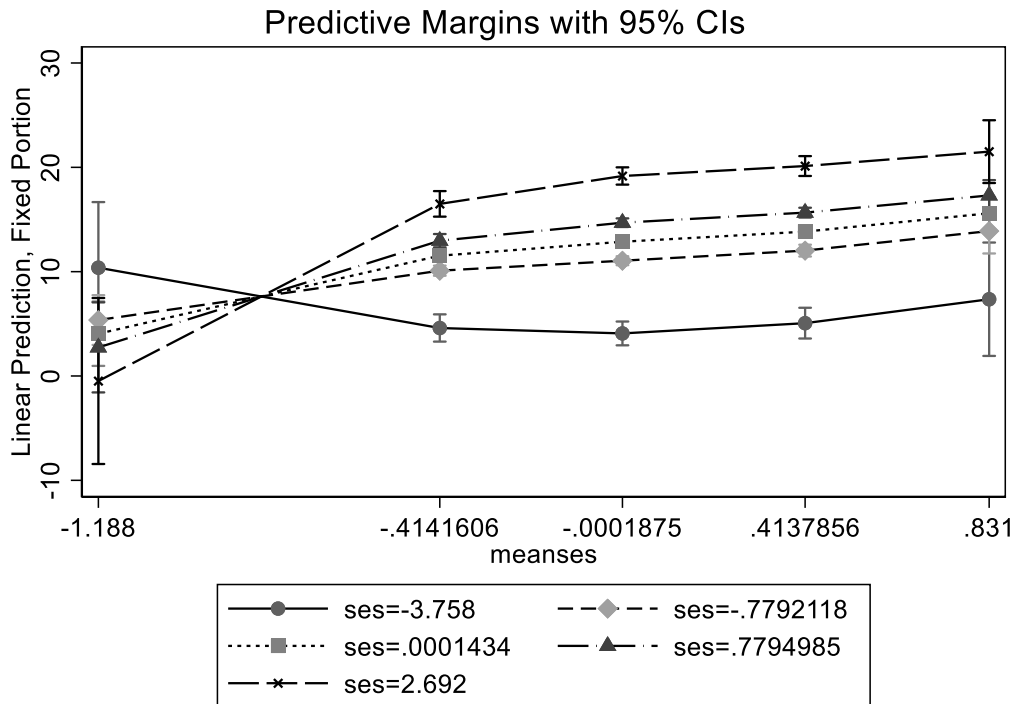
```
. marginsplot, x(meanses)
```

```
Variables that uniquely identify margins: ses meanses
```



```
. marginsplot, x(ses)
```

Variables that uniquely identify margins: ses meanse

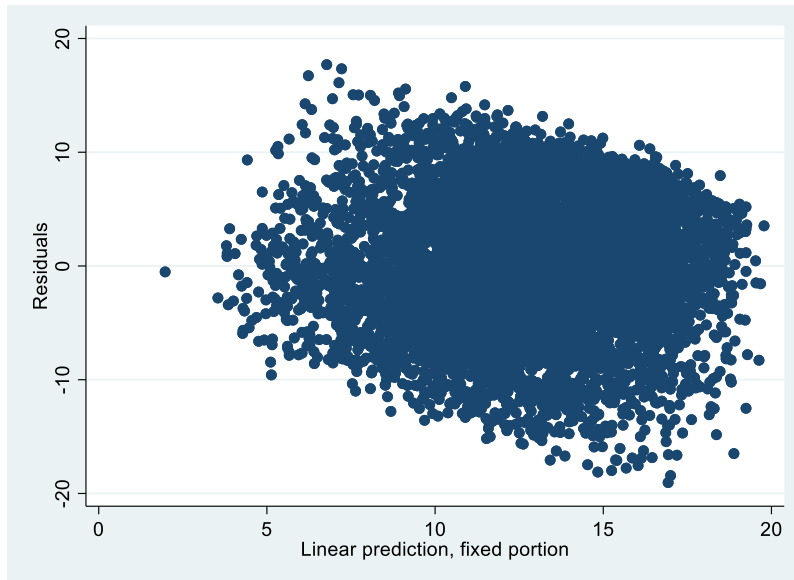


5. Homoscedasticity.

In HLM, the level-1 error terms should have equal variance across level-2 units (the assumption of homoscedasticity or homogeneity of variance) – e.g., all schools should have variances equal to the other schools in the sample.

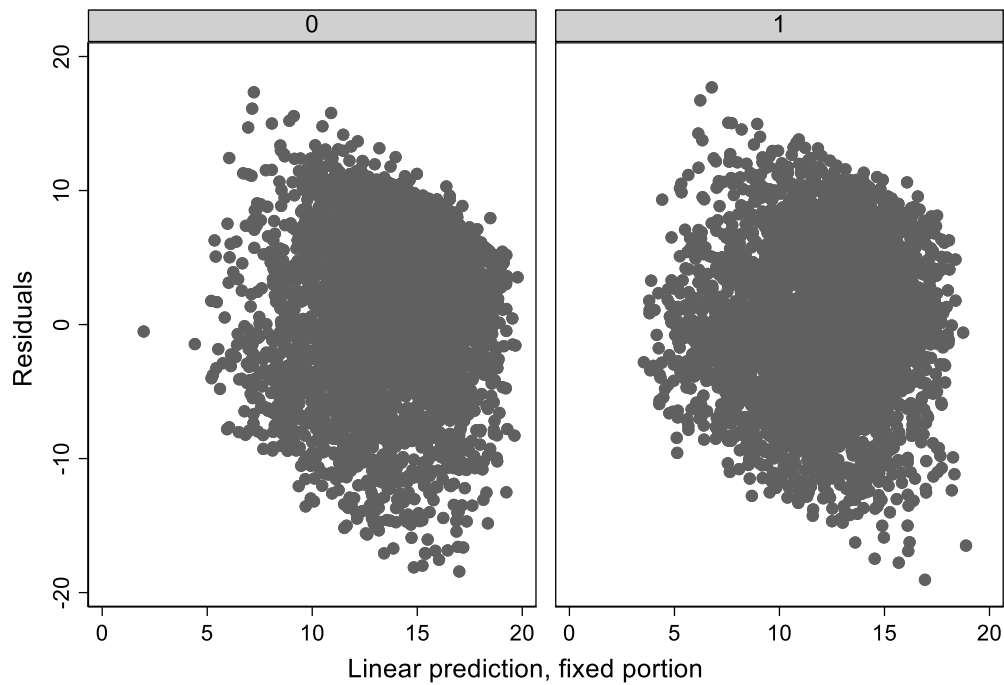
In order to graphically examine whether residual variance is heterogenous, we can look at an RVF plot – residuals vs. fitted values – we already constructed that plot above for the following model:

```
. qui mixed mathach c.ses##c.meanses c.ses##i.sector i.female##c.meanses  
i.female##i.sector || id: female, cov(unstr)  
  
. scatter llresid xb
```



We also already started to look at how residuals are distributed along the values of each individual predictor by constructing RVP plots -- residuals vs. predictors – for our continuous variables. However, for dichotomies, such plots are not very informative – however, we could create separate RVF plots by group to assess heterogeneity:

```
. graph twoway (scatter llresid xb, by(female))
```



Graphs by female

We can also assess residual variance by group using sum command:

```
. bysort female: sum llresid
```

```
-----
```

```
-> female = 0
```

Variable	Obs	Mean	Std. Dev.	Min	Max
llresid	3,390	-1.63e-09	6.137624	-18.42606	17.33541

```
-----
```

```
-> female = 1
```

Variable	Obs	Mean	Std. Dev.	Min	Max
llresid	3,795	3.99e-09	5.826746	-19.04129	17.70039

Heterogeneity of variance can be a nuisance. When it is a nuisance, the causes can be:

- One or more important level-1 predictors may have been omitted from the model.
- The effects of a level-1 predictor that is random or nonrandomly varying have been erroneously treated as fixed.
- Dependent variable is severely non-normal – skewed or kurtotic (has heavy tails).
- One (or more) of the independent variables has a nonlinear relationship to the dependent variable that we failed to model correctly.
- There are outliers or bad data.

We can deal with each of these problems; if that fails to remove heterogeneity, we can ultimately rely on robust SE. Robust standard errors are standard errors that are relatively insensitive to misspecification at the levels of the model and the distributional assumptions at each level. If the

robust and model-based standard errors differ substantially, that suggests that you have some problem with normality, homoscedasticity, or linearity, and you should further investigate those HLM assumptions. If it is not possible to correct the problem, you can report robust standard errors. Note, however, that the robust standard errors should be trusted only when the number of higher-level units is moderately large relative to the number of explanatory variables at the higher level.

Alternatively, heterogeneity of variance can be considered substantively interesting. In that case, we can model it using level 1 predictors – to see whether there are some predictors that seem to explain why level 1 variance is not uniform.

```
. mixed mathach c.ses##c.meanses c.ses##i.sector i.female##c.meanses
i.female##i.sector || id: female, cov(unstr) residuals(independent, by(female))
note: ses omitted because of collinearity
note: meanses omitted because of collinearity
```

```
Mixed-effects ML regression      Number of obs      =      7,185
Group variable: id              Number of groups   =      160

                                Obs per group:
                                min =      14
                                avg =     44.9
                                max =      67

                                Wald chi2(8)      =     855.62
                                Prob > chi2       =     0.0000
```

Log likelihood = -23213.302

mathach	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
ses	2.869241	.1473036	19.48	0.000	2.580531	3.157951
meanses	3.193401	.4964206	6.43	0.000	2.220435	4.166368
c.ses#						
c.meanses	.8249061	.2676124	3.08	0.002	.3003954	1.349417
ses	0	(omitted)				
1.sector	1.175409	.398656	2.95	0.003	.394058	1.956761
sector#c.ses						
1	-1.556612	.2219166	-7.01	0.000	-1.991561	-1.121664
1.female	-1.219793	.2381591	-5.12	0.000	-1.686577	-.7530101
meanses	0	(omitted)				
female#						
c.meanses						
1	-.0097155	.5062072	-0.02	0.985	-1.001863	.9824323
female#						
sector						
1 1	.0393173	.4180634	0.09	0.925	-.7800718	.8587064
_cons	12.71989	.2440476	52.12	0.000	12.24157	13.19822

Random-effects Parameters	Estimate	Std. Err.	[95% Conf. Interval]	
id: Unstructured				
var(female)	.9168451	.5663447	.2732121	3.076748

```

                var(_cons) |      2.79032   .5517962    1.893764    4.11133
                cov(female,_cons) |  -.9889858   .4754416   -1.920834   -.0571373
-----+-----
Residual: Independent,
  by female
                0: var(e) |      38.572   .9571921    36.74083    40.49443
                1: var(e) |      34.43275   .8047972    32.89096    36.04681
-----+-----

```

LR test vs. linear model: $\chi^2(4) = 201.88$ Prob > $\chi^2 = 0.0000$

Note: LR test is conservative and provided only for reference.

. estat ic

Akaike's information criterion and Bayesian information criterion

```

-----+-----
Model |          N   ll(null)  ll(model)    df        AIC        BIC
-----+-----
. |          7,185          .   -23213.3    14    46454.6    46550.92
-----+-----

```

Note: BIC uses N = number of observations. See [R] BIC note.

. est store efemale

. lrtest efemale baseline

```

Likelihood-ratio test                    LR  $\chi^2(1) =$     11.10
(Assumption: baseline nested in efemale) Prob >  $\chi^2 =$     0.0009

```

Note: The reported degrees of freedom assumes the null hypothesis is not on the boundary of the parameter space. If this is not true, then the reported test is conservative.

Looks like gender explains some heterogeneity – there is a higher amount of unexplained variance in math achievement among boys. We could assess model fit:

. est store egender

. estat ic

Akaike's information criterion and Bayesian information criterion

```

-----+-----
Model |          N   ll(null)  ll(model)    df        AIC        BIC
-----+-----
egender |          7,185          .   -23213.3    14    46454.6    46550.92
-----+-----

```

Note: BIC uses N = number of observations. See [R] BIC note.

```

. qui mixed mathach c.ses##c.meanses c.ses##i.sector i.female##c.meanses
i.female##i.sector || id: female, cov(unstr)

```

. est store baseline

. estat ic

Akaike's information criterion and Bayesian information criterion

```

-----+-----
Model |          N   ll(null)  ll(model)    df        AIC        BIC
-----+-----

```

```

. |          7,185          . -23218.85          13  46463.71  46553.14
-----

```

Note: BIC uses N = number of observations. See [R] BIC note.

```

. lrtest baseline egender

```

```

Likelihood-ratio test                    LR chi2(1) =    11.10
(Assumption: . nested in egender)       Prob > chi2 =    0.0009

```

Note: The reported degrees of freedom assumes the null hypothesis is not on the boundary of the parameter space. If this is not true, then the reported test is conservative.

LR test suggests that the model with heterogenous variance is preferred; however, BIC difference is only approximately 3, which is positive but not strong evidence in favor of that more complex model. We might want to explore what explains that higher unexplained variance among boys -- e.g., we could consider an interaction term of SES with gender (which would be an interaction of two level 1 variables), or examine additional predictors such as minority status etc.

We can also allow residual variance to vary by level 2 predictor groups:

```

. mixed mathach c.ses##c.meanses c.ses##i.sector i.female##c.meanses i.female#
> #i.sector || id: female, cov(unstr) residuals(independent, by(sector))
note: ses omitted because of collinearity
note: meanses omitted because of collinearity

```

```

Mixed-effects ML regression              Number of obs   =    7,185
Group variable: id                      Number of groups =    160

```

```

Obs per group:
      min =    14
      avg =   44.9
      max =    67

```

```

Log likelihood = -23205.16              Wald chi2(8)    =   822.03
                                          Prob > chi2     =    0.0000

```

mathach	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
ses	2.855303	.1528454	18.68	0.000	2.555731 3.154874	
meanses	3.193349	.5003151	6.38	0.000	2.21275 4.173949	
c.ses#						
c.meanses	.8257218	.2673987	3.09	0.002	.30163 1.349814	
ses	0	(omitted)				
1.sector	1.163307	.3995051	2.91	0.004	.3802913 1.946323	
sector#c.ses						
1	-1.550645	.2212156	-7.01	0.000	-1.98422 -1.117071	
1.female	-1.217677	.2453943	-4.96	0.000	-1.698641 -.7367136	
meanses	0	(omitted)				
female#						
c.meanses						
1	.0117788	.5131516	0.02	0.982	-.99398 1.017537	
female#						

```

sector |
  1 1 |   .051641   .417778   0.12   0.902   -.7671888   .8704709
      |
  _cons |  12.7195   .2479454   51.30   0.000   12.23354   13.20547
-----+-----
Random-effects Parameters |   Estimate   Std. Err.   [95% Conf. Interval]
-----+-----
id: Unstructured
      var(female) |   .9196108   .570399   .2726696   3.101498
      var(_cons) |   2.901038   .5536637   1.995728   4.217018
      cov(female,_cons) |  -1.080918   .4767802   -2.01539   -.1464463
-----+-----
Residual: Independent,
by sector
      0: var(e) |   39.55357   .9427131   37.74838   41.44509
      1: var(e) |   33.13314   .7991106   31.60336   34.73698
-----+-----
LR test vs. linear model: chi2(4) = 218.16          Prob > chi2 = 0.0000

```

Note: LR test is conservative and provided only for reference.

```
. estat ic
```

Akaike's information criterion and Bayesian information criterion

```

-----+-----
Model |           N   ll(null)   ll(model)   df       AIC       BIC
-----+-----
. |           7,185           .  -23205.16   14   46438.32   46534.64
-----+-----

```

Note: BIC uses N = number of observations. See [R] BIC note.

```
. est store esector
```

```
. lrtest baseline esector
```

```

Likelihood-ratio test                               LR chi2(1) =    27.39
(Assumption: baseline nested in esector)           Prob > chi2 =    0.0000

```

Note: The reported degrees of freedom assumes the null hypothesis is not on the boundary of the parameter space. If this is not true, then the reported test is conservative.