

Lab 1: Mixed Models

Georges Monette

Lab 1: Mixed Models

ICPSR Summer Program at York University

Georges Monette

May 2018

- Mixed Models Exercises Lab 1 —
— Setup —

- Introduction to Lab 1 —
 - Data: Math Achievement and SES —
 - * Exploring data —
 - Data: Levels and structure —
- Selecting a random subset of clusters —
- First look at variables —
 - Looking at Level 2 variables (invariant within schools) —
 - Creating additional Level 2 (and Level 1) variables with ‘capply’ —
 - Transformations of Level 1 variables within groups —
- Looking at data in 3D —
 - Looking at Level 1 and Level 2 data using Lattice graphics —
 - Visualizing fitted lines in beta space —
 - * Looking at between group effect —
 - Fitting a mixed model —
 - Convergence problems —
 - * In passing: handling NAs —
 - Hausman Test: Is the between effect different from the within effect? —
 - Fitting a model with a contextual mean —
 - Role of contextual variable for ses —
 - Interpreting the model with contextual effect —
 - Estimating the compositional effect (between effect) —

- Visualizing the fitted model —
 - * Effect plots with the effects package
 - * Handmade prediction plot
- Plotting error bars —
 - * Estimating the gap
- EXERCISES —
 - Using CWG instead of raw SES —
 - CWG vs raw in RE model —
- EXERCISES —
- REML vs ML —
 - Notes on testing: ML vs REML —
- Diagnostics —
 - Diagnostics with Level 1 residuals —
 - Scale - location plot: —
 - Diagnostics with Level 2 residuals —
 - Influence diagnostics – drop one row or one cluster at a time —
 - Exercises —

(Updated: May 27 2018 20:44)

Mixed Models Exercises Lab 1 —

Setup —

To install packages visit John Fox's <http://tinyurl.com/York-R-Course>

If necessary, update spida2:

```
#  
# devtools::install_github('gmonette/spida2')  
#  
# OR:  
#  
# On Windows  
# install.packages('http://blackwell.math.yorku.ca/R/spida2.zip', repos=NULL, type='source')  
#  
# On a Mac  
# install.packages("http://blackwell.math.yorku.ca/R/spida2.tgz", repos=NULL, type='source')  
#  
library(spida2)  
library(p3d)
```

Loading required package: rgl

Attaching package: 'rgl'

The following object is masked from 'package:spida2':

%>%

Attaching package: 'p3d'

The following objects are masked from 'package:spida2':

cell, center, ConjComp, dell, disp, ell, ell.conj, ellbox,
ellplus, ellpt, ellptc, ellpts, ellptsc, elltan, elltanc,
na.include, uv

library(car)

Loading required package: carData

Attaching package: 'car'

The following object is masked from 'package:p3d':

Identify3d

```
library(effects)
```

```
lattice theme set by effectsTheme()
```

```
See ?effectsTheme for details.
```

```
library(nlme)
```

Attaching package: 'nlme'

The following object is masked from 'package:spida2':

getData

```
library(lattice)
```

```
library(latticeExtra)
```

Loading required package: RColorBrewer

Introduction to Lab 1 —

At first, it's easier to learn by following a textbook example where nothing goes wrong. However, when you analyze real data lots of things tend to go wrong – or, more positively, many interesting things happen – and you wish you had experience with more realistic examples. These examples try to do both.

The first example is a gentle analysis in which we use the basic ideas of multilevel models with mixed models. We learn how to formulate and test general linear hypotheses that allow us to ask a much wider range of questions than those accessible through standard output.

The second example, using a different model with the same data, illustrates what to do when ‘everything’ goes wrong, oops, I meant “when a lot of interesting things happen.”

In the second example, the first model does not converge and we have to figure out what to do. We need to test random effects variance parameters that are not tested validly with the usual methods and we learn how to use simulation for this purpose.

Our analysis of BLUPS reveals problems with the model whose correction we consider.

#

The second sample analysis is quite long but it is amply annotated so that you can follow much of it on your own.

Data: Math Achievement and SES —

Math Achievement and SES in a sample of Public and Catholic high schools

Goal: Study the relationship between SES and Mathach in Public vs Catholic schools

Exploring data —

```
?hsfull
```

```
starting httpd help server ... done
```

```
dim( hsfull )
```

```
[1] 7185    9
```

```
head(hsfull)
```

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM
1	1224	5.876	-1.528	Female	No	842	Public	0.35	1.597
2	1224	19.708	-0.588	Female	No	842	Public	0.35	1.597
3	1224	20.349	-0.528	Male	No	842	Public	0.35	1.597
4	1224	8.781	-0.668	Male	No	842	Public	0.35	1.597
5	1224	17.898	-0.158	Male	No	842	Public	0.35	1.597

6	1224	4.583	0.022	Male	No	842	Public	0.35	1.597
---	------	-------	-------	------	----	-----	--------	------	-------

hsfull %>% head *# same as head(hsfull)*

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM
1	1224	5.876	-1.528	Female	No	842	Public	0.35	1.597
2	1224	19.708	-0.588	Female	No	842	Public	0.35	1.597
3	1224	20.349	-0.528	Male	No	842	Public	0.35	1.597
4	1224	8.781	-0.668	Male	No	842	Public	0.35	1.597
5	1224	17.898	-0.158	Male	No	842	Public	0.35	1.597
6	1224	4.583	0.022	Male	No	842	Public	0.35	1.597

hsfull %>% some *# a random selection of rows*

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM
285	1436	9.561	0.212	Male	No	185	Catholic	1.00	-1.141
1148	2526	9.843	0.762	Female	No	572	Catholic	0.97	-2.154
1425	2658	16.587	0.512	Female	No	780	Catholic	0.79	-0.961
1496	2768	12.812	0.092	Male	No	1680	Public	0.31	0.086
2255	3533	10.121	-0.418	Female	No	1015	Catholic	0.54	0.022
3019	4292	7.128	-1.148	Male	Yes	1328	Catholic	0.76	-0.674
3392	4530	8.898	-1.848	Female	Yes	435	Catholic	0.60	-0.245
4351	6170	10.991	-0.948	Male	No	2126	Public	0.25	0.432

4978	6990	4.756	1.292	Female	Yes	1650	Public	0.59	0.475
5192	7276	4.027	-0.438	Female	No	1630	Public	0.30	0.109

hsfull %>% tail

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM
7180	9586	20.967	1.612	Female	Yes	262	Catholic	1	-2.416
7181	9586	20.402	1.512	Female	No	262	Catholic	1	-2.416
7182	9586	14.794	-0.038	Female	No	262	Catholic	1	-2.416
7183	9586	19.641	1.332	Female	No	262	Catholic	1	-2.416
7184	9586	16.241	-0.008	Female	No	262	Catholic	1	-2.416
7185	9586	22.733	0.792	Female	No	262	Catholic	1	-2.416

A 'uniform quantile' plot. Data is lined up from shortest to tallest.

`xqplot( hsfull )` *# gives an idea of distribution, outliers, granularity, etc. a*

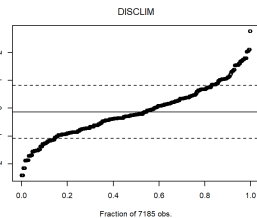
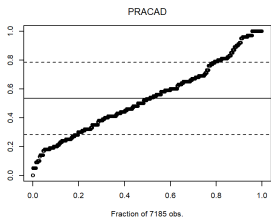
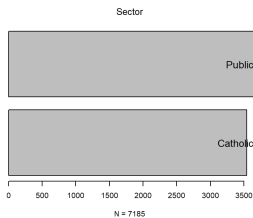
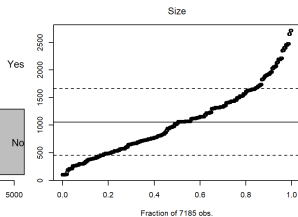
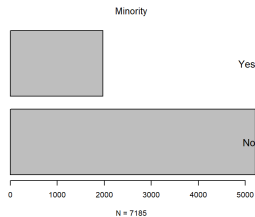
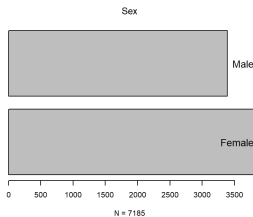
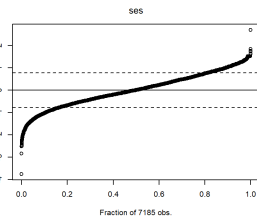
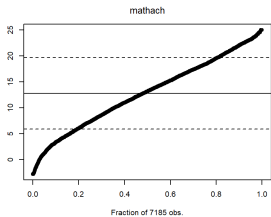
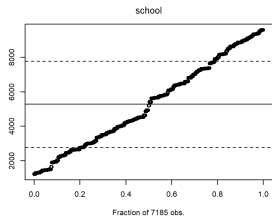


Figure: A **uniform quantile** plot. Data are lined up from shortest to tallest uniformly spaced on the x-axis. If a variable is close to uniform, its quantile plot will be close to a diagonal.

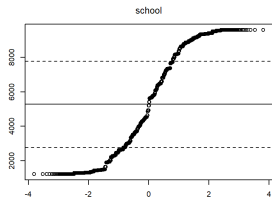
Think of a classroom photo with kids lined up from shortest to tallest

Solid line at mean, dashed lines at $\pm$ one standard deviation

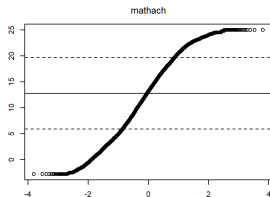
Find quantiles by selecting a proportion on x-axis and read off height of graph on y-axis (use the edge of sheet of paper) e.g. .8-quantile = 80th percentile of ses is ~ 0.8

Note: with an approximately normal distribution we expect mean $\pm$ std. dev. to be at approx the 32nd and 68th percentiles.

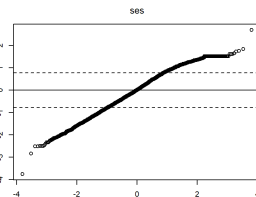
```
xqplot( hsfull , ptype='normal')
```



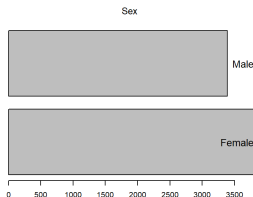
Normal quantile for 7185 obs.



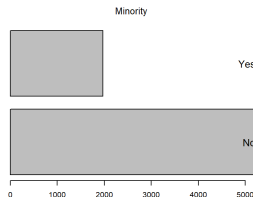
Normal quantile for 7185 obs.



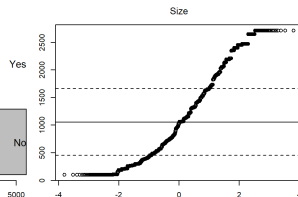
Normal quantile for 7185 obs.



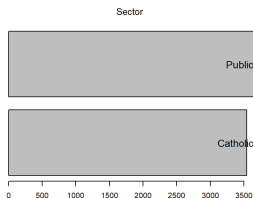
N = 7185



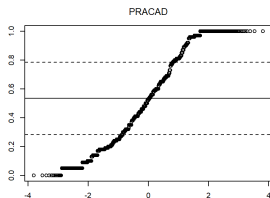
N = 7185



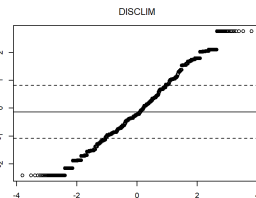
Normal quantile for 7185 obs.



N = 7185



Normal quantile for 7185 obs.



Normal quantile for 7185 obs.

Figure: A **normal quantile** plot. Data are lined up from shortest to tallest and x-axis is stretched at the ends to have shape of ideal normal. If a variable is close to normal, its quantile plot will be close to a diagonal

Data: Levels and structure —

Data structures, part 1: moving up one level

‘hsfull’ has one student per row.

It is a **long** file combining variables of all levels. If we want information on schools, we can create a data set that has one row per school, keeping only those variable that are invariant within schools, i.e. Level 2 variables

```
hsup <- up( hsfull, ~ school )  # one row per school with Level 2 variables  
some( hsup )
```

	school	Size	Sector	PRACAD	DISCLIM
3013	3013	760	Public	0.56	-0.213
3351	3351	1125	Public	0.35	0.383
3498	3498	388	Catholic	0.92	-1.556
4350	4350	1661	Public	0.62	0.512
4931	4931	603	Catholic	0.79	-1.172

5762	5762	1826	Public	0.24	0.364
7232	7232	1154	Public	0.20	0.975
7919	7919	1451	Public	0.50	-0.402
8775	8775	667	Public	0.28	1.783
9021	9021	472	Catholic	0.82	-0.817

```
dim( hsup )
```

```
[1] 160    5
```

```
tab(hsup, ~ Sector)
```

Sector		
Catholic	Public	Total
70	90	160

gicc: levels of variables in hsfull

Generalized intraclass correlation:

- intraclass correlation (variance between / total variance) of ranks for numerical variables
- Goodman-Kruskal's tau for factor and character variables

Interpretation:

- 1 means it's a level-2 or school-level variable,
- 0 means that it's mean is the same for each school, e.g. if it's balanced

```
gicc( hsfull, ~ school, type = 'raw' ) # uses variance between / total variance
```

	school	mathach	ses	Sex	Minority	Size	Sector
1.0000000	0.1955041	0.2749462	0.2869238	0.4644279	1.0000000	1.0000000	
PRACAD	DISCLIM						
1.0000000	1.0000000						

```
gicc( hsfull, ~ school ) # uses ranks by default
```

	school	mathach	ses	Sex	Minority	Size	Sector
1.0000000	0.1955041	0.2749462	0.2869238	0.4644279	1.0000000	1.0000000	
PRACAD	DISCLIM						
1.0000000	1.0000000						

```
tab_( hsfull, ~ school ) # school size: 'tab_' is a variant of 'tab' with
```

school															
1224	1288	1296	1308	1317	1358	1374	1433	1436	1461	1462	1477	1499	1637	1906	
47	25	48	20	48	30	28	35	44	33	57	62	53	27	53	
1909	1942	1946	2030	2208	2277	2305	2336	2458	2467	2526	2626	2629	2639	2651	
28	29	39	47	60	61	67	47	57	52	57	38	57	42	38	

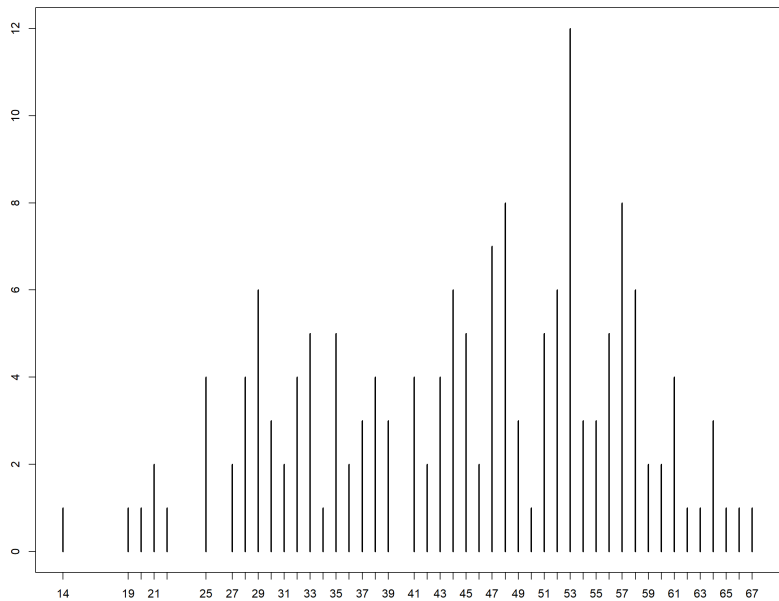
2655	2658	2755	2768	2771	2818	2917	2990	2995	3013	3020	3039	3088	3152	3332
52	45	47	25	55	42	43	48	46	53	59	21	39	52	38
3351	3377	3427	3498	3499	3533	3610	3657	3688	3705	3716	3838	3881	3967	3992
39	45	49	53	38	48	64	51	43	45	41	54	41	52	53
3999	4042	4173	4223	4253	4292	4325	4350	4383	4410	4420	4458	4511	4523	4530
46	64	44	45	58	65	53	33	25	41	32	48	58	47	63
4642	4868	4931	5192	5404	5619	5640	5650	5667	5720	5761	5762	5783	5815	5819
61	34	58	28	57	66	57	45	61	53	52	37	29	25	50
5838	5937	6074	6089	6144	6170	6291	6366	6397	6415	6443	6464	6469	6484	6578
31	29	56	33	43	21	35	58	60	54	30	29	57	35	56
6600	6808	6816	6897	6990	7011	7101	7172	7232	7276	7332	7341	7342	7345	7364
56	44	55	49	53	33	28	44	52	53	48	51	58	56	44
7635	7688	7697	7734	7890	7919	8009	8150	8165	8175	8188	8193	8202	8357	8367
51	54	32	22	51	37	47	44	49	33	30	43	35	27	14
8477	8531	8627	8628	8707	8775	8800	8854	8857	8874	8946	8983	9021	9104	9158
37	41	53	61	48	48	32	32	64	36	58	51	56	55	53
9198	9225	9292	9340	9347	9359	9397	9508	9550	9586					
31	36	19	29	57	53	47	35	29	59					

```
tab_( hsfull, ~ school ) %>% c %>% tab # distribution of school sizes
```

14	19	20	21	22	25	27	28	29	30	31	32
1	1	1	2	1	4	2	4	6	3	2	4
33	34	35	36	37	38	39	41	42	43	44	45
5	1	5	2	3	4	3	4	2	4	6	5
46	47	48	49	50	51	52	53	54	55	56	57
2	7	8	3	1	5	6	12	3	3	5	8
58	59	60	61	62	63	64	65	66	67	Total	
6	2	2	4	1	1	3	1	1	1	160	

```
tab_( hsfull, ~ school ) %>% c %>% tab_ %>% plot
```

```
tab_( hsfull, ~ school ) %>% as.vector %>% tab_ %>% plot
```



Level-2 variables

```
hsup <- up( hsfull, ~ school ) # 'up' one level in the multilevel hierarchy
```

To include summaries of selected Level-1 variables.

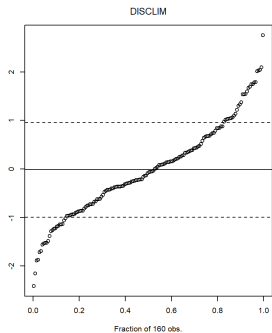
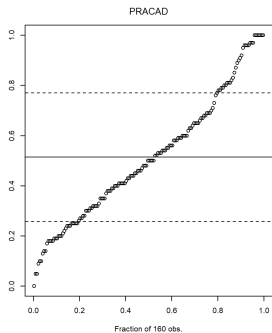
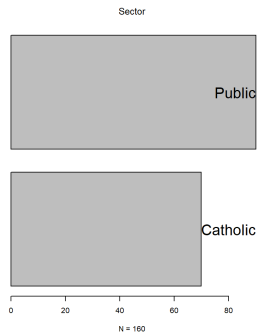
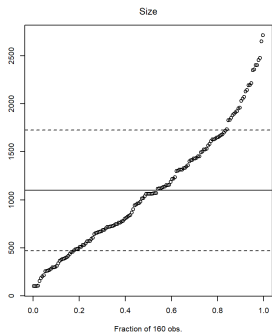
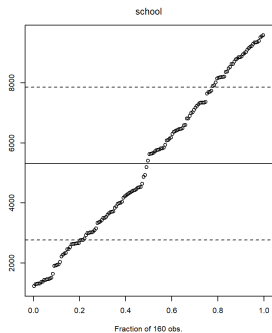
For factors, this produces the mean of the incidence matrix within each school, i.e. the proportion of students in each category of the factor.

For each factor, these proportions must sum to 1.

```
hsfull %>%  
  up( ~ school, agg = ~ Sex + ses ) %>%  
  head
```

	school	Size	Sector	PRACAD	DISCLIM	Sex_Female	Sex_Male	ses
1224	1224	842	Public	0.35	1.597	0.5957447	0.4042553	-0.43438298
1288	1288	1855	Public	0.27	0.174	0.4400000	0.5600000	0.12160000
1296	1296	1719	Public	0.32	-0.137	0.6458333	0.3541667	-0.42550000
1308	1308	716	Catholic	0.96	-0.622	0.0000000	1.0000000	0.52800000
1317	1317	455	Catholic	0.95	-1.694	1.0000000	0.0000000	0.34533333
1358	1358	1430	Public	0.25	1.535	0.3666667	0.6333333	-0.01966667

```
xqplot( hsup )
```



To speed up calculations, we will use a (pre-)randomly selected quarter of the schools.

However, since it's a good idea to know that it's easy to select a random subset of schools, **not students**, here's how it's done.

Selecting a random subset of clusters —

We can't randomly sample from the long file if we want each cluster (school) to be either all in or all out.

There are a few ways of selecting all the observations from a sample of clusters. The following seems fairly intuitive.

We first create the school level file and take a sample of school from that file. We then merge the sample file with the longfile. Merge will match with variables that have the same name. By default, it only uses records that match in both files, so it produces the result we want.

```
hsup <- up ( hsup, ~ school)  # as above
dim(hsup)                    # one row per school

[1] 160    5
```

We'll use a randomly selected subset of half the schools

This has another interesting consequence: we can do a **split-half analysis** in which the model developed on one half of the clusters is validated with the other half

The following code shows how to split the clusters into two halves although we'll be using two 'preselected' halves

How to select a random sample of 5 numbers from the numbers 1 to 10:

```
sample(1:10, 5)
```

```
[1] 5 9 7 3 6
```

Select a sample of half the row indices:

```
selected_rows <- sample( 1:nrow(hsup), round( nrow(hsup)/2 ))
```

```
selected_rows
```

```
[1] 113  47  36  59 126 138 145  28  57  46 129  62  60  19  90  38 150
[18]  31 149 106  30  61  54 102  91   6  32 157 139  67  68   9 140  11
[35]  82  72  27 128  78  87  56  13  85  58 124 101 121 155  97 119 100
[52]  49  79 103  39  99  84  76  40 156  75  88 123  69  33   4  80 125
[69] 111   8  66 152  77 110  55  18  53   3  71 122
```

```
hsu1 <- hsup[selected_rows, ]           # first half of schools
```

```
hsu2 <- hsup[ - selected_rows, ]       # other half
```

```
hs1 <- merge( hsu1, hsfull )      # long data for 1st half of schools
hs2 <- merge( hsu2, hsfull )      # long data for other half
```

```
dim(hs1)
```

```
[1] 3687    9
```

```
dim(hs2)
```

```
[1] 3498    9
```

```
length(unique(hs1$school))
```

```
[1] 80
```

```
length(unique(hs2$school))
```

```
[1] 80
```

But we will use an even smaller preselected subset of 40 schools

```
data(hs)
```

```
dd <- hs                      # easier to type repeatedly
```

```
dim( dd )
```

```
[1] 1977    9
```

```
some(dd)
```

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM
58	1906	14.171	0.602	Male	No	400	Catholic	0.87	-0.939
268	2629	14.691	-0.768	Male	No	1314	Catholic	0.81	-0.613
341	2639	6.166	-0.888	Female	Yes	2713	Public	0.14	-0.282
557	3610	13.942	0.602	Male	No	1431	Catholic	0.80	-0.621
915	5640	23.048	0.632	Male	No	1152	Public	0.41	0.256
916	5640	18.066	-0.038	Female	No	1152	Public	0.41	0.256
926	5640	18.225	0.332	Male	No	1152	Public	0.41	0.256
987	5650	20.907	-0.768	Female	Yes	720	Catholic	0.60	-0.070
1734	8627	13.946	-0.018	Male	No	2452	Public	0.25	0.742
1811	8707	8.432	-0.238	Male	Yes	1133	Public	0.48	1.542

First look at variables —

```
summary( dd )
```

school

mathach

ses

Sex

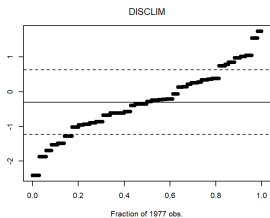
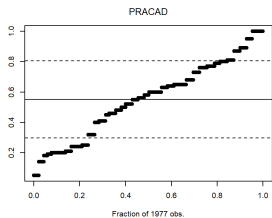
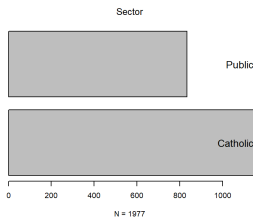
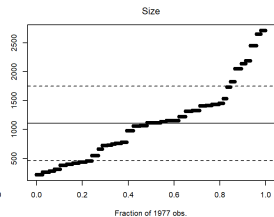
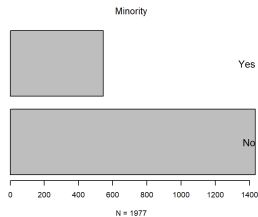
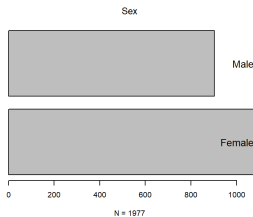
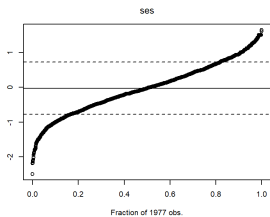
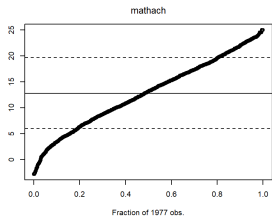
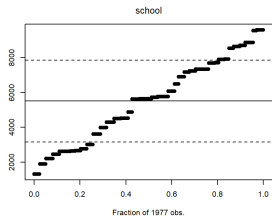
Min.	:1317	Min.	:-2.832	Min.	:-2.49800	Female:1074
1st Qu.:	3013	1st Qu.:	7.529	1st Qu.:	-0.55800	Male : 903
Median	:5650	Median	:13.095	Median	:-0.02800	
Mean	:5507	Mean	:12.783	Mean	:-0.02684	
3rd Qu.:	7345	3rd Qu.:	18.336	3rd Qu.:	0.52200	
Max.	:9586	Max.	:24.993	Max.	: 1.65200	

Minority	Size	Sector	PRACAD
No :1433	Min. : 215	Catholic:1144	Min. :0.0500
Yes: 544	1st Qu.: 545	Public : 833	1st Qu.:0.3200
	Median :1114		Median :0.5800
	Mean :1106		Mean :0.5513
	3rd Qu.:1415		3rd Qu.:0.7600
	Max. :2713		Max. :1.0000

DISCLIM

Min.	:-2.4160
1st Qu.:	-0.9390
Median	:-0.2820
Mean	:-0.3022
3rd Qu.:	0.3360
Max.	: 1.7420

xqplot(dd)



```
library(Hmisc, pos = 1000) # to avoid masking functions in other packages

Loading required package: survival
Loading required package: Formula
Loading required package: ggplot2

Attaching package: 'ggplot2'

The following object is masked from 'package:latticeExtra':

    layer

The following object is masked from 'package:spida2':

    labs

Attaching package: 'Hmisc'

The following object is masked _by_ 'package:p3d':

    na.include
```

The following objects are masked `_by_ 'package:spida2':`

`fillin, Lag, na.include`

The following objects are masked from `'package:base':`

`format.pval, units`

`describe(dd)`

`dd`

9 Variables 1977 Observations

`school`

n	missing	distinct	Info	Mean	Gmd	.05	.10
1977	0	40	0.999	5507	2695	1906	2458
.25	.50	.75	.90	.95			
3013	5650	7345	8707	8874			

lowest : 1317 1906 2208 2458 2626, highest: 8707 8854 8874 9550 9586

`mathach`

n	missing	distinct	Info	Mean	Gmd	.05	.10
1977	0	1860	1	12.78	7.863	1.363	3.369
.25	.50	.75	.90	.95			
7.529	13.095	18.336	21.892	23.220			

lowest : -2.832 -2.721 -2.658 -2.557 -2.550, highest: 24.488 24.545 24.707 24.707

ses

n	missing	distinct	Info	Mean	Gmd	.05	.10
1977	0	323	1	-0.02684	0.8577	-1.270	-0.998
.25	.50	.75	.90	.95			
-0.558	-0.028	0.522	0.952	1.202			

lowest : -2.498 -2.188 -2.178 -2.128 -2.108, highest: 1.452 1.462 1.512 1.512

Sex

n	missing	distinct
1977	0	2

Value	Female	Male
Frequency	1074	903

Proportion 0.543 0.457

Minority

	n	missing	distinct
1977		0	2

Value	No	Yes
-------	----	-----

Frequency	1433	544
-----------	------	-----

Proportion	0.725	0.275
------------	-------	-------

Size

	n	missing	distinct	Info	Mean	Gmd	.05	.10
1977		0	40	0.999	1106	714.7	262	311
	.25	.50	.75	.90	.95			
	545	1114	1415	2142	2452			

lowest : 215 262 280 311 381, highest: 2142 2190 2452 2650 2713

Sector

	n	missing	distinct
1977		0	2

Value	Catholic	Public
Frequency	1144	833
Proportion	0.579	0.421

PRACAD

n	missing	distinct	Info	Mean	Gmd	.05	.10
1977	0	34	0.999	0.5513	0.2903	0.180	0.200
.25	.50	.75	.90	.95			
0.320	0.580	0.760	0.878	0.950			

lowest : 0.05 0.14 0.18 0.19 0.20, highest: 0.81 0.87 0.89 0.95 1.00

DISCLIM

n	missing	distinct	Info	Mean	Gmd	.05	.10
1977	0	40	0.999	-0.3022	1.054	-1.872	-1.534
.25	.50	.75	.90	.95			
-0.939	-0.282	0.336	0.975	1.048			

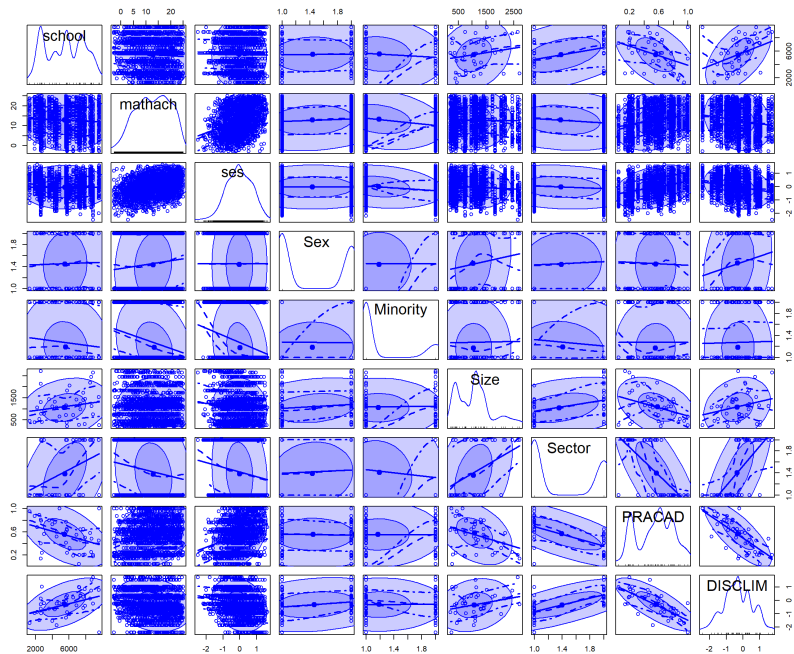
lowest : -2.416 -1.872 -1.694 -1.534 -1.484, highest: 0.975 1.013 1.048 1

QUESTIONS: looking at univariate distributions for this data:

Any problems with: NAs? Highly skewed distributions? Possible univariate outliers? Skewed factors: i.e. very rare category

Any actions to take?

```
library(car)  
scatterplotMatrix( dd , ellipse = TRUE)    # 2-dim summary
```



QUESTIONS: looking at bivariate relationships

Hint: focus on one column or one row at a time

Bivariate relationships that stand out?

Bivariate outliers?

Curvilinear bivariate relationships?

Which pair has the 'strongest' bivariate relationship?

Looking at Level 2 variables (invariant within schools) —

```
ddu <- up( dd, ~ school)  # only variables invariant within schools
dim(ddu)
```

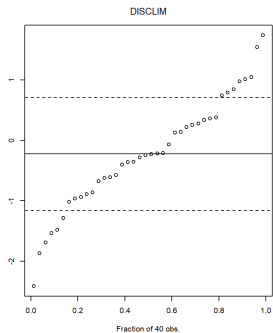
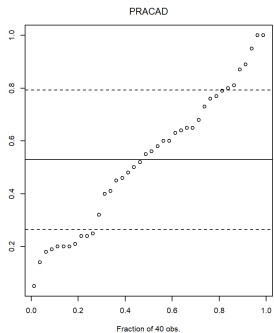
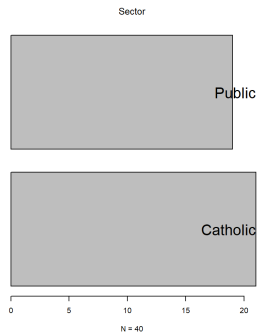
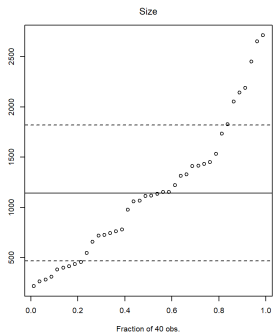
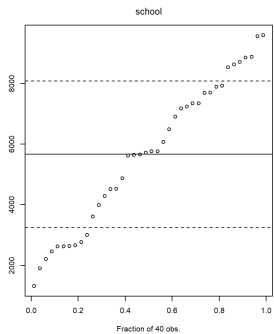
```
[1] 40  5
```

```
some(ddu)
```

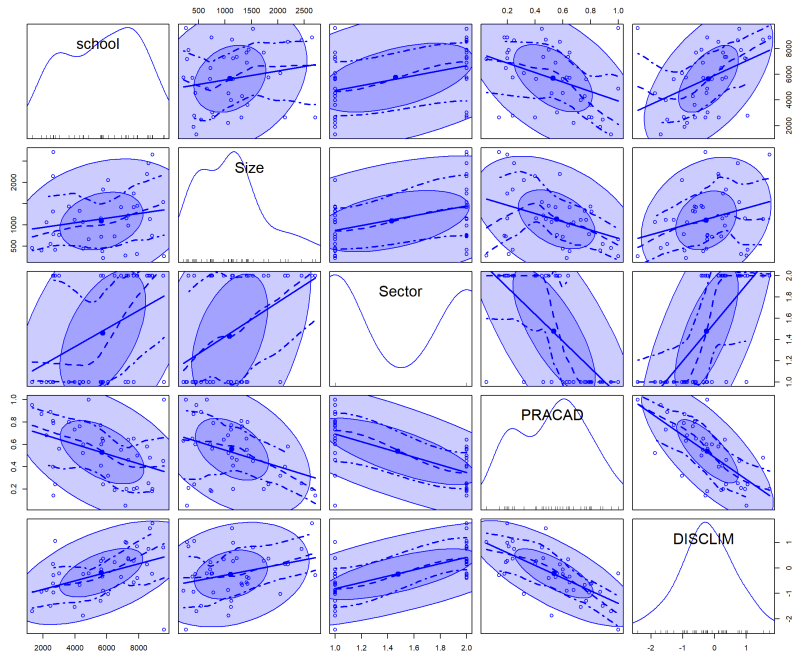
	school	Size	Sector	PRACAD	DISCLIM
2658	2658	780	Catholic	0.79	-0.961
2771	2771	415	Public	0.24	1.048
5650	5650	720	Catholic	0.60	-0.070
5761	5761	215	Catholic	0.63	-0.892

5762	5762	1826	Public	0.24	0.364
6074	6074	2051	Catholic	0.32	-1.018
7232	7232	1154	Public	0.20	0.975
7342	7342	1220	Catholic	0.46	0.380
8854	8854	745	Public	0.18	-0.228
9586	9586	262	Catholic	1.00	-2.416

```
xqplot( ddu )      # shows just variables invariant within schools
```



```
scatterplotMatrix( ddu , ell = T)    # between school variables
```



Include derived Level 2 summaries of Level 1 variables

- Level 2 means of Level 1 numerical variables
- Level 2 **modes** of Level 1 categorical variables (not very useful for analysis)
- Recall that **agg** returns the mean of incidence matrix

```
up(dd, ~school) %>% head
```

	school	Size	Sector	PRACAD	DISCLIM
1317	1317	455	Catholic	0.95	-1.694
1906	1906	400	Catholic	0.87	-0.939
2208	2208	1061	Catholic	0.68	-0.864
2458	2458	545	Catholic	0.89	-1.484
2626	2626	2142	Public	0.40	0.142
2629	2629	1314	Catholic	0.81	-0.613

```
ddall <- up (dd, ~school, all = T)
```

```
head(ddall)
```

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD
1317	1317	13.17769	0.34533333	Female	Yes	455	Catholic	0.95
1906	1906	15.98317	0.51162264	Female	No	400	Catholic	0.87
2208	2208	15.40467	0.42316667	Female	No	1061	Catholic	0.68
2458	2458	13.98568	0.22778947	Female	Yes	545	Catholic	0.89

2626	2626	13.39661	-0.06484211	Male	No	2142	Public	0.40
2629	2629	14.90777	-0.13764912	Male	No	1314	Catholic	0.81

DISCLIM

1317	-1.694
1906	-0.939
2208	-0.864
2458	-1.484
2626	0.142
2629	-0.613

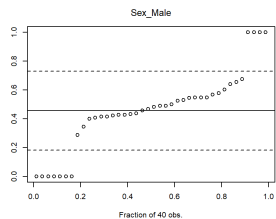
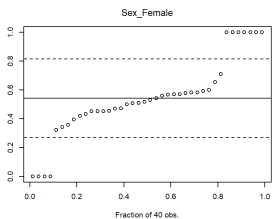
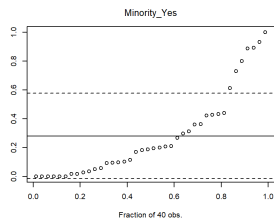
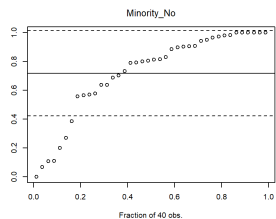
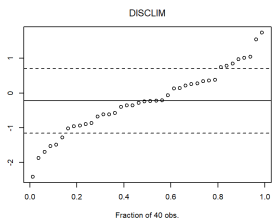
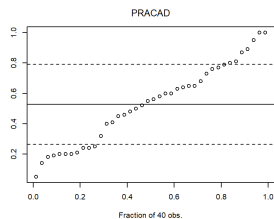
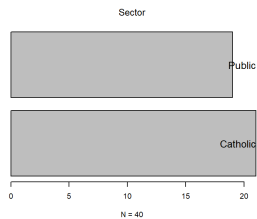
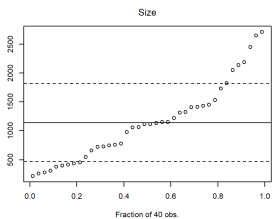
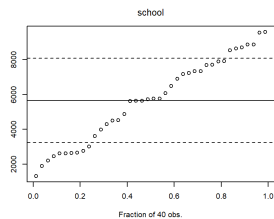
To get distribution of proportions for factors – not just modes

```
ddagg <- up(dd, ~school, agg = ~ Minority + Sex)
head(ddagg)
```

	school	Size	Sector	PRACAD	DISCLIM	Minority_No	Minority_Yes
1317	1317	455	Catholic	0.95	-1.694	0.2708333	0.72916667
1906	1906	400	Catholic	0.87	-0.939	0.9056604	0.09433962
2208	2208	1061	Catholic	0.68	-0.864	1.0000000	0.00000000
2458	2458	545	Catholic	0.89	-1.484	0.3859649	0.61403509
2626	2626	2142	Public	0.40	0.142	1.0000000	0.00000000
2629	2629	1314	Catholic	0.81	-0.613	0.7894737	0.21052632
	Sex_Female		Sex_Male				

1317	1.0000000	0.0000000
1906	0.5094340	0.4905660
2208	0.5833333	0.4166667
2458	1.0000000	0.0000000
2626	0.4736842	0.5263158
2629	0.0000000	1.0000000

```
xqplot( ddagg )
```



```
spm( ddagg, ell = T )  # spm = scatterplotMatrix for slow typists
```

```
Warning in MASS::cov.trob(cbind(x[use], y[use]), wt = weights[use]):  
Probable convergence failure
```

```
Warning in chol.default(shape, pivot = TRUE): the matrix is either rank-  
deficient or indefinite
```

```
Warning in chol.default(shape, pivot = TRUE): the matrix is either rank-  
deficient or indefinite
```

```
Warning in MASS::cov.trob(cbind(x[use], y[use]), wt = weights[use]):  
Probable convergence failure
```

```
Warning in MASS::cov.trob(cbind(x[use], y[use]), wt = weights[use]):  
Probable convergence failure
```

```
Warning in chol.default(shape, pivot = TRUE): the matrix is either rank-  
deficient or indefinite
```

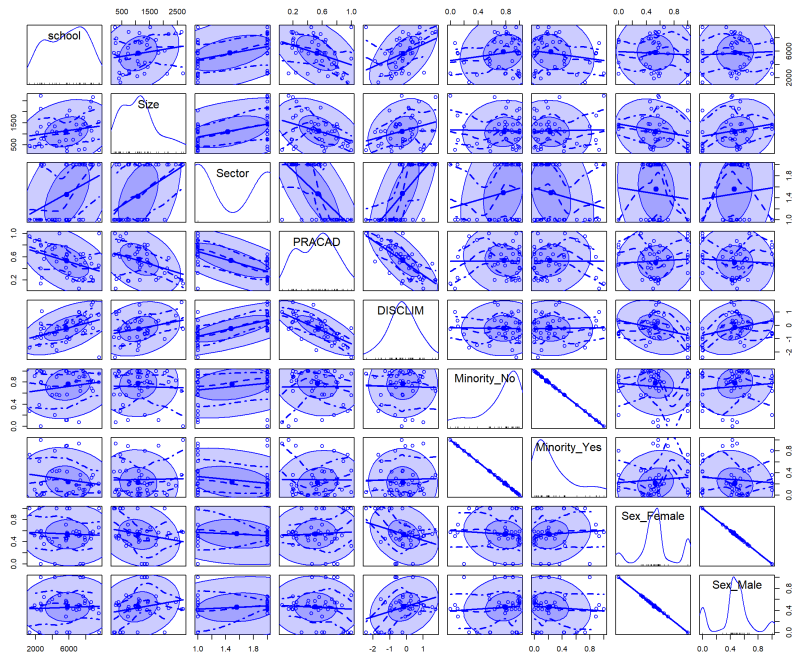
```
Warning in chol.default(shape, pivot = TRUE): the matrix is either rank-  
deficient or indefinite
```

```
Warning in MASS::cov.trob(cbind(x[use], y[use]), wt = weights[use]):
```

Probable convergence failure

Warning in chol.default(shape, pivot = TRUE): the matrix is either rank-deficient or indefinite

Warning in chol.default(shape, pivot = TRUE): the matrix is either rank-deficient or indefinite



QUESTIONS:

- Note that there is a relationship between minority and ses as well as between minority and mathach
- Does anything stand out in ‘between-school’ relationships?
- Classify variables:
 - Level 1 variables
 - Level 2 variables:
 - * contextual variables: natural (properties of cluster) and
 - * compositional: derived cluster values from individual values
- Does this suggest any multilevel questions?
- Are there derived level 2 (compositional) variables that could be relevant?

Creating additional Level 2 (and Level 1) variables with ‘capply’ —

An important part of modeling is being able to easily create the variables that capture the phenomena you want to study. Often, these variables are not in the raw data set.

Examples:

Sample size:

`capply(y, cluster, fun)`

1. divides 'y' into chunks for each value of 'cluster'
2. applies the function 'fun' to each chunk
3.
 - a. if 'fun' returns a vector of the same length as the chunk of 'y' then 'capply' returns that vector in the positions corresponding to the values of 'y'
 - b. if 'fun' returns a single value, that value gets returned in every position of 'y'

How many observations within each school?

```
dd$n <- capply( dd$school, dd$school, length)
```

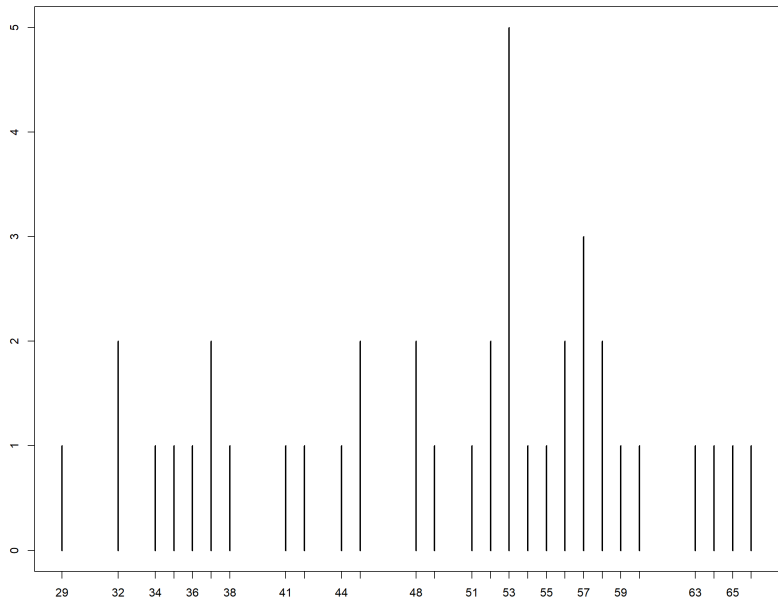
```
head( dd ) # to see that new variable is constant in first school
```

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM	n
1	1317	12.862	0.882	Female	No	455	Catholic	0.95	-1.694	48
2	1317	8.961	0.932	Female	Yes	455	Catholic	0.95	-1.694	48
3	1317	4.756	-0.158	Female	Yes	455	Catholic	0.95	-1.694	48
4	1317	21.405	0.362	Female	Yes	455	Catholic	0.95	-1.694	48
5	1317	20.748	1.372	Female	No	455	Catholic	0.95	-1.694	48
6	1317	18.362	0.132	Female	Yes	455	Catholic	0.95	-1.694	48

```
some( dd )
```

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM	n
--	--------	---------	-----	-----	----------	------	--------	--------	---------	---


```
dd %>%  
  {up(., ~school)$n} %>%  
  tab_ %>%  
  plot
```



mean ses by school (sample compositional ses)

```
dd$ses.m <- with( dd, capply(ses, school, mean, na.rm = TRUE ))  
head( dd )
```

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM	n
1	1317	12.862	0.882	Female	No	455	Catholic	0.95	-1.694	48
2	1317	8.961	0.932	Female	Yes	455	Catholic	0.95	-1.694	48
3	1317	4.756	-0.158	Female	Yes	455	Catholic	0.95	-1.694	48
4	1317	21.405	0.362	Female	Yes	455	Catholic	0.95	-1.694	48
5	1317	20.748	1.372	Female	No	455	Catholic	0.95	-1.694	48
6	1317	18.362	0.132	Female	Yes	455	Catholic	0.95	-1.694	48

```
      ses.m  
1 0.3453333  
2 0.3453333  
3 0.3453333  
4 0.3453333  
5 0.3453333  
6 0.3453333
```

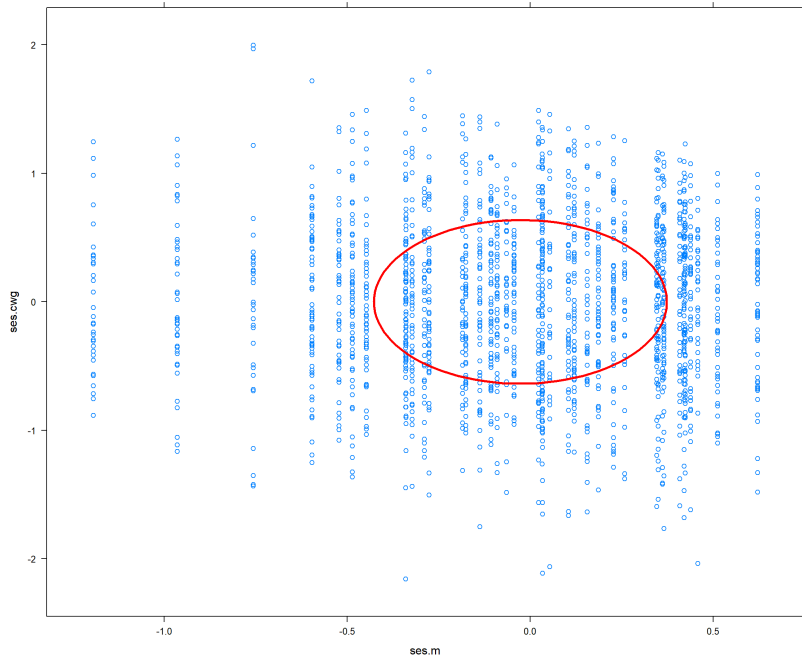
ses centered within school (CWG = centered with groups) This is the deviation of a student from the mean of the sample in her school

```
dd$ses.cwg <- dd$ses - dd$ses.m
head( dd )
```

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM	n
1	1317	12.862	0.882	Female	No	455	Catholic	0.95	-1.694	48
2	1317	8.961	0.932	Female	Yes	455	Catholic	0.95	-1.694	48
3	1317	4.756	-0.158	Female	Yes	455	Catholic	0.95	-1.694	48
4	1317	21.405	0.362	Female	Yes	455	Catholic	0.95	-1.694	48
5	1317	20.748	1.372	Female	No	455	Catholic	0.95	-1.694	48
6	1317	18.362	0.132	Female	Yes	455	Catholic	0.95	-1.694	48

	ses.m	ses.cwg
1	0.3453333	0.53666667
2	0.3453333	0.58666667
3	0.3453333	-0.50333333
4	0.3453333	0.01666667
5	0.3453333	1.02666667
6	0.3453333	-0.21333333

```
layer <- latticeExtra::layer
xyplot(ses.cwg ~ ses.m, dd) + layer(panel.dell(... ,col='red',lwd=3))
```



Note that ses.cwg and ses.m are uncorrelated

ses heterogeneity as measured by the IQR within schools

```
dd$ses.iqr <- capply( dd$ses, dd$school,  
                      function ( x ) {  
                        qs <- quantile(x, c(25,75)/100)  
                        qs[2] - qs[1]  
                      }  
)
```

Note: There is an “IQR” function to compute the inter-quartile range but this does it from scratch to illustrate the use of an ‘anonymous’ function, i.e. a function defined on the fly.

We could have done:

```
dd$ses.iqr <- capply(dd$ses, dd$school, IQR, na.rm = FALSE)
```

but we could modify the anonymous function it to

```
dd$ses.trange <- capply( dd$ses, dd$school, # where 'trange' stands for  
                                                                # 'truncated range'  
                      function ( x ) {  
                        qs <- quantile(x, c(10,90)/100, na.rm = FALSE)  
                        qs[2] - qs[1]  
                      }
```

}

)

consider variations like

```
dd$ses.sd <- capply( dd$ses, dd$school, sd, na.rm = FALSE)
```

```
some( dd )
```

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM	n
663	4292	12.956	0.852	Male	Yes	1328	Catholic	0.76	-0.674	65
677	4292	7.128	-1.148	Male	Yes	1328	Catholic	0.76	-0.674	65
763	4530	6.295	-0.738	Female	Yes	435	Catholic	0.60	-0.245	63
943	5640	4.037	-0.818	Female	No	1152	Public	0.41	0.256	57
1001	5650	8.507	1.262	Female	No	720	Catholic	0.60	-0.070	45
1107	5761	14.872	-0.278	Female	Yes	215	Catholic	0.63	-0.892	52
1121	5762	10.810	-1.498	Male	Yes	1826	Public	0.24	0.364	37
1163	6074	15.573	-0.438	Female	No	2051	Catholic	0.32	-1.018	56
1820	8707	18.470	1.512	Female	Yes	1133	Public	0.48	1.542	48
1905	9550	22.802	0.302	Male	No	1532	Public	0.45	0.791	29
	ses.m	ses.cwg	ses.iqr	ses.trange	ses.sd					
663	-0.48615385	1.3381538	0.8600	1.708	0.6511382					
677	-0.48615385	-0.6618462	0.8600	1.708	0.6511382					

763	-0.59688889	-0.1411111	0.8850	1.544	0.6210062
943	-0.17659649	-0.6414035	0.8500	1.492	0.5830261
1001	0.02244444	1.2395556	1.0600	2.114	0.7777414
1107	-0.32300000	0.0450000	0.8000	1.917	0.7122389
1121	-1.19394595	-0.3040541	0.6500	1.272	0.5154149
1163	-0.27675000	-0.1612500	0.8175	1.440	0.6271235
1820	0.15512500	1.3568750	1.1350	2.169	0.8042577
1905	0.05303448	0.2489655	1.0200	1.824	0.7847035

head(dd)

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM	n
1	1317	12.862	0.882	Female	No	455	Catholic	0.95	-1.694	48
2	1317	8.961	0.932	Female	Yes	455	Catholic	0.95	-1.694	48
3	1317	4.756	-0.158	Female	Yes	455	Catholic	0.95	-1.694	48
4	1317	21.405	0.362	Female	Yes	455	Catholic	0.95	-1.694	48
5	1317	20.748	1.372	Female	No	455	Catholic	0.95	-1.694	48
6	1317	18.362	0.132	Female	Yes	455	Catholic	0.95	-1.694	48

	ses.m	ses.cwg	ses.iqr	ses.trange	ses.sd
1	0.3453333	0.53666667	0.7825	1.166	0.5561583
2	0.3453333	0.58666667	0.7825	1.166	0.5561583
3	0.3453333	-0.50333333	0.7825	1.166	0.5561583

4	0.3453333	0.01666667	0.7825	1.166	0.5561583
5	0.3453333	1.02666667	0.7825	1.166	0.5561583
6	0.3453333	-0.21333333	0.7825	1.166	0.5561583

Transformations of Level 1 variables within groups —

`capply` will also create a Level 1 variable if the FUN returns a vector of length greater than 1. The vector gets recycled to match the length of the argument, i.e. the number of the rows corresponding to a particular id.

The following example shows the calculation of ses rank within schools:

```
dd$ses.rank <- capply( dd$ses, dd$school, rank , na.last = 'keep' )
some( dd )
```

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM	n
12	1317	23.355	0.502	Female	No	455	Catholic	0.95	-1.694	48
62	1906	21.122	0.382	Female	No	400	Catholic	0.87	-0.939	53
722	4511	20.566	-0.258	Female	No	1068	Catholic	0.52	-1.872	58
946	5640	17.085	0.012	Female	No	1152	Public	0.41	0.256	57
1027	5720	15.896	-0.078	Male	No	381	Catholic	0.65	-0.352	53
1150	5762	2.617	-1.368	Male	No	1826	Public	0.24	0.364	37

1216	6484	15.296	1.122	Male	No	726	Public	0.19	0.218	35
1394	7342	23.271	-0.748	Male	No	1220	Catholic	0.46	0.380	58
1479	7345	5.926	-1.048	Female	No	978	Public	0.64	0.336	56
1827	8854	-2.832	-0.408	Male	No	745	Public	0.18	-0.228	32

	ses.m	ses.cwg	ses.iqr	ses.trange	ses.sd	ses.rank
12	0.34533333	0.1566667	0.7825	1.166	0.5561583	30.0
62	0.51162264	-0.1296226	1.0100	1.586	0.6135833	24.5
722	-0.10713793	-0.1508621	0.8875	1.476	0.5813363	26.0
946	-0.17659649	0.1885965	0.8500	1.492	0.5830261	37.5
1027	0.03256604	-0.1105660	1.0100	1.730	0.6641693	23.0
1150	-1.19394595	-0.1740541	0.6500	1.272	0.5154149	15.0
1216	-0.18514286	1.3071429	0.7400	1.932	0.6958345	33.0
1394	-0.44782759	-0.3001724	0.7125	1.323	0.5648459	19.0
1479	0.03325000	-1.0812500	1.3550	1.930	0.8257296	6.0
1827	-0.75675000	0.3487500	0.8125	1.732	0.8036439	26.0

```
head( dd )
```

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM	n
1	1317	12.862	0.882	Female	No	455	Catholic	0.95	-1.694	48
2	1317	8.961	0.932	Female	Yes	455	Catholic	0.95	-1.694	48
3	1317	4.756	-0.158	Female	Yes	455	Catholic	0.95	-1.694	48

4	1317	21.405	0.362	Female	Yes	455	Catholic	0.95	-1.694	48
5	1317	20.748	1.372	Female	No	455	Catholic	0.95	-1.694	48
6	1317	18.362	0.132	Female	Yes	455	Catholic	0.95	-1.694	48
		ses.m	ses.cwg	ses.iqr	ses.trange		ses.sd	ses.rank		
1	0.3453333	0.53666667	0.7825		1.166	0.5561583		39		
2	0.3453333	0.58666667	0.7825		1.166	0.5561583		40		
3	0.3453333	-0.50333333	0.7825		1.166	0.5561583		6		
4	0.3453333	0.01666667	0.7825		1.166	0.5561583		25		
5	0.3453333	1.02666667	0.7825		1.166	0.5561583		47		
6	0.3453333	-0.21333333	0.7825		1.166	0.5561583		20		

Using the rank instead of the raw value could be useful for highly skewed variables where you don't want extreme values to have too much influence

The following example shows the use of 'capply' to compute a value that depends on more than one variable in the data frame. If the first argument is a data frame, then 'capply' splits the data frame into data frames with just the rows belonging to each id. Using the 'with' function on these data frames allows you to write an expression as the fourth argument which becomes the second argument of with.

ses discrepancy of Minority in school (requires more than one variable)

```
dd$minority.diff <- capply( dd, ~ school, with,
```

```
mean( ses[ Minority == "Yes" ] , na.rm = T) -
      mean( ses[ Minority == "No" ] , na.rm = T) )
```

Beware: this way of using ‘capply’ can be slow with very large files such as the NPHS long file.

```
some( dd )
```

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM	n	
95	1906	-1.603	0.282	Male	Yes	400	Catholic	0.87	-0.939	53	
141	2208	6.078	0.772	Female	No	1061	Catholic	0.68	-0.864	60	
200	2458	20.689	0.502	Female	No	545	Catholic	0.89	-1.484	57	
278	2629	13.913	0.352	Male	No	1314	Catholic	0.81	-0.613	57	
281	2629	19.111	-0.768	Male	No	1314	Catholic	0.81	-0.613	57	
1358	7232	17.001	0.332	Female	No	1154	Public	0.20	0.975	52	
1469	7345	16.652	0.292	Male	No	978	Public	0.64	0.336	56	
1509	7688	20.928	0.222	Male	No	1410	Catholic	0.65	-0.575	54	
1556	7688	19.567	-1.278	Male	No	1410	Catholic	0.65	-0.575	54	
1754	8627	16.660	-0.648	Male	No	2452	Public	0.25	0.742	53	
	ses.m		ses.cwg		ses.iqr		ses.trange		ses.sd		ses.rank
95	0.51162264		-0.22962264		1.0100		1.586		0.6135833		21.0
141	0.42316667		0.34883333		0.9825		1.505		0.5981188		38.5
200	0.22778947		0.27421053		0.7900		1.948		0.6584097		41.0
278	-0.13764912		0.48964912		1.0600		1.728		0.7063209		43.0

281	-0.13764912	-0.63035088	1.0600	1.728	0.7063209	11.5
1358	-0.09011538	0.42211538	0.7975	1.327	0.5743482	40.0
1469	0.03325000	0.25875000	1.3550	1.930	0.8257296	29.0
1509	0.18588889	0.03611111	0.8275	1.378	0.5644347	28.0
1556	0.18588889	-1.46388889	0.8275	1.378	0.5644347	1.0
1754	0.10483019	-0.75283019	0.8400	1.788	0.7077276	7.0

minority.diff

95	-0.3661667
141	NaN
200	-0.5820390
278	0.7141667
281	0.7141667
1358	-0.0107971
1469	-1.0707154
1509	-0.1387347
1556	-0.1387347
1754	-0.7930556

head(dd)

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM	n
1	1317	12.862	0.882	Female	No	455	Catholic	0.95	-1.694	48

2	1317	8.961	0.932	Female	Yes	455	Catholic	0.95	-1.694	48
3	1317	4.756	-0.158	Female	Yes	455	Catholic	0.95	-1.694	48
4	1317	21.405	0.362	Female	Yes	455	Catholic	0.95	-1.694	48
5	1317	20.748	1.372	Female	No	455	Catholic	0.95	-1.694	48
6	1317	18.362	0.132	Female	Yes	455	Catholic	0.95	-1.694	48

	ses.m	ses.cwg	ses.iqr	ses.trange	ses.sd	ses.rank
1	0.3453333	0.53666667	0.7825	1.166	0.5561583	39
2	0.3453333	0.58666667	0.7825	1.166	0.5561583	40
3	0.3453333	-0.50333333	0.7825	1.166	0.5561583	6
4	0.3453333	0.01666667	0.7825	1.166	0.5561583	25
5	0.3453333	1.02666667	0.7825	1.166	0.5561583	47
6	0.3453333	-0.21333333	0.7825	1.166	0.5561583	20

	minority.diff
1	-0.2676044
2	-0.2676044
3	-0.2676044
4	-0.2676044
5	-0.2676044
6	-0.2676044

Looking at data in 3D —

Visualizing the school level data again

```
if(interactive) {  
  up( dd, ~ school) %>% xqplot  
  up( dd, ~ school) %>% spm  
}
```

QUESTIONS:

1. Would sample size be a reasonable proxy for school size if we did not have school size?
2. Can you think of other ways of summarizing or visualizing the data: what do you see?

1d - 2d -> 3d

We can see 3 continuous variable + 1 categorical variable (really 4d)

```
if(interactive){  
  library( p3d )  
  Init3d(family = 'serif', cex = 1.2)  
  dda <- up(dd, ~ school, ~ mathach + ses + Sex)  
  head(dda)  
  Plot3d ( mathach ~ ses + Sex_Female | Sector, dda)
```

```
fg()
}
```

Note gender composition of Catholic schools

```
# just for fun
if(interactive) {
  fit <- lm( mathach ~ ses * Sex_Female * Sector +
             I(ses*(Sector == 'Catholic')*Sex_Female^2), dda)
  summary(fit)
  Fit3d ( fit )
  Id3d(pad=2)
}
```

EXERCISE:

Try other possibilities. What do you find?

Looking at Level 1 and Level 2 data using Lattice graphics —

Preparing for multilevel modelling:

Visualizing data at Level 1 and Level 2

From the mixed model to the hierarchical models:

With software like HLM, MLWin, you need to use separate data sets for the different levels.

With nlme and SAS PROC MIXED, we use the combined long data set with all the variables together.

We first need to separate the Levels of the model.

1. Level 1 model:

```
mathach ~ 1 + ses | school
```

2. Level 2 model:

- Using coefficients of Level 1 Model:

- `B.0 ~ 1 + Sector`

- `B.ses ~ 1 + Sector`

- Combining with Level 1:

- `~ 1 * ( 1 + Sector ) + ses * ( 1 + Sector )`

3. Mixed model: Simplify combined model

```
mathach ~ 1 + ses + Sector + ses:Sector
```

OR (using R's rules for generating marginal effects)

```
mathach ~ ses * Sector
```

4. Random effects model:

```
~ 1 + ses | id
```

– all terms usually contained in Level 1 model

Often, a simpler model is used, e.g. ‘~ 1 | id’

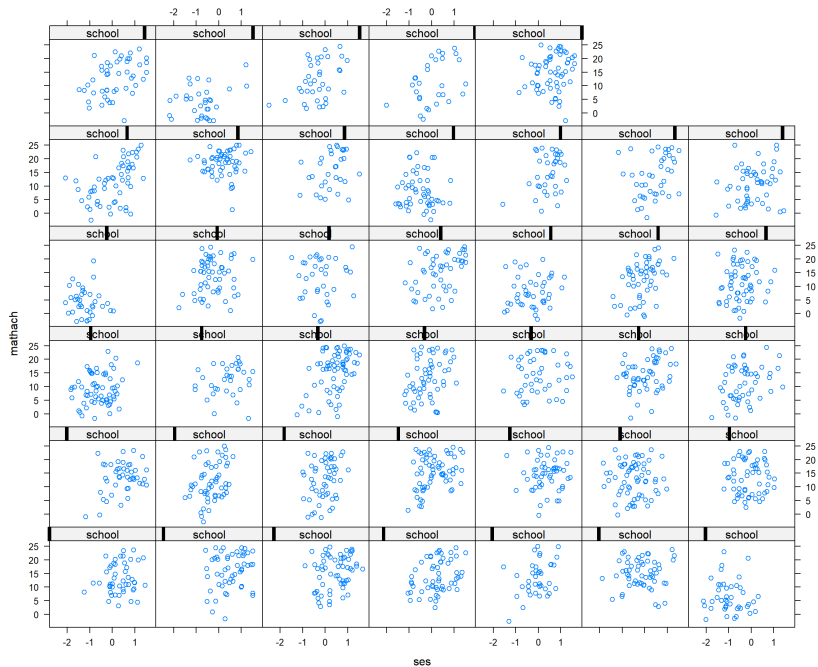
5. Compositional variable:

Mean ses | school i.e `cvar( ses , school)`

Mixed model with compositional variable:

```
mathach ~ 1 + ses + Sector + ses:Sector + cvar( ses, school) + ...
```

```
xyplot( mathach ~ ses | school, dd )
```



```
# Make an informative label and ordering for schools
```

```
dd$id <- factor( paste( substring( dd$Sector, 1,1), dd$school, sep =','))
some( dd )
```

	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD	DISCLIM	n
147	2208	20.628	0.792	Male	No	1061	Catholic	0.68	-0.864	60
362	2658	12.860	0.952	Female	No	780	Catholic	0.79	-0.961	45
842	4868	9.368	-0.158	Male	Yes	657	Catholic	1.00	-0.219	34
886	5619	1.701	0.472	Female	No	1118	Catholic	0.77	-1.286	66
1210	6074	19.882	1.512	Female	No	2051	Catholic	0.32	-1.018	56
1484	7345	9.098	-0.358	Female	Yes	978	Public	0.64	0.336	56
1495	7345	23.134	0.472	Female	No	978	Public	0.64	0.336	56
1584	7697	11.620	0.872	Male	No	1734	Public	0.20	0.279	32
1910	9550	-1.005	-0.508	Female	Yes	1532	Public	0.45	0.791	29
1958	9586	24.292	1.002	Female	No	262	Catholic	1.00	-2.416	59
	ses.m	ses.cwg	ses.iqr	ses.trange	ses.sd	ses.rank				
147	0.42316667	0.36883333	0.9825	1.505	0.5981188	41.5				
362	0.43844444	0.51355556	0.9100	1.698	0.6402846	37.0				
842	0.36111765	-0.51911765	1.0475	1.920	0.7100080	10.0				
886	0.42033333	0.05166667	0.7550	1.405	0.5972748	31.0				
1210	-0.27675000	1.78875000	0.8175	1.440	0.6271235	56.0				

1484	0.03325000	-0.39125000	1.3550	1.930	0.8257296	18.0
1495	0.03325000	0.43875000	1.3550	1.930	0.8257296	35.5
1584	0.25825000	0.61375000	0.8625	1.438	0.6133712	29.0
1910	0.05303448	-0.56103448	1.0200	1.824	0.7847035	9.0
1958	0.62115254	0.38084746	0.8550	1.390	0.5949914	43.0

	minority.diff	id
147	NaN	C2208
362	-0.18305556	C2658
842	-0.77505263	C4868
886	NaN	C5619
1210	-0.32796296	C6074
1484	-1.07071545	P7345
1495	-1.07071545	P7345
1584	NaN	P7697
1910	-0.56239130	P9550
1958	0.08517857	C9586

ordering used for panels in xyplot:

```
dd$id <- reorder( dd$id, dd$ses)      # or
```

```
dd$id.sec <- reorder( dd$id, dd$ses + 1000* (dd$Sector == "Catholic"))
```

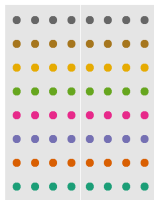
Note: If you are irritated by caps in variable names use:

```
names(dd) <- tolower( names(dd) )
```

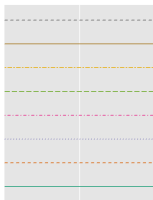
Make sure this won't create duplicates. Variable names are case sensitive in R

```
gd() # sets 8 default colours for multiple groups using colour-blind friendly co
```

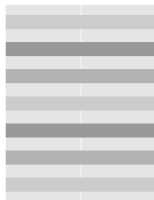
```
show.settings() # first two panels show 'superpose' colours for multiple groups
```



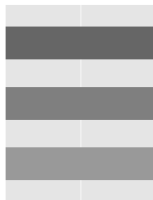
superpose.symbol



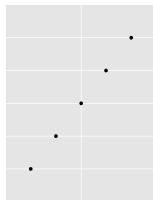
superpose.line



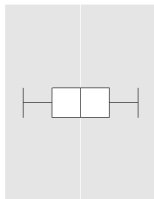
strip.background



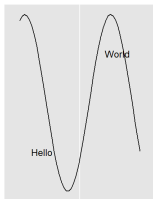
strip.shingle



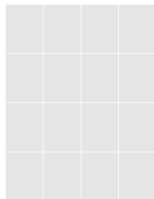
dot.[symbol, line]



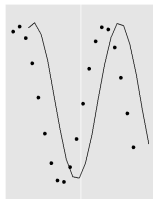
box.[dot, rectangle, umbrella]



add.[line, text]



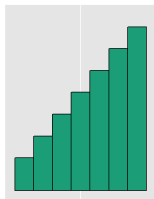
reference.line



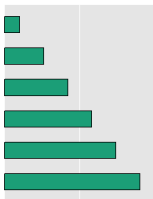
plot.[symbol, line]



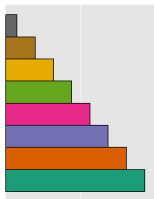
plot.shingle[plot.polygon]



histogram[plot.polygon]



barchart[plot.polygon]

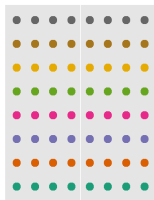


superpose.polygon

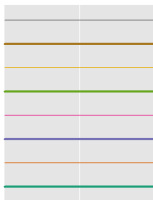


regions

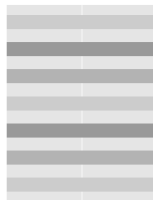
```
gd(lwd=c(3,1), lty = c(1,1)) # if arguments have length > 1, superpose options a  
show.settings() # first two panels show 'superpose' colours for multiple groups
```



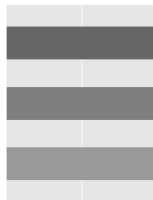
superpose.symbol



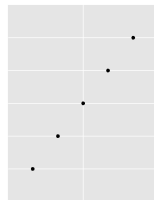
superpose.line



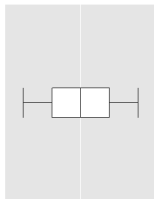
strip.background



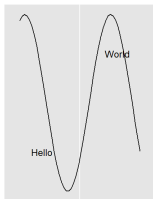
strip.shingle



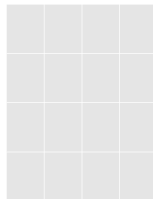
dot.[symbol, line]



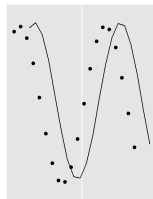
box.[dot, rectangle, umbrella]



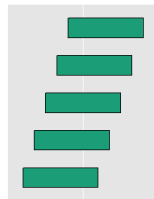
add.[line, text]



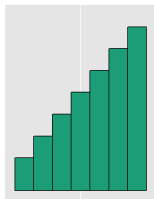
reference.line



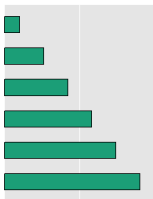
plot.[symbol, line]



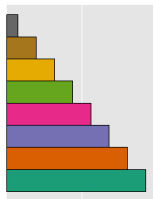
plot.shingle[plot.polygon]



histogram[plot.polygon]



barchart[plot.polygon]



superpose.polygon



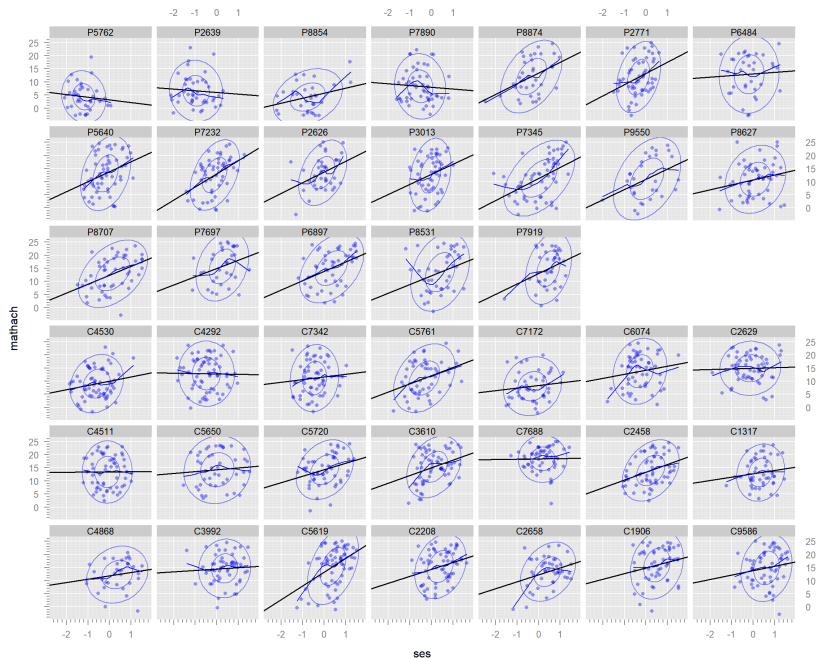
regions

```

gd(col = 'blue', pch = 16, alpha = 1)      # options for single group
xyplot( mathach ~ ses | id.sec, dd, layout = c(7,6),
  main = list('schools ordered by sector and mean ses', cex = 1, font = 1),
  skip = c(rep(FALSE, 19), TRUE, TRUE, rep(FALSE, 21) ),
  as.table = TRUE,
  panel = function( x,y ,..., type, alpha, lwd) {
    panel.xyplot( x, y, ... , alpha = .4)
    panel.lines( dell( x, y, radius = 1:2), type = 'l', alpha = .5,...)
    panel.lmline( x, y, ... , lwd = 1.5, alpha = 1)
    panel.lines( lowess( x, y),..., type = 'l')
  }
)

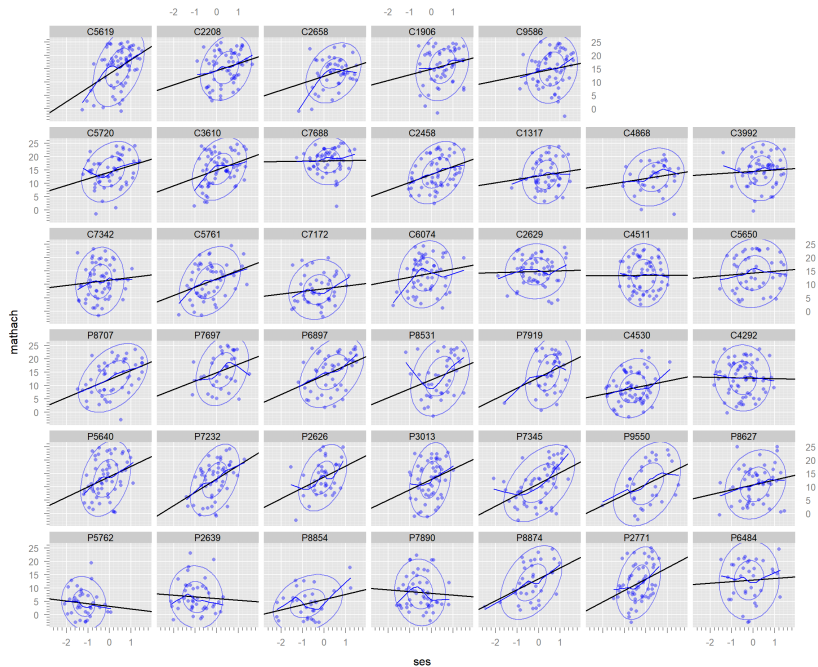
```

schools ordered by sector and mean ses



If you like a particular panel function, you can make it yours: R is easy to customize:

```
mypanel = function( x,y ,..., type, alpha, lwd) {  
  panel.xyplot( x, y, ... , alpha = .4)  
  panel.lines( dell( x, y, radius = 1:2), type = 'l', alpha = .5,...)  
  panel.lmline( x, y, ..., lwd = 1.5, alpha = 1)  
  panel.lines( lowess( x, y),..., type = 'l')  
}  
  
xyplot( mathach ~ ses | id.sec, dd,  
        panel = mypanel)
```

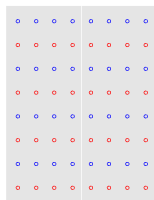


ses

Use 'groups' to have more than one group per panel

Set options for groups by specifying vectors of length > 1 , short vectors get recycled

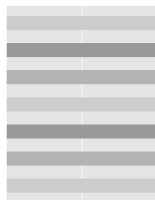
```
gd(col = c('red', 'blue'), pch = c(1, 1), cex = c(.7, .7))  
show.settings()
```



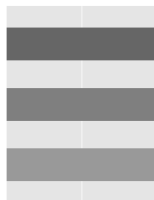
superpose.symbol



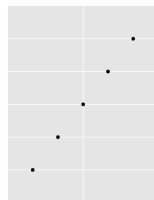
superpose.line



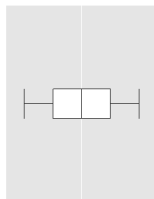
strip.background



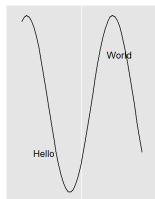
strip.shingle



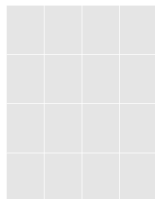
dot.[symbol, line]



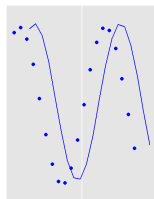
box.[dot, rectangle, umbrella]



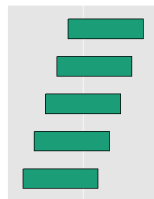
add.[line, text]



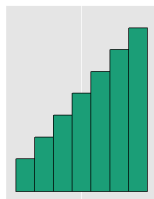
reference.line



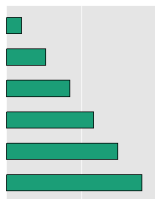
plot.[symbol, line]



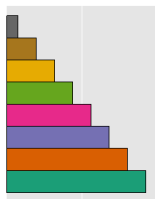
plot.shingle[plot.polygon]



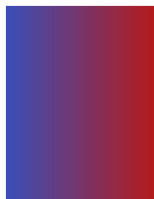
histogram[plot.polygon]



barchart[plot.polygon]



superpose.polygon

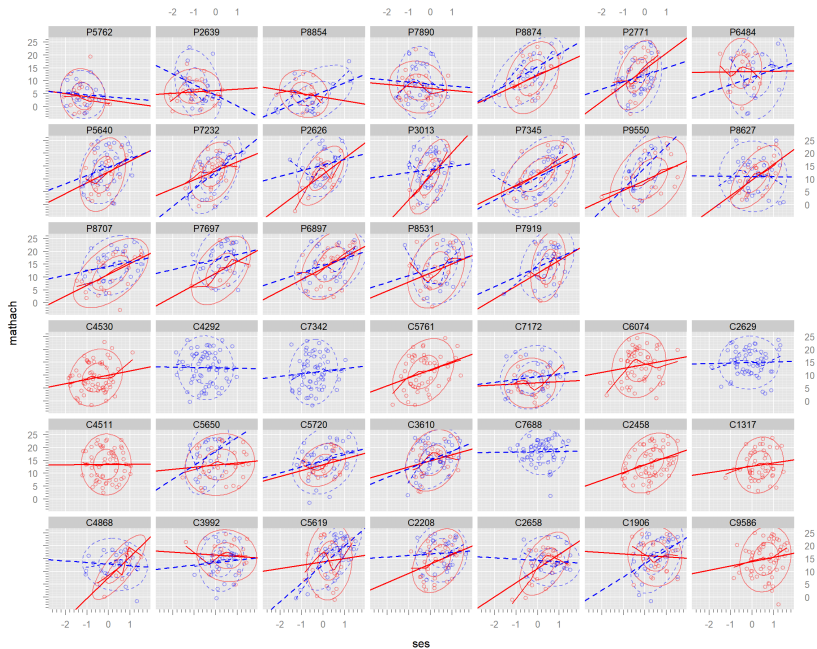


regions

```
xyplot( mathach ~ ses | id.sec, dd, layout = c(7,6),  
        groups = Sex,  
        main = list('schools ordered by sector and mean ses', cex = 1, font = 1),  
        skip = c(rep(FALSE, 19), TRUE, TRUE, rep(FALSE, 21) ),  
        as.table = TRUE,  
        panel = panel.superpose,  
        panel.groups = mypanel,  
        auto.key = list(columns = 2, lines = T)  
)
```

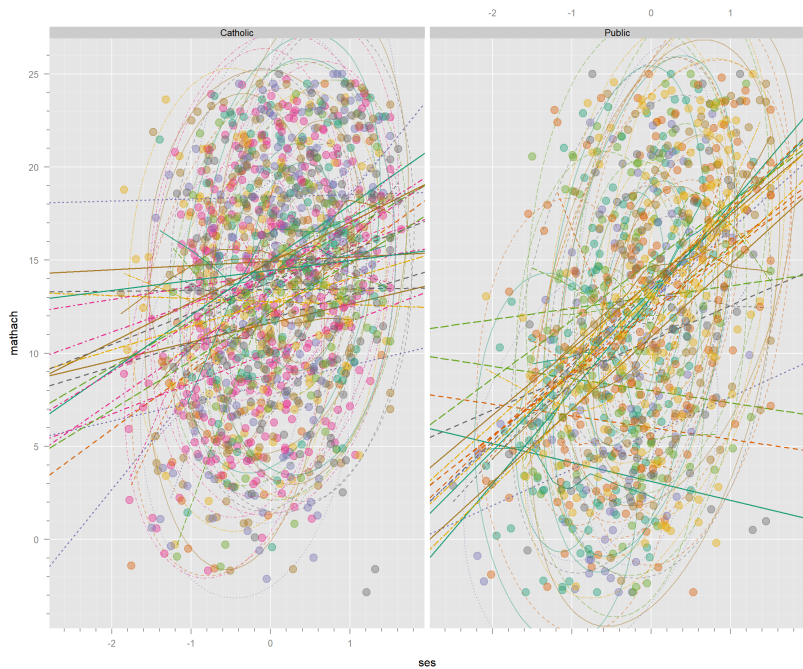
schools ordered by sector and mean ses

Female ○ — Male ○ - - -



Combine panels and groups to see data in different ways:

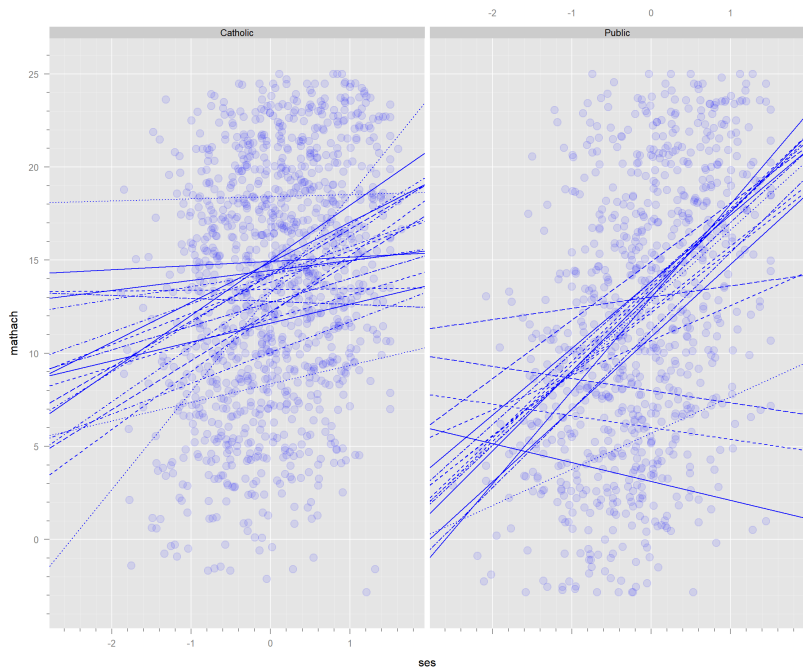
```
gd()  
xyplot( mathach ~ ses | Sector, dd,    groups = id,  
        panel = panel.superpose,  
        panel.groups = mypanel)
```



too much! Just data and fitted line:

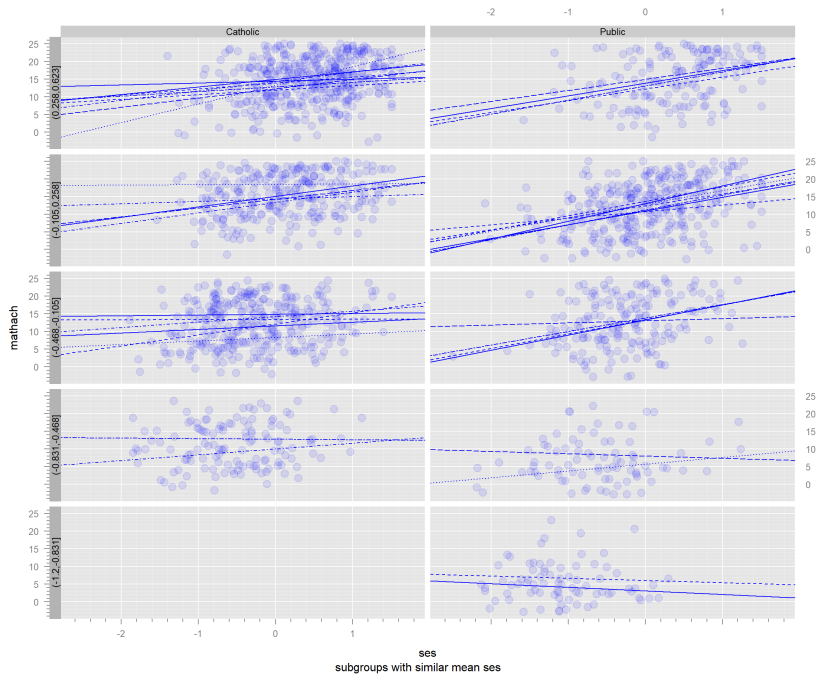
```
mypanel2 <- function( x,y ,..., type, alpha, col) {  
  panel.xyplot( x, y, ... ,alpha = alpha)  
  panel.lmline( x, y, ..., alpha = 1, superpose.col = 'black')  
}
```

```
gd(col=c('blue','blue'),alpha=c(.1,.1))  
xyplot( mathach ~ ses | Sector, dd, groups = id,  
        panel = panel.superpose,  
        panel.groups = mypanel2)
```



fitted lines generally higher and flatter in Catholic sector

```
xyplot( mathach ~ ses | Sector * cut(ses.m, 5), dd, groups = id,  
        sub = list('subgroups with similar mean ses', font=1, cex = 1),  
        panel = panel.superpose,  
        panel.groups = mypanel2) %>%  
useOuterStrips
```

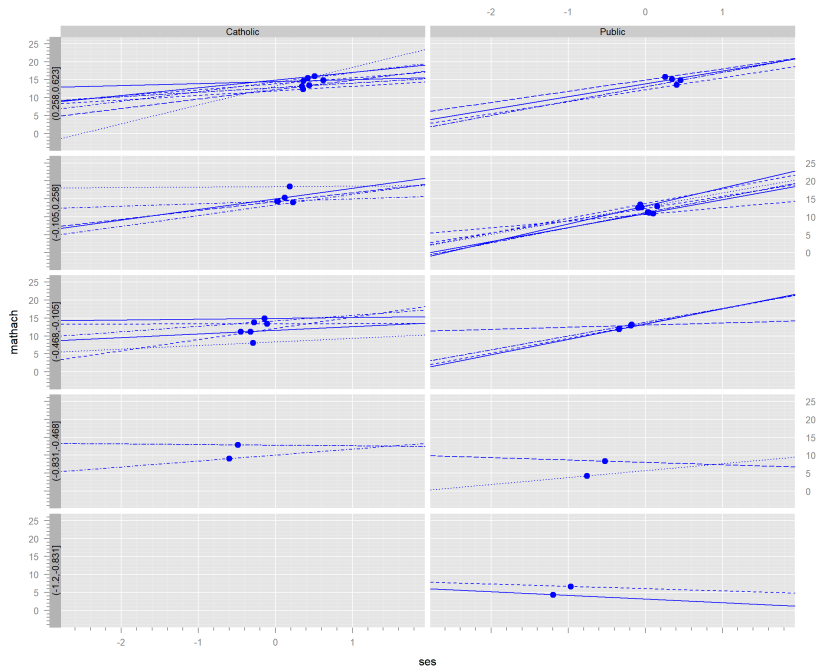


Note:

- fitted lines generally higher and flatter in Catholic sector

Center point and fitted lines

```
mypanel3 <- function( x,y ,..., type) {  
  panel.xyplot( mean(x), mean(y), ... )  
  panel.lmline( x, y, ...)  
}  
gd(col=c('blue','blue'), alpha = 1, cex = 1)  
xyplot( mathach ~ ses | Sector * cut( ses.m, 5), dd, groups = id,  
        panel = panel.superpose,  
        panel.groups = mypanel3) %>%  
useOuterStrips
```



Visualizing fitted lines in beta space —

Fit an OLS model in each school

```
# fit.list <- lmList( mathach ~ ses | id, dd)
```

```
# may produce an error due to missing values in some variables in 'dd' that  
# 'lmList' does not use. To work around this problem, you can use:
```

```
fit.list <- lmList( mathach ~ ses | id, dd[,c('mathach','ses','id')])  
fit.list
```

Call:

```
Model: mathach ~ ses | id  
Data: dd[, c("mathach", "ses", "id")]
```

Coefficients:

	(Intercept)	ses
P5762	3.114085	-1.01409923
P2639	6.007785	-0.63010463
P8854	5.707002	1.93884461
C4530	10.039029	1.64742597

P7890	7.998220	-0.65596791
C4292	12.786274	-0.16060800
C7342	11.619820	1.01245787
P8874	13.450957	4.09630389
P2771	13.292965	4.26818800
C5761	12.141945	3.10801055
C7172	8.355037	0.99448053
C6074	14.202264	1.52908771
P6484	13.024537	0.60567730
P5640	13.836071	3.82774230
C2629	14.938378	0.22234915
C4511	13.413589	0.04251038
P7232	12.993356	5.00160032
P2626	13.662437	4.09967961
P3013	12.780651	3.83979913
C5650	14.258257	0.68061888
C5720	14.201984	2.46630669
P7345	11.198507	4.21192377
P9550	10.882731	3.89193777
P8627	10.687731	1.86955959
C3610	14.999882	2.95584910

P8707	12.357826	3.39153230
C7688	18.400688	0.11634493
C2458	13.312180	2.95669443
P7697	14.911853	3.13621972
C1317	12.737763	1.27391282
P6897	13.846070	3.58048769
C4868	11.845609	1.28647122
C3992	14.448670	0.53787533
P8531	12.174522	3.31822792
C5619	13.206326	5.25753341
C2208	14.288928	2.63664069
C2658	12.243090	2.62990138
P7919	13.024135	3.98937031
C1906	14.885481	2.14550546
C9586	13.825077	1.67208118

Degrees of freedom: 1977 total; 1897 residual

Residual standard error: 6.106947

`levels( dd$id )`

[1] "P5762" "P2639" "P8854" "C4530" "P7890" "C4292" "C7342" "P8874"

```

[9] "P2771" "C5761" "C7172" "C6074" "P6484" "P5640" "C2629" "C4511"
[17] "P7232" "P2626" "P3013" "C5650" "C5720" "P7345" "P9550" "P8627"
[25] "C3610" "P8707" "C7688" "C2458" "P7697" "C1317" "P6897" "C4868"
[33] "C3992" "P8531" "C5619" "C2208" "C2658" "P7919" "C1906" "C9586"

```

```

beta <- expand.grid( id = levels( dd$id ))
beta <- cbind( beta, coef( fit.list))
some ( beta )

```

	id (Intercept)	ses
C7342 C7342	11.619820	1.0124579
P8874 P8874	13.450957	4.0963039
C5761 C5761	12.141945	3.1080106
C7172 C7172	8.355037	0.9944805
C2629 C2629	14.938378	0.2223492
C5720 C5720	14.201984	2.4663067
C3610 C3610	14.999882	2.9558491
C2458 C2458	13.312180	2.9566944
C1317 C1317	12.737763	1.2739128
C2658 C2658	12.243090	2.6299014

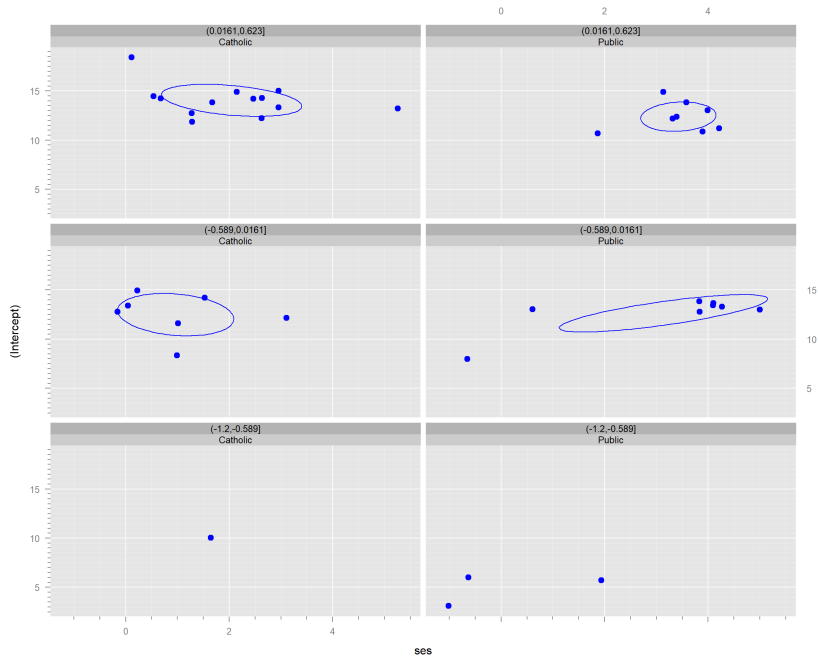
```

beta <- merge( beta, up(dd, ~ id))

```

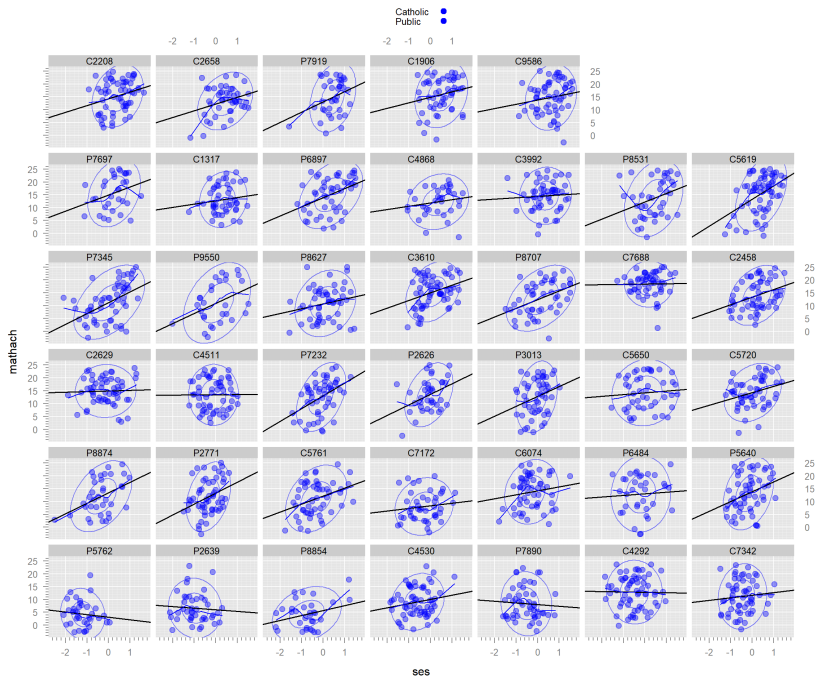
```
xyplot( '(Intercept)' ~ ses | Sector * cut( ses.m, 3), beta, groups = Sector,  
  panel = function(x, y, ..., type){  
    panel.xyplot( x,y , ...)  
    if ( length(x) > 3)panel.lines( dell( x, y), ..., type = 'l')  
  },  
  main = 'beta space'  
)
```

beta space



Looking at between group effect —

```
xyplot( mathach ~ ses | id, dd, layout = c(7,6), groups = Sector,  
        panel = mypanel, auto.key = T)
```



ses

```
dda <- up(dd, ~ id, all = T) # means of numerical variables, modes of factors
some(dda)
```

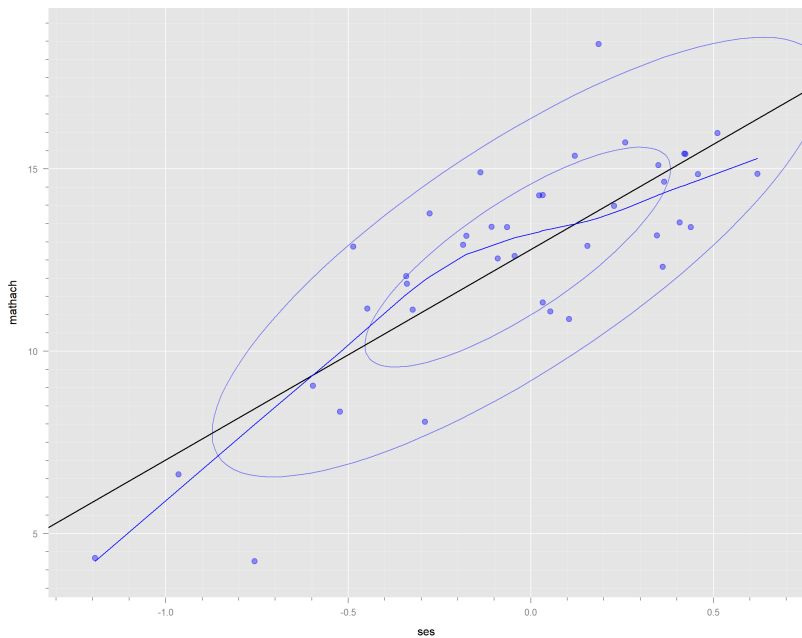
	school	mathach	ses	Sex	Minority	Size	Sector	PRACAD
C7342	7342	11.166414	-0.44782759	Male	No	1220	Catholic	0.46
C5761	5761	11.138058	-0.32300000	Female	No	215	Catholic	0.63
C7172	7172	8.066818	-0.28981818	Female	Yes	280	Catholic	0.05
P2626	2626	13.396605	-0.06484211	Male	No	2142	Public	0.40
P3013	3013	12.610830	-0.04422642	Male	No	760	Public	0.56
P8627	8627	10.883717	0.10483019	Male	No	2452	Public	0.25
P8707	8707	12.883938	0.15512500	Female	No	1133	Public	0.48
C4868	4868	12.310176	0.36111765	Male	No	657	Catholic	1.00
P8531	8531	13.528683	0.40809756	Female	No	2190	Public	0.58
P7919	7919	14.849973	0.45767568	Male	No	1451	Public	0.50

	DISCLIM	n	ses.m	ses.cwg	ses.iqr	ses.trange	ses.sd
C7342	0.380	58	-0.44782759	-4.974128e-19	0.7125	1.323	0.5648459
C5761	-0.892	52	-0.32300000	-5.358461e-18	0.8000	1.917	0.7122389
C7172	1.013	44	-0.28981818	-3.027573e-17	1.0275	1.686	0.6764417
P2626	0.142	38	-0.06484211	6.211337e-18	0.5975	1.384	0.5601067
P3013	-0.213	53	-0.04422642	1.112228e-17	0.7300	1.168	0.4799328
P8627	0.742	53	0.10483019	-6.563514e-18	0.8400	1.788	0.7077276
P8707	1.542	48	0.15512500	8.673617e-19	1.1350	2.169	0.8042577

C4868	-0.219	34	0.36111765	-6.555484e-18	1.0475	1.920	0.7100080
P8531	0.132	41	0.40809756	-1.152428e-17	1.0600	1.650	0.6829747
P7919	-0.402	37	0.45767568	-1.351883e-17	0.6700	1.070	0.5367005
	ses.rank	minority.diff	id	id.sec			
C7342	29.5	0.1027413	C7342	C7342			
C5761	26.5	0.1197576	C5761	C5761			
C7172	22.5	0.3915447	C7172	C7172			
P2626	19.5	NaN	P2626	P2626			
P3013	27.0	-0.6765385	P3013	P3013			
P8627	27.0	-0.7930556	P8627	P8627			
P8707	24.5	-0.5873333	P8707	P8707			
C4868	17.5	-0.7750526	C4868	C4868			
P8531	21.0	-0.4734848	P8531	P8531			
P7919	19.0	0.3127778	P7919	P7919			

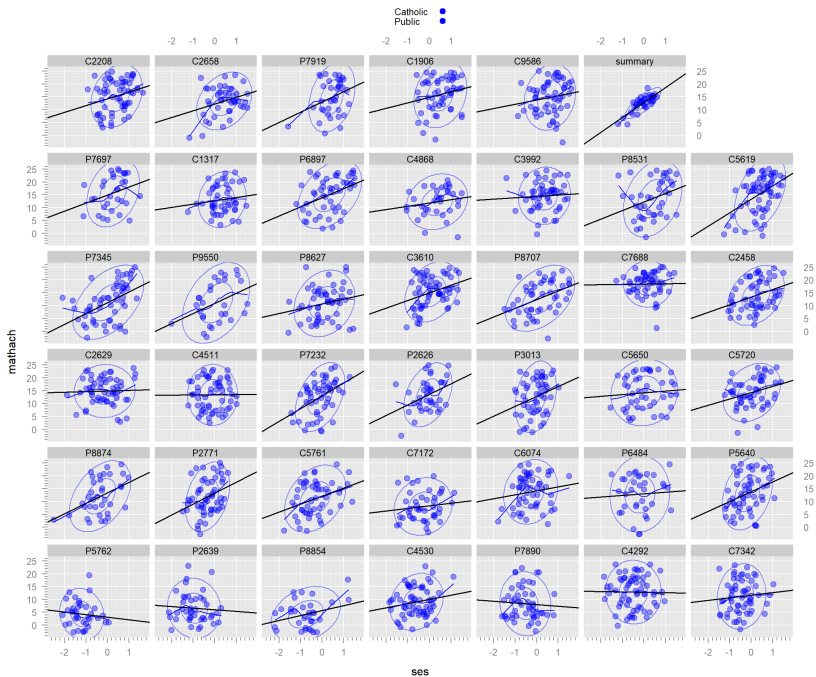
```
xyplot( mathach ~ ses, dda , groups = Sector,
        panel = mypanel, auto.key = T)
```

Catholic
Public



Seeing both together:

```
dda$id <- 'summary'  
gd(pch='.', cex = 2)  
xyplot( mathach ~ ses | id, rbind(dd, dda), layout = c(7, 6),  
        groups = Sector,  
        panel = mypanel, auto.key = T)
```



ses

EXERCISE:

Try groups = Sex or Minority and try 'id.sec

Do these plots reveal anything useful?

Fitting a mixed model —

Question to investigate:

mathach ~ ses in different Sectors

Level 1 model:

mathach ~ 1 + ses

Level 1 + 2:

mathach ~ 1 + ses + Sector + Sector:sese

Full random model:

~ 1 + ses | id

dd\$id <- factor(dd\$school)

fit <- lme(mathach ~ ses * Sector, dd, random = ~ 1 + ses | id)

If this fails to converge, increase the number of iteration

Convergence problems —

```
fit <- lme( mathach ~ ses * Sector, dd, random = ~ 1 + ses | id,  
           control = list(msMaxIter=200, msVerbose=T))
```

0:	11105.191:	1.22722	2.27855	-3.36583
1:	11105.172:	1.23706	3.08949	-3.39049
2:	11105.171:	1.19748	3.08072	-3.39188
3:	11105.161:	1.21737	3.07678	-3.39235
4:	11105.096:	1.22151	2.70896	-3.41493
5:	11105.089:	1.22011	2.64468	-3.43555
6:	11105.085:	1.22389	2.55034	-3.50479
7:	11105.083:	1.22385	2.57106	-3.54636
8:	11105.076:	1.22534	2.60016	-3.72983
9:	11105.057:	1.23381	2.66214	-4.47027
10:	11105.033:	1.25234	2.72319	-5.87758
11:	11105.019:	1.27696	2.75593	-7.28573
12:	11105.013:	1.30246	2.78019	-8.50888
13:	11105.011:	1.31965	2.79844	-9.36213

14:	11105.011:	1.33116	2.81675	-9.93866
15:	11105.011:	1.33714	2.82776	-10.3398
16:	11105.011:	1.33993	2.83733	-10.5086
17:	11105.011:	1.33984	2.84079	-10.5414
18:	11105.011:	1.33908	2.84283	-10.5393
19:	11105.011:	1.33903	2.84278	-10.5371

```
print(fit)
```

Linear mixed-effects model fit by REML

Data: dd

Log-restricted-likelihood: -6419.695

Fixed: mathach ~ ses * Sector

(Intercept)	ses	SectorPublic	ses:SectorPublic
13.479564	1.805347	-1.599373	1.388797

Random effects:

Formula: ~1 + ses | id

Structure: General positive-definite, Log-Cholesky parametrization

StdDev	Corr
--------	------

(Intercept)	1.8816109	(Intr)
-------------	-----------	--------

ses	0.3564559	0.523
-----	-----------	-------

Residual 6.1180179

Number of Observations: 1977

Number of Groups: 40

Things we can do with a model

`summary( fit )`

Linear mixed-effects model fit by REML

Data: dd

	AIC	BIC	logLik
	12855.39	12900.09	-6419.695

Random effects:

Formula: ~1 + ses | id

Structure: General positive-definite, Log-Cholesky parametrization

	StdDev	Corr
(Intercept)	1.8816109	(Intr)
ses	0.3564559	0.523
Residual	6.1180179	

Fixed effects: mathach ~ ses * Sector

	Value	Std.Error	DF	t-value	p-value
(Intercept)	13.479564	0.4502225	1935	29.939782	0.0000
ses	1.805347	0.2905759	1935	6.212995	0.0000
SectorPublic	-1.599373	0.6628232	38	-2.412971	0.0208
ses:SectorPublic	1.388797	0.4359721	1935	3.185518	0.0015

Correlation:

	(Intr)	ses	SctrPb
ses	0.094		
SectorPublic	-0.679	-0.064	
ses:SectorPublic	-0.063	-0.667	0.163

Standardized Within-Group Residuals:

Min	Q1	Med	Q3	Max
-3.06175080	-0.74559046	0.03508812	0.78804832	2.73786882

Number of Observations: 1977

Number of Groups: 40

Sequential tests

anova(fit)

numDF	denDF	F-value	p-value
-------	-------	---------	---------

(Intercept)	1	1935	1376.8723	<.0001
ses	1	1935	129.5570	<.0001
Sector	1	38	8.8271	0.0051
ses:Sector	1	1935	10.1475	0.0015

Type II tests

- Each term added last EXCLUDING interactions that include it
- Test that respect the principle of marginality

Anova(fit)

Analysis of Deviance Table (Type II tests)

Response: mathach

	Chisq	Df	Pr(>Chisq)	
ses	125.0347	1	< 2.2e-16	***
Sector	8.8271	1	0.002968	**
ses:Sector	10.1475	1	0.001445	**

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Overall test for a variable including all its interactions

- Quick and rough: using spida2::wald
- Conservative denominator DFs

```
wald(fit, 'ses')           # overall test for 'ses'
```

	numDF	denDF	F.value	p.value
ses	2	1935	67.59113	<.00001

Coefficients	Estimate	Std.Error	DF	t-value	p-value	Lower	Upper
ses	1.805347	0.290576	1935	6.212995	<.00001	1.235472	2.375222
ses:SectorPublic	1.388797	0.435972	1935	3.185518	0.00147	0.533773	2.243822

Coefficients	Upper
ses	2.375222
ses:SectorPublic	2.243822

```
wald(fit, 'Sector')       # overall test for 'Sector'
```

	numDF	denDF	F.value	p.value
Sector	2	38	9.487323	0.00045

Coefficients	Estimate	Std.Error	DF	t-value	p-value	Lower	Upper
SectorPublic	-1.599373	0.662823	38	-2.412971	0.02075	-2.941188	0.533773

```
ses:SectorPublic  1.388797  0.435972 1935  3.185518 0.00147  0.533773
```

```
Coefficients      Upper 0.95  
SectorPublic      -0.257558  
ses:SectorPublic   2.243822
```

Using car::lht

```
fixef(fit)
```

(Intercept)	ses	SectorPublic	ses:SectorPublic
13.479564	1.805347	-1.599373	1.388797

```
coefs <- names(fixef(fit))  
coefs
```

```
[1] "(Intercept)"      "ses"              "SectorPublic"  
[4] "ses:SectorPublic"
```

```
lht(fit, coefs[c(2,4)])
```

Linear hypothesis test

Hypothesis:
ses = 0

```
ses:SectorPublic = 0
```

```
Model 1: restricted model
```

```
Model 2: mathach ~ ses * Sector
```

```
      Df  Chisq Pr(>Chisq)
```

```
1
```

```
2  2 135.18  < 2.2e-16 ***
```

```
---
```

```
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
lht(fit, coefs[c(3,4)])
```

```
Linear hypothesis test
```

```
Hypothesis:
```

```
SectorPublic = 0
```

```
ses:SectorPublic = 0
```

```
Model 1: restricted model
```

```
Model 2: mathach ~ ses * Sector
```

```

      Df   Chisq Pr(>Chisq)
1
2  2 18.975   7.581e-05 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

Note that this model tacitly assumes that the between-school effect of ses is the same as the within-school effect of ses

In passing: handling NAs —

This is a big topic but note that deleting rows with NAs does not delete entire clusters, only units within clusters. This is appropriate under a wider range of assumptions than deleting clusters – as one would have to do with a repeated measures analysis. More on this in Lab 4.R

Syntax

```

fit <- lme( mathach ~ ses * Sector, dd, random = ~ 1 + ses | id,
            na.action = na.exclude,
            control = list(msMaxIter=200, msVerbose=T))

```

0:	11105.191:	1.22722	2.27855	-3.36583
1:	11105.172:	1.23706	3.08949	-3.39049

2:	11105.171:	1.19748	3.08072	-3.39188
3:	11105.161:	1.21737	3.07678	-3.39235
4:	11105.096:	1.22151	2.70896	-3.41493
5:	11105.089:	1.22011	2.64468	-3.43555
6:	11105.085:	1.22389	2.55034	-3.50479
7:	11105.083:	1.22385	2.57106	-3.54636
8:	11105.076:	1.22534	2.60016	-3.72983
9:	11105.057:	1.23381	2.66214	-4.47027
10:	11105.033:	1.25234	2.72319	-5.87758
11:	11105.019:	1.27696	2.75593	-7.28573
12:	11105.013:	1.30246	2.78019	-8.50888
13:	11105.011:	1.31965	2.79844	-9.36213
14:	11105.011:	1.33116	2.81675	-9.93866
15:	11105.011:	1.33714	2.82776	-10.3398
16:	11105.011:	1.33993	2.83733	-10.5086
17:	11105.011:	1.33984	2.84079	-10.5414
18:	11105.011:	1.33908	2.84283	-10.5393
19:	11105.011:	1.33903	2.84278	-10.5371

Note: generally 'na.exclude' is preferable to 'na.omit' It allows residuals and predicted values to match the original data frame, not just the observations that were not missing.

In passing: What to do if the model does not converge

We will revisit this when we need it:

Summary:

1. If Iteration limit reached: try increase iterations.

```
?lmeControl
```

```
lmeControl
```

```
fit <- lme( mathach ~ ses * Sector, dd, random = ~ 1 + ses | id,  
          control = list( maxIter = 200, msMaxIter = 200, niterEM = 100,  
                          msVerbose = TRUE , returnObject = TRUE ))  
summary( fit )
```

2. If singular convergence, the frequent cause is that the random model is more complex than the cluster size or the variability between clusters can support. I.e. the variability from cluster to cluster that you are trying to account for with the random part of the model is actually accounted for by the ‘epsilons’, the level 1 or within-cluster variability. Consequently a variance is estimated to be 0 or the G matrix is non-singular, i.e. it’s a flat pancake.

As a rule of thumb, if the random effects model has K terms, you need, for a balanced

within cluster design, a good deal more than K clusters with at least K observations and some with more than K observations. Otherwise, the G matrix and the within-cluster variance may not be identifiable.

Also, the G matrix might be relatively flat because of scaling OR centering. For example, one dimension is so much more stretched than another.

Try:

1. Rescale variables so they have similar variability or similar variability in slopes.
2. Global center x 's in RE model (not necessarily at the mean but that's a good first try. The 'ideal' place is the point of minimum variance of the response. Plot OLS lines to get an idea.
3. Also try centering within clusters (CWC or CWG). The RE model is not equivalent but it could often be better depending on the process. Use AIC to compare.,
4. If nothing works, start with simple RE model and add components using a manual forward stepwise approach. Use 'anova' and 'simulate' to test additional components. If a model does not converge, use `..., control = (..., returnObject = TRUE, msVerbose = TRUE)` to judge whether the likelihood ratio converges even though the parameters don't. In this case, anova can give a usable p-value that can be adjusted with 'simulate'.

```
'fit.big <- lme( . . . . , control = list( maxIter = 200,  
'          msMaxIter = 200, niterEM = 100,  
'          msVerbose = TRUE , returnObject = TRUE ))'  
'anova ( fit.simple, fit.big )'  
'zsim <- simulate( fit.simple, nsim = 1000, m2 = fit.big)'  
'plot( zsim )'
```

to test whether 'fit.simple' is adequate.

Handling non-convergence is discussed in greater detail in Lab Session 3

Hausman Test: Is the between effect different from the within effect?

—

First diagnostics: Hausman Test

After fitting a model: diagnostics

- Historically: The Hausman test asks whether the mixed model correctly estimates the effect of ses
- But in fact:
 - Do we need a contextual variable? i.e.

- Is the between-school effect of ses not significantly different from the within-school effect of ses
- If so, then we should include a separate effect for the between-school effect of ses and we will be able to estimate the within school effect correctly.

Model with contextual variable for ses, i.e. each school's mean ses: Note: we can use the `cvar` function for convenience or we can create a variable in the data frame, i.e. `ses.m` above. We will use `cvar` because it's always easy to use even if the variable has not been defined.

Moreover, `cvar` works correctly with factors generating mean values of indicators for each level of the factor omitting the reference level. We will illustrate this later.

Fitting a model with a contextual mean —

```
fitc <- lme( mathach ~ (ses + cvar(ses,id))* Sector, dd,
            random = ~ 1 + ses | id )
summary( fitc )
```

Linear mixed-effects model fit by REML

Data: dd

AIC	BIC	logLik
12839.6	12895.47	-6409.801

Random effects:

Formula: ~1 + ses | id

Structure: General positive-definite, Log-Cholesky parametrization

StdDev Corr

(Intercept) 1.4561519 (Intr)

ses 0.5190639 -0.217

Residual 6.1140010

Fixed effects: mathach ~ (ses + cvar(ses, id)) * Sector

	Value	Std.Error	DF	t-value	p-value
(Intercept)	13.356053	0.3753744	1935	35.58062	0.0000
ses	1.699397	0.3095543	1935	5.48982	0.0000
cvar(ses, id)	2.407785	1.0629298	36	2.26523	0.0296
SectorPublic	-0.934345	0.5628780	36	-1.65994	0.1056
ses:SectorPublic	1.173092	0.4677016	1935	2.50821	0.0122
cvar(ses, id):SectorPublic	1.414896	1.4652174	36	0.96566	0.3407

Correlation:

	(Intr) ses	cv(,i) SctrPb	ss:ScP
ses	-0.071		
cvar(ses, id)	-0.186	-0.248	

SectorPublic	-0.667	0.047	0.124		
ses:SectorPublic	0.047	-0.662	0.164	-0.065	
cvar(ses, id):SectorPublic	0.135	0.180	-0.725	0.056	-0.284

Standardized Within-Group Residuals:

Min	Q1	Med	Q3	Max
-3.09882513	-0.75336271	0.03194452	0.78173508	2.79671295

Number of Observations: 1977

Number of Groups: 40

wald(fitc)

numDF	denDF	F.value	p.value
6	36	407.3664	<.00001

Coefficients	Estimate	Std.Error	DF	t-value	p-value
(Intercept)	13.356053	0.375374	1935	35.580620	<.00001
ses	1.699397	0.309554	1935	5.489821	<.00001
cvar(ses, id)	2.407785	1.062930	36	2.265235	0.02961
SectorPublic	-0.934345	0.562878	36	-1.659942	0.10561
ses:SectorPublic	1.173092	0.467702	1935	2.508206	0.01222

```
cvar(ses, id):SectorPublic  1.414896  1.465217  36  0.965656 0.34066
```

Coefficients	Lower 0.95	Upper 0.95
(Intercept)	12.619872	14.092234
ses	1.092302	2.306492
cvar(ses, id)	0.252064	4.563507
SectorPublic	-2.075914	0.207225
ses:SectorPublic	0.255840	2.090344
cvar(ses, id):SectorPublic	-1.556703	4.386494

```
wald( fitc, -1)      # overall test of FE model
```

	numDF	denDF	F.value	p.value
1	5	36	33.98978	<.00001

	Estimate	Std.Error	DF	t-value	p-value
ses	1.699397	0.309554	1935	5.489821	<.00001
cvar(ses, id)	2.407785	1.062930	36	2.265235	0.02961
SectorPublic	-0.934345	0.562878	36	-1.659942	0.10561
ses:SectorPublic	1.173092	0.467702	1935	2.508206	0.01222
cvar(ses, id):SectorPublic	1.414896	1.465217	36	0.965656	0.34066

	Lower 0.95	Upper 0.95
ses	1.092302	2.306492
cvar(ses, id)	0.252064	4.563507
SectorPublic	-2.075914	0.207225
ses:SectorPublic	0.255840	2.090344
cvar(ses, id):SectorPublic	-1.556703	4.386494

```
wald( fitc, 'cvar')
```

	numDF	denDF	F.value	p.value
cvar	2	36	9.749657	0.00041

Coefficients	Estimate	Std.Error	DF	t-value	p-value
cvar(ses, id)	2.407785	1.062930	36	2.265235	0.02961
cvar(ses, id):SectorPublic	1.414896	1.465217	36	0.965656	0.34066

Coefficients	Lower 0.95	Upper 0.95
cvar(ses, id)	0.252064	4.563507
cvar(ses, id):SectorPublic	-1.556703	4.386494

```
wald( fitc, 'ses') # overall test of all 'ses' effects
```

	numDF	denDF	F.value	p.value
--	-------	-------	---------	---------

```
ses      4      36 39.23784 <.00001
```

Coefficients	Estimate	Std.Error	DF	t-value	p-value
ses	1.699397	0.309554	1935	5.489821	<.00001
cvar(ses, id)	2.407785	1.062930	36	2.265235	0.02961
ses:SectorPublic	1.173092	0.467702	1935	2.508206	0.01222
cvar(ses, id):SectorPublic	1.414896	1.465217	36	0.965656	0.34066

Coefficients	Lower 0.95	Upper 0.95
ses	1.092302	2.306492
cvar(ses, id)	0.252064	4.563507
ses:SectorPublic	0.255840	2.090344
cvar(ses, id):SectorPublic	-1.556703	4.386494

```
wald( fitc, 'Sector')  # overall test of all 'Sector' effects
```

	numDF	denDF	F.value	p.value
Sector	3	36	3.939573	0.01576

Coefficients	Estimate	Std.Error	DF	t-value	p-value
SectorPublic	-0.934345	0.562878	36	-1.659942	0.10561
ses:SectorPublic	1.173092	0.467702	1935	2.508206	0.01222

```
cvar(ses, id):SectorPublic  1.414896  1.465217   36  0.965656 0.34066
```

Coefficients	Lower 0.95	Upper 0.95
SectorPublic	-2.075914	0.207225
ses:SectorPublic	0.255840	2.090344
cvar(ses, id):SectorPublic	-1.556703	4.386494

```
wald( fitc, ':' )  # overall test of all interaction effects
```

	numDF	denDF	F.value	p.value
:	2	36	4.675875	0.01566

Coefficients	Estimate	Std.Error	DF	t-value	p-value
ses:SectorPublic	1.173092	0.467702	1935	2.508206	0.01222
cvar(ses, id):SectorPublic	1.414896	1.465217	36	0.965656	0.34066

Coefficients	Lower 0.95	Upper 0.95
ses:SectorPublic	0.255840	2.090344
cvar(ses, id):SectorPublic	-1.556703	4.386494

```
  # i.e. are lines parallel between sectors
```

Role of contextual variable for ses —

Including `cvar(ses, id)` has two important consequences:

- 1) It unbias the estimated effect of `ses` so it estimates the WITHIN-SCHOOL effect of `ses` unbiased by the between school effect of `ses`
- 2) It provides an estimate of the 'contextual effect' of `ses` and of the BETWEEN-SCHOOL (= compositional) effect of `ses` which is equal to the sum of the contextual and the within-effect

Interpreting the model with contextual effect —

```
wald( fitc )
```

numDF	denDF	F.value	p.value
6	36	407.3664	<.00001

Coefficients	Estimate	Std.Error	DF	t-value	p-value
(Intercept)	13.356053	0.375374	1935	35.580620	<.00001

ses	1.699397	0.309554	1935	5.489821	<.00001
cvar(ses, id)	2.407785	1.062930	36	2.265235	0.02961
SectorPublic	-0.934345	0.562878	36	-1.659942	0.10561
ses:SectorPublic	1.173092	0.467702	1935	2.508206	0.01222
cvar(ses, id):SectorPublic	1.414896	1.465217	36	0.965656	0.34066

Coefficients	Lower 0.95	Upper 0.95
(Intercept)	12.619872	14.092234
ses	1.092302	2.306492
cvar(ses, id)	0.252064	4.563507
SectorPublic	-2.075914	0.207225
ses:SectorPublic	0.255840	2.090344
cvar(ses, id):SectorPublic	-1.556703	4.386494

wald(fitc, 'cvar')

	numDF	denDF	F.value	p.value
cvar	2	36	9.749657	0.00041

Coefficients	Estimate	Std.Error	DF	t-value	p-value
cvar(ses, id)	2.407785	1.062930	36	2.265235	0.02961
cvar(ses, id):SectorPublic	1.414896	1.465217	36	0.965656	0.34066

Coefficients	Lower 0.95	Upper 0.95
cvar(ses, id)	0.252064	4.563507
cvar(ses, id):SectorPublic	-1.556703	4.386494

Anova(fitc)

Analysis of Deviance Table (Type II tests)

Response: mathach

	Chisq	Df	Pr(>Chisq)
ses	90.9731	1	< 2.2e-16 ***
cvar(ses, id)	18.5668	1	1.641e-05 ***
Sector	2.4670	1	0.11626
ses:Sector	6.2911	1	0.01213 *
cvar(ses, id):Sector	0.9325	1	0.33422

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Note that although neither 'cvar' component in the model seems overwhelmingly significant, the joint test that both are 0 is much more significant. This is a common kind of occurrence and illustrates how individual component p-values should not be used individually to judge the overall contribution of a variable.

Interpretation of the coefficients of in this model:

```
> wald( fitc )
```

```
numDF denDF  F.value p.value
```

```
6      76 555.2492 <.00001
```

```
-- This is a test that ALL parameters (including the intercept)
   are equal to 0. It is generally of no interest. The
   command 'wald(fitc, -1)' tests all parameters except the
   intercept and provides a test that model as a whole is
   not predictive.
```

```
numDF denDF  F.value p.value
```

```
6      36 407.3664 <.00001
```

Coefficients	Estimate	Std.Error	DF	t-value	p-value
(Intercept)	13.356053	0.375374	1935	35.580620	<.00001

```
-- E(Y) when ses = 0 for a Catholic school
   mean ses = 0
```

ses	1.699397	0.309554	1935	5.489821	<.00001
-----	----------	----------	------	----------	---------

-- Expected difference in $E(Y)$ associated with
a difference of 1 unit of individual ses
within a given Catholic school. Often called the
'WITHIN-SCHOOL' effect of ses.

cvar(ses, id)	2.407785	1.062930	36	2.265235	0.02961
---------------	----------	----------	----	----------	---------

-- Expected different in $E(Y)$ associated with
a difference in a Catholic school mean ses of 1 unit
with individual ses constant. Often call the
'CONTEXTUAL' effect of ses.

The sum of the 'WITHIN-SCHOOL' and the 'CONTEXTUAL'
effects is the 'COMPOSITIONAL' or 'BETWEEN-SCHOOL' effect

SectorPublic	-0.934345	0.562878	36	-1.659942	0.10561
--------------	-----------	----------	----	-----------	---------

-- Public intercept minus Catholic intercept

ses:SectorPublic	1.173092	0.467702	1935	2.508206	0.01222
------------------	----------	----------	------	----------	---------

```
-- coef of 'ses' in Public schools minus value in  
Catholic schools
```

```
cvar(ses, id):SectorPublic  1.414896  1.465217   36  0.965656 0.34066
```

```
-- ditto for coefficient of 'cvar(ses, id)'
```

Estimating the compositional effect (between effect) —

```
wald( fitc )
```

```
numDF denDF  F.value p.value  
   6    36 407.3664 <.00001
```

Coefficients	Estimate	Std.Error	DF	t-value	p-value
(Intercept)	13.356053	0.375374	1935	35.580620	<.00001
ses	1.699397	0.309554	1935	5.489821	<.00001
cvar(ses, id)	2.407785	1.062930	36	2.265235	0.02961
SectorPublic	-0.934345	0.562878	36	-1.659942	0.10561
ses:SectorPublic	1.173092	0.467702	1935	2.508206	0.01222
cvar(ses, id):SectorPublic	1.414896	1.465217	36	0.965656	0.34066

Coefficients	Lower 0.95	Upper 0.95
(Intercept)	12.619872	14.092234
ses	1.092302	2.306492
cvar(ses, id)	0.252064	4.563507
SectorPublic	-2.075914	0.207225
ses:SectorPublic	0.255840	2.090344
cvar(ses, id):SectorPublic	-1.556703	4.386494

All 'possible' effects of ses:

They're actually 'infinite' consisting of ALL linear combinations of the row space of the following matrix which has dimension 4.

SO: infinite in number but finite in dimension

```
L <- rbind(
  'Within-school effect of ses | Cath' = c(0, 1, 0, 0, 0, 0),
  'Contextual effect of ses | Cath'   = c(0, 0, 1, 0, 0, 0),
  'Between-school effect of ses | Cath' = c(0, 1, 1, 0, 0, 0),
  'Within-school effect of ses | Pub'  = c(0, 1, 0, 0, 1, 0),
  'Contextual effect of ses | Pub'     = c(0, 0, 1, 0, 0, 1),
  'Between-school effect of ses | Pub' = c(0, 1, 1, 0, 1, 1),
```

```

'Within-school effect of ses | Pub - Cath' = c(0, 0, 0, 0, 1, 0),
'Contextual effect of ses | Pub - Cath'    = c(0, 0, 0, 0, 0, 1),
'Between-school effect of ses | Pub - Cath' = c(0, 0, 0, 0, 1, 1))
wald( fitc, L)

```

```

      numDF denDF  F.value p.value
1         4     36 39.23784 <.00001

```

	Estimate	Std.Error	DF
Within-school effect of ses Cath	1.699397	0.309554	1935
Contextual effect of ses Cath	2.407785	1.062930	36
Between-school effect of ses Cath	4.107183	1.030638	36
Within-school effect of ses Pub	2.872489	0.350601	1935
Contextual effect of ses Pub	3.822681	1.008485	36
Between-school effect of ses Pub	6.695170	0.956243	36
Within-school effect of ses Pub - Cath	1.173092	0.467702	1935
Contextual effect of ses Pub - Cath	1.414896	1.465217	36
Between-school effect of ses Pub - Cath	2.587987	1.405921	36

	t-value	p-value	Lower	0.95
Within-school effect of ses Cath	5.489821	<.00001	1.092302	
Contextual effect of ses Cath	2.265235	0.02961	0.252064	

Between-school effect of ses Cath	3.985090	0.00031	2.016953
Within-school effect of ses Pub	8.193048	<.00001	2.184894
Contextual effect of ses Pub	3.790518	0.00055	1.777378
Between-school effect of ses Pub	7.001537	<.00001	4.755820
Within-school effect of ses Pub - Cath	2.508206	0.01222	0.255840
Contextual effect of ses Pub - Cath	0.965656	0.34066	-1.556703
Between-school effect of ses Pub - Cath	1.840777	0.07391	-0.263353

Upper 0.95

Within-school effect of ses Cath	2.306492
Contextual effect of ses Cath	4.563507
Between-school effect of ses Cath	6.197413
Within-school effect of ses Pub	3.560084
Contextual effect of ses Pub	5.867984
Between-school effect of ses Pub	8.634521
Within-school effect of ses Pub - Cath	2.090344
Contextual effect of ses Pub - Cath	4.386494
Between-school effect of ses Pub - Cath	5.439328

We can ask 'subquestions' with multiple dfs

Question: Is there evidence of an effect of ses in Catholic schools?

```
wald(fitc, list('ses in Catholic schools' = L[1:3,]))
```

	numDF	denDF	F.value	p.value
ses in Catholic schools	2	36	22.08524	<.00001

	Estimate	Std.Error	DF	t-value
Within-school effect of ses Cath	1.699397	0.309554	1935	5.489821
Contextual effect of ses Cath	2.407785	1.062930	36	2.265235
Between-school effect of ses Cath	4.107183	1.030638	36	3.985090

	p-value	Lower 0.95	Upper 0.95
Within-school effect of ses Cath	<.00001	1.092302	2.306492
Contextual effect of ses Cath	0.02961	0.252064	4.563507
Between-school effect of ses Cath	0.00031	2.016953	6.197413

Question: Is there evidence of an effect of ses in Public schools?

```
wald(fitc, list('ses in Public schools' = L[4:6,]))
```

	numDF	denDF	F.value	p.value
ses in Public schools	2	36	56.39045	<.00001

	Estimate	Std.Error	DF	t-value
--	----------	-----------	----	---------

Within-school effect of ses Pub	2.872489	0.350601	1935	8.193048
Contextual effect of ses Pub	3.822681	1.008485	36	3.790518
Between-school effect of ses Pub	6.695170	0.956243	36	7.001537

	p-value	Lower 0.95	Upper 0.95
Within-school effect of ses Pub	<.00001	2.184894	3.560084
Contextual effect of ses Pub	0.00055	1.777378	5.867984
Between-school effect of ses Pub	<.00001	4.755820	8.634521

Question: Is there evidence of a difference between Public and Catholic?

```
wald(fitc, list('Difference between effect of ses between Sectors' = L[7:9,]))
```

	numDF	denDF	F.value
Difference between effect of ses between Sectors	2	36	4.675875

p.value

Difference between effect of ses between Sectors	0.01566
--	---------

	Estimate	Std.Error	DF
Within-school effect of ses Pub - Cath	1.173092	0.467702	1935
Contextual effect of ses Pub - Cath	1.414896	1.465217	36
Between-school effect of ses Pub - Cath	2.587987	1.405921	36

	t-value	p-value	Lower 0.95
Within-school effect of ses Pub - Cath	2.508206	0.01222	0.255840
Contextual effect of ses Pub - Cath	0.965656	0.34066	-1.556703
Between-school effect of ses Pub - Cath	1.840777	0.07391	-0.263353

	Upper 0.95
Within-school effect of ses Pub - Cath	2.090344
Contextual effect of ses Pub - Cath	4.386494
Between-school effect of ses Pub - Cath	5.439328

or

```
wald(fitc, ':')
```

	numDF	denDF	F.value	p.value
:	2	36	4.675875	0.01566

Coefficients	Estimate	Std.Error	DF	t-value	p-value
ses:SectorPublic	1.173092	0.467702	1935	2.508206	0.01222
cvar(ses, id):SectorPublic	1.414896	1.465217	36	0.965656	0.34066

Coefficients	Lower 0.95	Upper 0.95
ses:SectorPublic	0.255840	2.090344

```
cvar(ses, id):SectorPublic  -1.556703  4.386494
```

Note: that the estimated different in contextual effect is larger than the difference in within effect, yet the p-values are ‘reversed’.

This illustrates the fact that there is generally more information – i.e. smaller SEs – for within effects than for contextual effects – although not always.

Although the interaction is not significant, there is much less power to detect between-school effects than within-school effects and depending on the purposes of the analysis it might not be reasonable to assume that the slopes are equal just because the p-value does not provide evidence that they are not.

QUESTIONS:

1. How would you test whether there is ANY difference between the two sectors – not just in the effect of SES.
2. Why does the second to last F test have 2 numDFs although there are three lines in the L matrix?

Visualizing the fitted model —

Effect plots with the effects package

The **effects** package, provides an excellent way of visualizing the effects of terms in a model. If a model has

- categorical variables with more than 2 levels or
- interactions

the table of estimated coefficients and their standard errors give a very incomplete picture of the model.

Effect plots as implemented in the ‘effects’ package let you see the effect of categorical variables at all levels and the conditional effect of interacting variables at various levels of the other variables with which they interact. They convey a great deal more than the numerical summary of estimates and standard deviations and they should be used routinely.

There is one caveat in using effect plots with mixed models in that they may, when looking at effects of level-1 variables, give the impression that an effect is less important than it actually is. The effect plot shows error bands around the predicted value of the response. With mixed models, there could be a large standard error around predicted values but a relatively small standard error around the difference of two predicted values.

Thus, testing specific hypotheses with linear hypothesis matrices or constructing graphical estimates of quantities other than predicted values can play a role in the analysis and reporting of results.

```
library(effects)
```

We need to refit with a computed variable for `cvar(ses, id)`

```
dd$ses.cvar <- with(dd, cvar(ses, id))  
fitc2 <- update(fitc, . ~ (ses + ses.cvar) * Sector)
```

Range of predicted values given levels of interacting variables

```
predictorEffects(fitc2, 'ses')
```

```
ses predictor effect
```

```
ses*Sector effect
```

```
Sector
```

ses	Catholic	Public
-2	9.892629	6.574122
-1	11.592026	9.446611
-0.4	12.611665	11.170105

0.6	14.311062	14.042594
2	16.690219	18.064079

```
predictorEffects(fitc2, 'ses.cvar')
```

ses.cvar predictor effect

ses.cvar*Sector effect

	Sector	
ses.cvar	Catholic	Public
-1	10.90265	8.521924
-0.7	11.62499	9.668729
-0.3	12.58810	11.197801
0.2	13.79200	13.109142
0.6	14.75511	14.638214

```
predictorEffects(fitc2, 'Sector')
```

Sector predictor effect

Sector*ses*ses.cvar effect

, , ses.cvar = -1

	ses				
Sector	-2	-1	-0.4	0.6	2
Catholic	7.549473	9.248870	10.268509	11.96791	14.34706
Public	2.854048	5.726538	7.450031	10.32252	14.34401

, , ses.cvar = -0.7

	ses				
Sector	-2	-1	-0.4	0.6	2
Catholic	8.271808	9.971206	10.990844	12.69024	15.06940
Public	4.000853	6.873342	8.596835	11.46932	15.49081

, , ses.cvar = -0.3

	ses				
Sector	-2	-1	-0.4	0.6	2
Catholic	9.234922	10.934320	11.95396	13.65336	16.03251
Public	5.529925	8.402414	10.12591	12.99840	17.01988

```
, , ses.cvar = 0.2
```

	ses				
Sector	-2	-1	-0.4	0.6	2
Catholic	10.438815	12.13821	13.15785	14.85725	17.23640
Public	7.441266	10.31375	12.03725	14.90974	18.93122

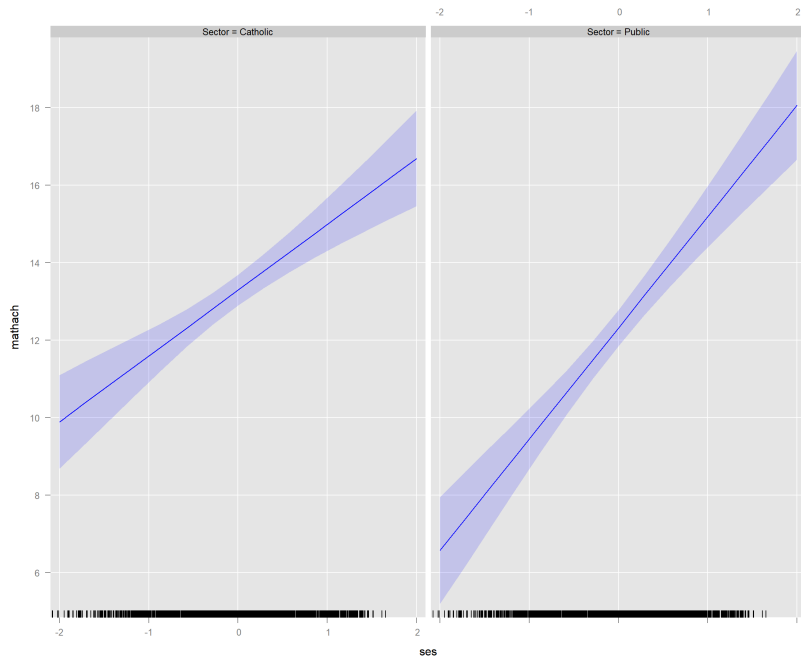
```
, , ses.cvar = 0.6
```

	ses				
Sector	-2	-1	-0.4	0.6	2
Catholic	11.401929	13.10133	14.12097	15.82036	18.19952
Public	8.970338	11.84283	13.56632	16.43881	20.46029

Plot:

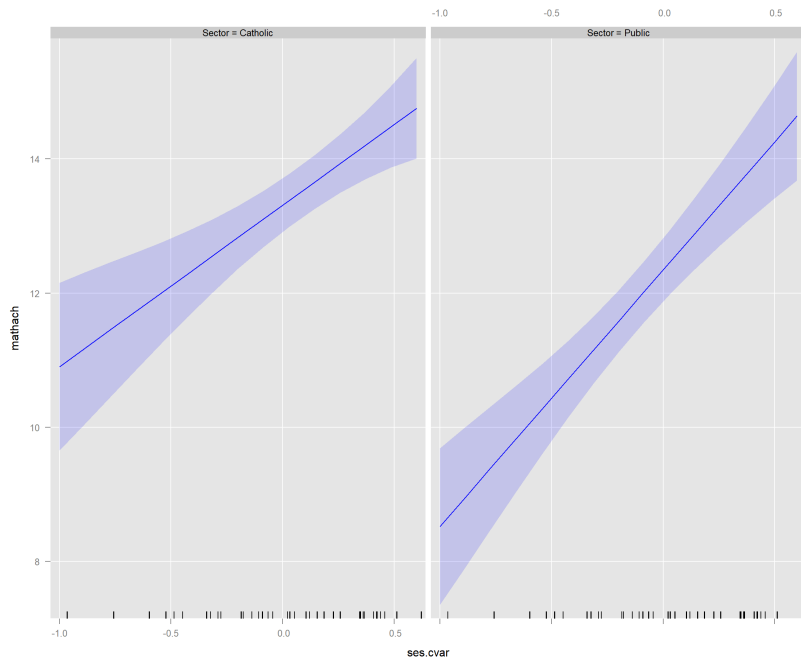
```
predictorEffects(fitc2, 'ses') %>% plot
```

ses predictor effect plot



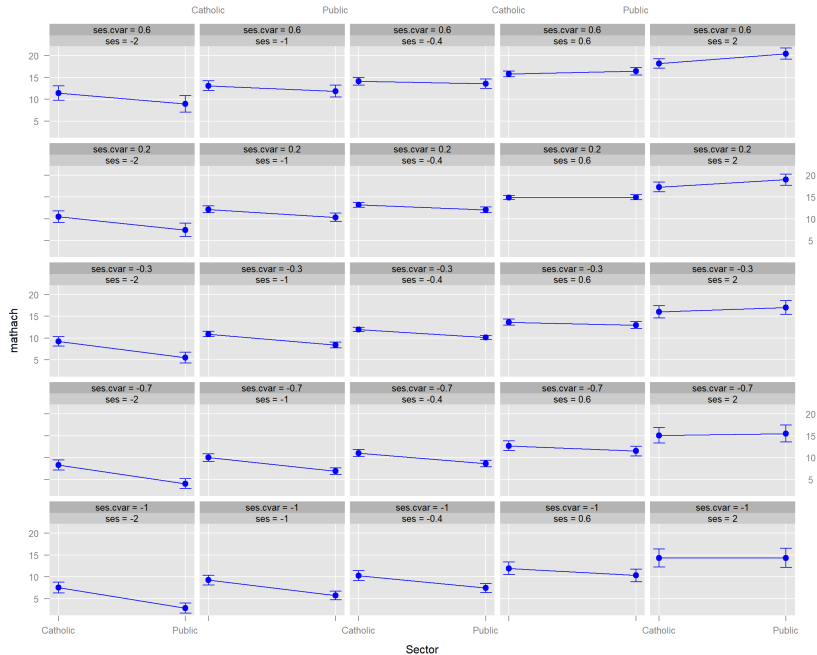
```
predictorEffects(fitc2, 'ses.cvar') %>% plot
```

ses.cvar predictor effect plot



```
predictorEffects(fitc2, 'Sector') %>% plot
```

Sector predictor effect plot



Handmade prediction plot

We create a data frame with predictor values for which we want to visualize the predicted values for the model.

Since the data have two levels, we create a data set with school characteristics and deviations of students within schools

```
pred <- expand.grid(ses.mean = c(-1,0,1),  
                  Sector = levels(dd$Sector), # this ensure that the factor is  
                  ses.dev = seq(-1,1,.1))    # properly  
pred$ses <- with(pred, ses.mean + ses.dev)  
pred$m.ses <- with(pred, paste0(Sector, " mean ses: ", ses.mean))  
some(pred)
```

	ses.mean	Sector	ses.dev	ses	m.ses
2	0	Catholic	-1.0	-1.0	Catholic mean ses: 0
17	0	Public	-0.8	-0.8	Public mean ses: 0
30	1	Public	-0.6	0.4	Public mean ses: 1
31	-1	Catholic	-0.5	-1.5	Catholic mean ses: -1
32	0	Catholic	-0.5	-0.5	Catholic mean ses: 0
42	1	Public	-0.4	0.6	Public mean ses: 1
67	-1	Catholic	0.1	-0.9	Catholic mean ses: -1

78	1	Public	0.2	1.2	Public mean ses: 1
92	0	Catholic	0.5	0.5	Catholic mean ses: 0
107	0	Public	0.7	0.7	Public mean ses: 0

We can now use the wald function to generate predicted values and standard errors

The 'getX' function generates the design matrix over the data frame pred

Note that it is possible to use the 'predict' method but it does not provide standard errors

```
round( head( getX(fitc, pred)),2)
```

```
Error in capply.default(x, id, mean, na.rm = na.rm, ...): object 'id' not found
```

```
ww <- as.data.frame( wald( fitc, getX(fitc, pred)) )
```

```
Error in capply.default(x, id, mean, na.rm = na.rm, ...): object 'id' not found
```

```
head(ww)
```

```
Error in head(ww): object 'ww' not found
```

```
xyplot( coef ~ ses | m.ses, ww, type = 'l')
```

```
Error in eval(substitute(groups), data, environment(x)): object 'ww' not found
```

```
gd(6, lwd = 2) # select colours
```

```
xyplot( coef ~ ses | Sector, ww, groups = m.ses, type = 'l')
```

Error in eval(substitute(groups), data, environment(x)): object 'ww' not found

```
xyplot( coef ~ ses | Sector, ww, groups = m.ses,  
        auto.key = list(columns = 2, lines = T, points = F), type = 'l')
```

Error in eval(substitute(groups), data, environment(x)): object 'ww' not found

```
xyplot( coef ~ ses , ww, groups = m.ses,  
        auto.key = list(columns = 2, lines = T, points = F), type = 'l')
```

Error in eval(substitute(groups), data, environment(x)): object 'ww' not found

```
gd(col=rep(c('red','blue'),each = 3), lty = rep(1:3,2), lwd = 2)
```

```
xyplot( coef ~ ses , ww, groups = m.ses,  
        ylab = 'predicted math achievement',  
        auto.key = list(columns = 2, lines = T, points = F), type = 'l') +  
layer(panel.abline(v=c(-1,0,1), lty = 2))
```

Error in eval(substitute(groups), data, environment(x)): object 'ww' not found

EXERCISE:

- Show how the estimated coefficients are related to the last plot.

Plotting error bars —

The data frame created by `wald` includes the information necessary to plot error bounds on the predicted lines

The ‘panel function’ `panel.fit` will produce the error bars. The easiest way to use it is with ‘layers’ provided by the ‘`latticeExtra`’ package

In the call to `xyplot`, provide arguments for the following parameters:

- `fit`: the fitted value for which a line or curve will be plotted
- `upper`: the upper boundary of the error bar
- `lower`: the lower boundary
- `subscripts = T` (so `xyplot` will pass the necessary information to the panel function)

Then you need to add a ‘+’ at the end of the ‘`xyplot`’ command and follow that with:

`layer(panel.fit(...))` if you did not use groups, and

`glayer(panel.fit(...))` if you did use groups

`head(ww)`

```
Error in head(ww): object 'ww' not found
```

```
xyplot( coef ~ ses | Sector,
        ww, groups = m.ses,
        ylab = 'predicted math achievement',
        fit = ww$coef,
        upper = ww$U2, lower = ww$L2,
        subscripts = T,
        sub = 'predicted mean with appx. 95% error bars (NOT prediction bars)',
        auto.key = list(columns = 2, lines = T, points = F), type = 'l') +
glayer(panel.fit(...))
```

Error in xyplot.formula(coef ~ ses | Sector, ww, groups = m.ses, ylab = "pred

```
xyplot( coef ~ ses | ses.mean,
        ww, groups = m.ses,
        ylab = 'predicted math achievement',
        fit = ww$coef,
        upper = ww$U2, lower = ww$L2,
        subscripts = T,
        layout = c(1,3),
        sub = 'predicted mean with appx. 95% error bars (NOT prediction bars)',
        auto.key = list(columns = 2, lines = T, points = F), type = 'l') +
glayer(panel.fit(...))
```

```
Error in xyplot.formula(coef ~ ses | ses.mean, ww, groups = m.ses, ylab = "pr
xyplot( coef ~ ses | paste0('Mean school ses = ',ses.mean),
  ww, groups = m.ses,
  ylab = 'predicted math achievement',
  fit = ww$coef,
  upper = ww$U2, lower = ww$L2,
  subscripts = T,
  layout = c(1,3),
  sub = 'predicted mean with appx. 95% error bars (NOT prediction bars)',
  auto.key = list(columns = 2, lines = T, points = F), type = 'l') +
glayer(panel.fit(...))
```

```
Error in xyplot.formula(coef ~ ses | paste0("Mean school ses = ", ses.mean),
```

Estimating the gap

We would like to estimate the intersector gap at the same value of ses.mean and ses.

```
head(ww)
```

```
Error in head(ww): object 'ww' not found
```

```
vals <- ww[, c('ses', 'm.ses', 'ses.mean', 'Sector')]
```

Error in eval(expr, envir, enclos): object 'ww' not found

Construct an L matrix to estimate the gap

This is like estimating the 'derivative' of each term wrt Sector

```
wald(fitc)
```

numDF	denDF	F.value	p.value
6	36	407.3664	<.00001

Coefficients	Estimate	Std.Error	DF	t-value	p-value
(Intercept)	13.356053	0.375374	1935	35.580620	<.00001
ses	1.699397	0.309554	1935	5.489821	<.00001
cvar(ses, id)	2.407785	1.062930	36	2.265235	0.02961
SectorPublic	-0.934345	0.562878	36	-1.659942	0.10561
ses:SectorPublic	1.173092	0.467702	1935	2.508206	0.01222
cvar(ses, id):SectorPublic	1.414896	1.465217	36	0.965656	0.34066

Coefficients	Lower 0.95	Upper 0.95
(Intercept)	12.619872	14.092234
ses	1.092302	2.306492

cvar(ses, id)	0.252064	4.563507
SectorPublic	-2.075914	0.207225
ses:SectorPublic	0.255840	2.090344
cvar(ses, id):SectorPublic	-1.556703	4.386494

```
Lgap <- with(vals,
             cbind(0, 0, 0, 1, ses, ses.mean)
)
```

Error in with(vals, cbind(0, 0, 0, 1, ses, ses.mean)): object 'vals' not found

```
some(Lgap)
```

Error in some(Lgap): object 'Lgap' not found

```
ww <- as.data.frame(wald(fitc, Lgap, data = vals ))
```

Error in wald(fitc, Lgap, data = vals): object 'Lgap' not found

```
some(ww)
```

Error in some(ww): object 'ww' not found

```
gd(col = 'blue')
```

```
xyplot( coef ~ ses | paste0('Mean school ses = ',ses.mean),
        ww,
```

```

ylim = c(-8,8), xlim = c(-2,2),
ylab = 'predicted math achievement',
fit = ww$coef,
upper = ww$U2, lower = ww$L2,
subscripts = T,
layout = c(1,3),
type = 'l',
sub = 'estimated difference (Public - Catholic) with appx. 95% error bars'
layer(panel.abline(h = 0))+
layer(panel.fit(...))

```

Error in `xyplot.formula(coef ~ ses | paste0("Mean school ses = ", ses.mean),`

```

xyplot( coef ~ ses | paste0('Mean school ses = ',ses.mean),
  ww,
  ylim = c(-8,8), xlim = c(-2,2),
  ylab = 'predicted math achievement',
  fit = ww$coef,
  upper = ww$U2, lower = ww$L2,
  subscripts = T,
  layout = c(3,1),
  type = 'l',

```

```

      sub = 'estimated difference (Public - Catholic) with appx. 95% error bar
layer(panel.abline(h = 0, lwd = 2 ))+
layer(panel.fit(...))

```

Error in `xyplot.formula(coef ~ ses | paste0("Mean school ses = ", ses.mean),`

We can combine the bands in one panel, although it isn't very effective here.

Note the changes:

- the `| panel` argument becomes a `'groups'` argument
- remove the `'layout'` argument
- add `'auto.key'`
- `'layer'` become `'glayer'`
- everything else can stay the same

```

gd(3)
xyplot( coef ~ ses ,
        ww,
        groups = paste0('Mean school ses = ',ses.mean),
        ylim = c(-8,8), xlim = c(-2,2),
        ylab = 'predicted math achievement',
        fit = ww$coef,
        upper = ww$U2, lower = ww$L2,

```

```

subscripts = T,
type = 'l',
auto.key = T,
sub = 'estimated difference (Public - Catholic) with appx. 95% error bar'
glayer(panel.abline(h = 0, lwd = 2 ))+
glayer(panel.fit(...))

```

Error in xyplot.formula(coef ~ ses, ww, groups = paste0("Mean school ses = ",

Figure: Combining bands in one panel. Tastes may vary!

EXERCISES —

1. Starting with the model above, test whether Sex improves prediction in the model. Consider including a contextual variable for Sex. What is its interpretation?
2. Is it reasonable to use the contextual variable for Sex as the only regressor for the contextual effect of Sex or should other variables be used as well? Can you create these variables and include them in your analysis? What do they reveal?

See Visualizing the coefficients

Using CWG instead of raw SES —

```
fitcd <- update( fitc, . ~ (cvar(ses,id) + dvar(ses,id)*Sector))  
summary( fitcd )
```

Linear mixed-effects model fit by REML

Data: dd

AIC	BIC	logLik
12843.4	12893.68	-6412.7

Random effects:

Formula: ~1 + ses | id

Structure: General positive-definite, Log-Cholesky parametrization

	StdDev	Corr
(Intercept)	1.5088268	(Intr)
ses	0.4215203	-0.202
Residual	6.1173112	

Fixed effects: mathach ~ cvar(ses, id) + dvar(ses, id) + Sector + dvar(ses,

	Value	Std.Error	DF	t-value	p-value
(Intercept)	13.257428	0.3803559	1935	34.85533	0.0000
cvar(ses, id)	5.477723	0.7136326	37	7.67583	0.0000

```
dvar(ses, id)          1.711278 0.3023289 1935  5.66032  0.0000
SectorPublic          -0.981203 0.5765720   37 -1.70179  0.0972
dvar(ses, id):SectorPublic 1.161565 0.4571695 1935  2.54078  0.0111
  Correlation:
                        (Intr) cv(,i) dv(,i) SctrPb
cvar(ses, id)          -0.140
dvar(ses, id)          -0.052  0.020
SectorPublic           -0.682  0.251  0.038
dvar(ses, id):SectorPublic 0.033 -0.003 -0.661 -0.052
```

Standardized Within-Group Residuals:

	Min	Q1	Med	Q3	Max
	-3.12476554	-0.74613310	0.02214392	0.78204903	2.76676038

Number of Observations: 1977

Number of Groups: 40

summary(fitc)

Linear mixed-effects model fit by REML

Data: dd

AIC	BIC	logLik
-----	-----	--------

12839.6 12895.47 -6409.801

Random effects:

Formula: ~1 + ses | id

Structure: General positive-definite, Log-Cholesky parametrization

	StdDev	Corr
(Intercept)	1.4561519	(Intr)
ses	0.5190639	-0.217
Residual	6.1140010	

Fixed effects: mathach ~ (ses + cvar(ses, id)) * Sector

	Value	Std.Error	DF	t-value	p-value
(Intercept)	13.356053	0.3753744	1935	35.58062	0.0000
ses	1.699397	0.3095543	1935	5.48982	0.0000
cvar(ses, id)	2.407785	1.0629298	36	2.26523	0.0296
SectorPublic	-0.934345	0.5628780	36	-1.65994	0.1056
ses:SectorPublic	1.173092	0.4677016	1935	2.50821	0.0122
cvar(ses, id):SectorPublic	1.414896	1.4652174	36	0.96566	0.3407

Correlation:

	(Intr) ses	cv(,i) SctrPb	ss:ScP
ses	-0.071		

```

cvar(ses, id)                -0.186 -0.248
SectorPublic                  -0.667  0.047  0.124
ses:SectorPublic              0.047 -0.662  0.164 -0.065
cvar(ses, id):SectorPublic    0.135  0.180 -0.725  0.056 -0.284

```

Standardized Within-Group Residuals:

	Min	Q1	Med	Q3	Max
	-3.09882513	-0.75336271	0.03194452	0.78173508	2.79671295

Number of Observations: 1977

Number of Groups: 40

wald(fitc, L)

	numDF	denDF	F.value	p.value
1	4	36	39.23784	<.00001

	Estimate	Std.Error	DF
Within-school effect of ses Cath	1.699397	0.309554	1935
Contextual effect of ses Cath	2.407785	1.062930	36
Between-school effect of ses Cath	4.107183	1.030638	36
Within-school effect of ses Pub	2.872489	0.350601	1935

Contextual effect of ses Pub	3.822681	1.008485	36
Between-school effect of ses Pub	6.695170	0.956243	36
Within-school effect of ses Pub - Cath	1.173092	0.467702	1935
Contextual effect of ses Pub - Cath	1.414896	1.465217	36
Between-school effect of ses Pub - Cath	2.587987	1.405921	36

	t-value	p-value	Lower 0.95
Within-school effect of ses Cath	5.489821	<.00001	1.092302
Contextual effect of ses Cath	2.265235	0.02961	0.252064
Between-school effect of ses Cath	3.985090	0.00031	2.016953
Within-school effect of ses Pub	8.193048	<.00001	2.184894
Contextual effect of ses Pub	3.790518	0.00055	1.777378
Between-school effect of ses Pub	7.001537	<.00001	4.755820
Within-school effect of ses Pub - Cath	2.508206	0.01222	0.255840
Contextual effect of ses Pub - Cath	0.965656	0.34066	-1.556703
Between-school effect of ses Pub - Cath	1.840777	0.07391	-0.263353

	Upper 0.95
Within-school effect of ses Cath	2.306492
Contextual effect of ses Cath	4.563507
Between-school effect of ses Cath	6.197413

Within-school effect of ses Pub	3.560084
Contextual effect of ses Pub	5.867984
Between-school effect of ses Pub	8.634521
Within-school effect of ses Pub - Cath	2.090344
Contextual effect of ses Pub - Cath	4.386494
Between-school effect of ses Pub - Cath	5.439328

QUESTIONS:

1. what's the same and what's different?
2. How are the various effects of ses related
3. Create an L matrix to estimate the 3 effects of ses with fitted as in the previous example

Note: replacing ses with dvar(ses,id) in the RE model does give an **mathematically** equivalent model but **NOT** a **numerically** equivalent model.

That is, with calculations using infinite precision the the two models would give equivalent results but *with finite precision* the 'dvar(ses,id)' model will, in general, converge more easily.

CWG vs raw in RE model —

Alternative but non-equivalent RE parametrization

Using 'dvar(ses,id)' or raw 'ses' in the FE model produces an equivalent model.

This is not so in the RE model. The

'random = ~ 1 + ses | id' (1)

model is essentially different from the

'random = ~ 1 + dvar(ses,id) | id' (2)

model.

In model (1) the 'point of minimum variance' of school regression lines corresponds to a *common* value of ses for all schools.

In model (2) the 'point of minimum variance' of school regression lines has a consistent value *relative* to each school's mean ses.

```
fitca <- update( fitc, random = ~ 1 + dvar( ses, id ) | id )  
summary( fitca )
```

Linear mixed-effects model fit by REML

Data: dd

AIC	BIC	logLik
12839.22	12895.09	-6409.612

Random effects:

Formula: ~1 + dvar(ses, id) | id

Structure: General positive-definite, Log-Cholesky parametrization

	StdDev	Corr
(Intercept)	1.4625200	(Intr)
dvar(ses, id)	0.6802261	-0.071
Residual	6.1084863	

Fixed effects: mathach ~ (ses + cvar(ses, id)) * Sector

	Value	Std.Error	DF	t-value	p-value
(Intercept)	13.374349	0.3732778	1935	35.82948	0.0000
ses	1.695976	0.3241495	1935	5.23208	0.0000
cvar(ses, id)	2.398991	1.0734653	36	2.23481	0.0317
SectorPublic	-0.939334	0.5625274	36	-1.66985	0.1036
ses:SectorPublic	1.171880	0.4888990	1935	2.39698	0.0166
cvar(ses, id):SectorPublic	1.238194	1.4581049	36	0.84918	0.4014

Correlation:

	(Intr) ses	cv(,i) SctrPb	ss:ScP
ses	-0.028		
cvar(ses, id)	-0.159	-0.300	
SectorPublic	-0.664	0.019	0.106

```
ses:SectorPublic          0.019 -0.663  0.199 -0.026
cvar(ses, id):SectorPublic 0.117  0.221 -0.736  0.078 -0.336
```

Standardized Within-Group Residuals:

	Min	Q1	Med	Q3	Max
	-3.10199875	-0.75105077	0.03298316	0.78240512	2.80416484

Number of Observations: 1977

Number of Groups: 40

summary(fitc)

Linear mixed-effects model fit by REML

Data: dd

	AIC	BIC	logLik
	12839.6	12895.47	-6409.801

Random effects:

Formula: ~1 + ses | id

Structure: General positive-definite, Log-Cholesky parametrization

	StdDev	Corr
--	--------	------

(Intercept)	1.4561519	(Intr)
-------------	-----------	--------

```
ses          0.5190639 -0.217
Residual     6.1140010
```

Fixed effects: mathach ~ (ses + cvar(ses, id)) * Sector

	Value	Std.Error	DF	t-value	p-value
(Intercept)	13.356053	0.3753744	1935	35.58062	0.0000
ses	1.699397	0.3095543	1935	5.48982	0.0000
cvar(ses, id)	2.407785	1.0629298	36	2.26523	0.0296
SectorPublic	-0.934345	0.5628780	36	-1.65994	0.1056
ses:SectorPublic	1.173092	0.4677016	1935	2.50821	0.0122
cvar(ses, id):SectorPublic	1.414896	1.4652174	36	0.96566	0.3407

Correlation:

	(Intr) ses	cv(,i)	SctrPb	ss:ScP
ses	-0.071			
cvar(ses, id)	-0.186	-0.248		
SectorPublic	-0.667	0.047	0.124	
ses:SectorPublic	0.047	-0.662	0.164	-0.065
cvar(ses, id):SectorPublic	0.135	0.180	-0.725	0.056 -0.284

Standardized Within-Group Residuals:

Min	Q1	Med	Q3	Max
-----	----	-----	----	-----

-3.09882513 -0.75336271 0.03194452 0.78173508 2.79671295

Number of Observations: 1977

Number of Groups: 40

Which is better? It will depend on the true state of nature.

What do the data tell us?

Neither model is nested in the other – in fact they have the same DFs – so we can't get a p-value for a test. We can compare with AIC:

```
anova(fitca, fitc)
```

	Model	df	AIC	BIC	logLik
fitca	1	10	12839.22	12895.09	-6409.612
fitc	2	10	12839.60	12895.47	-6409.801

fitca looks better, barely!

More importantly, it seems to make more sense.

EXERCISES —

1. Now that we have another Level 2 variable (`cvar(ses, id)`), we could consider whether there are interesting interactions between it and other variables.

Fit a model with interactions between `cvar(ses,id)` and other variables.

2. Is there evidence that these interactions are non-zero in the population?
 - BEWARE: Do not merely look at the individual p-value Either remove effects one at a time or test effects simultaneously with ‘wald’ or anova (need to refit with `method = “ML”`)
 - What do you conclude wrt interactions?

REML vs ML —

Previous fits with REML (default)

Here we use a ‘sub-par’ model (no contextual effect) to illustrate convergence issues.

FE: $\sim 1 + \text{ses} + \text{Sector} + \text{ses}:\text{Sector}$

RE: $\sim 1 + \text{ses} \mid \text{id}$

Fitting with 'REML'

Maximum likelihood optimization is done only on the RE model, i.e. the G matrix and sigma – or G and R. The FE model is then estimated using GLS. This makes optimization easier because there are fewer parameters – in this case 4:

1. 2 SD parameters for REs: 'beta_1' and 'beta_ses' (diagonal of G)
2. covariance parameter for REs: between 'beta_1', and 'beta_ses' (off-diagonal of G)
3. the SD parameter for individual error

Fit with 'ML'

Maximum likelihood optimization done on all parameters together. Generally more numerically difficult.

Pros and Cons:

REML: better estimates of variances, good for Wald tests. But, can use Likelihood-Ratio Tests (LRT with anova) only to compare models with identical FEs and nested RE models. However, beware that LRTs for RE models are problematic near the 'edge' of the model, e.g.

- if the null model is on the 'edge' of the full model
- if an estimated G matrix is nearly singular

ML: Not so good estimates of variances, not so good for Wald tests. But LRTs can compare models that are nested wrt to FEs and or REs.

REML fit (default):

```
fit <- lme( mathach ~ ses * Sector, dd,  
            random = ~ 1 + ses | id,  
            na.action = na.exclude,  
            control = list(msMaxIter=200, msVerbose=T))
```

0:	11105.191:	1.22722	2.27855	-3.36583
1:	11105.172:	1.23706	3.08949	-3.39049
2:	11105.171:	1.19748	3.08072	-3.39188
3:	11105.161:	1.21737	3.07678	-3.39235
4:	11105.096:	1.22151	2.70896	-3.41493
5:	11105.089:	1.22011	2.64468	-3.43555
6:	11105.085:	1.22389	2.55034	-3.50479
7:	11105.083:	1.22385	2.57106	-3.54636
8:	11105.076:	1.22534	2.60016	-3.72983
9:	11105.057:	1.23381	2.66214	-4.47027
10:	11105.033:	1.25234	2.72319	-5.87758
11:	11105.019:	1.27696	2.75593	-7.28573
12:	11105.013:	1.30246	2.78019	-8.50888

13:	11105.011:	1.31965	2.79844	-9.36213
14:	11105.011:	1.33116	2.81675	-9.93866
15:	11105.011:	1.33714	2.82776	-10.3398
16:	11105.011:	1.33993	2.83733	-10.5086
17:	11105.011:	1.33984	2.84079	-10.5414
18:	11105.011:	1.33908	2.84283	-10.5393
19:	11105.011:	1.33903	2.84278	-10.5371

ML fit:

```
# fit.ml <- lme( mathach ~ ses * Sector, dd,  
#               random = ~ 1 + ses | id,  
#               na.action = na.exclude,  
#               method = "ML",  
#               control = list(msMaxIter=200, msVerbose=T))
```

equivalent:

```
fit.ml <- update(fit, method = 'ML')
```

0:	11116.488:	1.26387	2.36435	-3.50300
1:	11116.369:	1.27189	3.26343	-3.52690
2:	11116.362:	1.25371	3.26152	-3.52738
3:	11116.357:	1.26159	3.19060	-3.54350

4:	11116.344:	1.26820	3.00294	-3.58152
5:	11116.336:	1.27032	2.81688	-3.62716
6:	11116.332:	1.26359	2.81941	-3.65385
7:	11116.330:	1.25804	2.81817	-3.71167
8:	11116.329:	1.25586	2.81904	-3.76973
9:	11116.324:	1.25192	2.81640	-4.00208
10:	11116.314:	1.24864	2.82902	-4.61237
11:	11116.304:	1.25175	2.85724	-5.22213
12:	11116.277:	1.27266	2.93789	-7.00650
13:	11116.252:	1.30765	3.05638	-9.17520
14:	11116.240:	1.34226	3.01159	-11.3467
15:	11116.229:	1.37860	3.10898	-13.5164
16:	11116.224:	1.41455	3.14663	-15.6880
17:	11116.221:	1.45113	3.15186	-17.8599
18:	11116.220:	1.48586	3.18124	-20.0316
19:	11116.219:	1.51994	3.20406	-22.2035
20:	11116.218:	1.55064	3.22386	-24.2324
21:	11116.218:	1.57972	3.24631	-26.2613
22:	11116.217:	1.60724	3.26437	-28.2902
23:	11116.217:	1.63097	3.28564	-30.0739
24:	11116.217:	1.65191	3.30750	-31.8577

25:	11116.216:	1.67476	3.34690	-34.5078
26:	11116.215:	1.68926	3.38050	-36.5249
27:	11116.215:	1.68666	3.39839	-37.2087
28:	11116.215:	1.69364	3.40859	-37.8926
29:	11116.215:	1.69991	3.41753	-38.5765
30:	11116.215:	1.70677	3.42223	-39.1666
31:	11116.215:	1.73242	3.43322	-41.1845
32:	11116.215:	1.74434	3.43038	-41.8849
33:	11116.215:	1.76468	3.43539	-43.4545
34:	11116.214:	1.78907	3.44632	-45.5732
35:	11116.214:	1.81430	3.45958	-47.9453
36:	11116.214:	1.82514	3.46566	-48.9606
37:	11116.214:	1.86235	3.48610	-52.5741
38:	11116.214:	1.85876	3.48527	-52.3932
39:	11116.214:	1.87752	3.49341	-54.1692
40:	11116.214:	1.88572	3.49664	-54.9450
41:	11116.214:	1.90242	3.50224	-56.4968
42:	11116.214:	1.91953	3.50653	-58.0486
43:	11116.214:	1.93706	3.50907	-59.6003
44:	11116.214:	1.95337	3.51374	-61.1521
45:	11116.214:	1.96914	3.51822	-62.7039

46:	11116.214:	1.98756	3.52451	-64.6271
47:	11116.214:	2.00629	3.53088	-66.5502
48:	11116.214:	2.01839	3.53556	-67.9328
49:	11116.214:	2.03148	3.53962	-69.3154
50:	11116.214:	2.04463	3.54299	-70.6979
51:	11116.214:	2.06259	3.54637	-72.5577
52:	11116.214:	2.08052	3.55034	-74.4175
53:	11116.214:	2.08561	3.55078	-74.9460
54:	11116.214:	2.10570	3.55444	-77.0600
55:	11116.214:	2.12038	3.55726	-78.6633
56:	11116.214:	2.13299	3.56012	-80.1147
57:	11116.214:	2.14597	3.56298	-81.5662
58:	11116.214:	2.15858	3.56589	-83.0177
59:	11116.214:	2.16736	3.56817	-84.0919
60:	11116.214:	2.17658	3.57007	-85.1662
61:	11116.214:	2.19976	3.57416	-87.8635
62:	11116.214:	2.21172	3.57556	-89.2594
63:	11116.214:	2.22380	3.57736	-90.6552
64:	11116.214:	2.23579	3.57908	-92.0511
65:	11116.214:	2.23585	3.57893	-92.0595

Here ML takes more iterations but **used to** give up with ‘singular convergence’ perhaps

because the estimated RE variance is closer to singular, which, combined with the larger number of parameter, leads 'nlminb' to declare singular convergence: the peak is too flat.

Note: increasing msMaxIter won't help with 'singular convergence'

REML maximizes only over RE parameters, while ML maximizes over all parameters, hence more challenging optimization

To see the fit at the last iteration, use 'returnObject = TRUE' in the 'control' list:

```
# fit.ml <- lme( mathach ~ ses * Sector, dd, random = ~ 1 + ses | id,  
#               na.action = na.exclude,  
#               method = "ML",  
#               control = list(msMaxIter=200,  
#                             msVerbose=T,  
#                             returnObject = TRUE))  
summary(fit.ml)
```

Linear mixed-effects model fit by maximum likelihood

Data: dd

AIC	BIC	logLik
12854.79	12899.51	-6419.397

Random effects:

Formula: ~1 + ses | id

Structure: General positive-definite, Log-Cholesky parametrization

	StdDev	Corr
(Intercept)	1.8032583	(Intr)
ses	0.1707538	0.932
Residual	6.1189789	

Fixed effects: mathach ~ ses * Sector

	Value	Std.Error	DF	t-value	p-value
(Intercept)	13.489366	0.4345233	1935	31.044061	0.0000
ses	1.815089	0.2821217	1935	6.433711	0.0000
SectorPublic	-1.590550	0.6400743	38	-2.484946	0.0175
ses:SectorPublic	1.398797	0.4234030	1935	3.303700	0.0010

Correlation:

	(Intr)	ses	SctrPb
ses	0.076		
SectorPublic	-0.679	-0.052	
ses:SectorPublic	-0.051	-0.666	0.149

Standardized Within-Group Residuals:

Min	Q1	Med	Q3	Max
-----	----	-----	----	-----

-3.06217668 -0.74613002 0.03736602 0.78804322 2.74091602

Number of Observations: 1977

Number of Groups: 40

Shows high correlation in the RE model between b_1 and b_ses . This suggests that the ‘point of minimum variance’ is far from the observed values of ses and that the random true school regression lines seem close to parallel.

Possible remedies for non-convergence:

Alas now ‘academic’ here because the model now converges

- drop random slope
- recenter ses
- add a contextual effect and CWG random effect: $\sim 1 + dvar(ses, id) \mid id$
- explore different models
- change the criterion for lme to declare singularity, i.e. excessive flatness near the likelihood peak with ‘sing.tol’

```
fit.ml <- lme( mathach ~ ses * Sector + Sex, dd,  
              random = ~ 1 + ses + Sex | id,  
              na.action = na.exclude,  
              method = "ML",
```

```
control = list(msMaxIter=200, msVerbose=T,  
               sing.tol=1e-20))
```

0:	11109.615:	1.41833	2.41824	1.88935	-3.96692	-0.973608	-1.36731
1:	11109.477:	1.44252	2.70884	2.15340	-3.97460	-0.990959	-1.37504
2:	11109.400:	1.35156	2.71273	2.13702	-3.97739	-0.998252	-1.37783
3:	11109.376:	1.36793	2.79269	2.14010	-3.99193	-1.03470	-1.39820
4:	11109.362:	1.40683	2.80828	2.01850	-4.03354	-1.14572	-1.46064
5:	11109.320:	1.42023	2.99532	2.08100	-4.13910	-1.43142	-1.62290
6:	11109.285:	1.43114	3.10075	2.08836	-4.24835	-1.75102	-1.80401
7:	11109.185:	1.48413	3.36028	2.12353	-4.68678	-3.05622	-2.55469
8:	11109.123:	1.53966	3.48380	2.21745	-5.12894	-4.36430	-3.32730
9:	11109.076:	1.59585	3.55853	2.24802	-5.56928	-5.67298	-4.11118
10:	11109.021:	1.71368	3.55344	2.39068	-6.49460	-8.40163	-5.77104
11:	11108.954:	1.81242	3.40201	2.40801	-7.25146	-10.6234	-7.15512
12:	11108.937:	1.90548	3.27424	2.47237	-8.01006	-12.8474	-8.53624
13:	11108.932:	1.90942	3.30006	2.46637	-8.09936	-13.0958	-8.69635
14:	11108.923:	1.93768	3.37975	2.48650	-8.44823	-14.1038	-9.32179
15:	11108.912:	2.04458	3.52641	2.55840	-9.63736	-17.5463	-11.4635
16:	11108.905:	2.05586	3.54962	2.56007	-9.83938	-18.1168	-11.8326
17:	11108.899:	2.11950	3.62610	2.59167	-10.6409	-20.4159	-13.2863
18:	11108.894:	2.18970	3.69114	2.62024	-11.5585	-23.0377	-14.9574

19:	11108.889:	2.24429	3.73449	2.63533	-12.3256	-25.2168	-16.3618
20:	11108.887:	2.30215	3.79270	2.65515	-13.0915	-27.4009	-17.7587
21:	11108.885:	2.34188	3.83961	2.66230	-13.6605	-29.0131	-18.8015
22:	11108.883:	2.39691	3.90956	2.66522	-14.4695	-31.2977	-20.2890
23:	11108.882:	2.45426	3.97424	2.67403	-15.2774	-33.5868	-21.7705
24:	11108.880:	2.50176	4.02582	2.67524	-15.9609	-35.5165	-23.0288
25:	11108.879:	2.55864	4.08471	2.66951	-16.7983	-37.8728	-24.5771
26:	11108.878:	2.61557	4.14578	2.67017	-17.6350	-40.2328	-26.1198
27:	11108.878:	2.64907	4.18248	2.66874	-18.1610	-41.7119	-27.0935
28:	11108.877:	2.69285	4.23159	2.66654	-18.9217	-43.8454	-28.5079
29:	11108.876:	2.73814	4.28140	2.66685	-19.6820	-45.9818	-29.9180
30:	11108.876:	2.77826	4.32515	2.66935	-20.4426	-48.1145	-31.3339
31:	11108.875:	2.81108	4.36009	2.67593	-21.2034	-50.2423	-32.7574
32:	11108.874:	2.86012	4.41024	2.68995	-22.4337	-53.6809	-35.0625
33:	11108.874:	2.87997	4.42669	2.71002	-23.3420	-56.2078	-36.7800
34:	11108.873:	2.91449	4.46188	2.72112	-24.2499	-58.7441	-38.4832
35:	11108.873:	2.93380	4.48241	2.72586	-24.7517	-60.1447	-39.4251
36:	11108.873:	2.93473	4.48449	2.72325	-24.7335	-60.0938	-39.3907
37:	11108.873:	2.97031	4.52788	2.71283	-25.3758	-61.8861	-40.5955
38:	11108.873:	3.03861	4.60421	2.70580	-26.8211	-65.9184	-43.3097
39:	11108.873:	3.03513	4.59758	2.70602	-26.7677	-65.7680	-43.2114

40:	11108.873:	3.03679	4.59937	2.70571	-26.8211	-65.9168	-43.3123
41:	11108.872:	3.05342	4.61500	2.70623	-27.2488	-67.1083	-44.1186
42:	11108.872:	3.11825	4.67979	2.70999	-28.9597	-71.8741	-47.3440
43:	11108.872:	3.11142	4.67338	2.71520	-28.9620	-71.8786	-47.3515
44:	11108.872:	3.11117	4.67349	2.71485	-28.9660	-71.8897	-47.3590
45:	11108.872:	3.13222	4.69632	2.71875	-29.6371	-73.7580	-48.6261
46:	11108.872:	3.15772	4.72426	2.72096	-30.3932	-75.8633	-50.0525
47:	11108.872:	3.18344	4.75266	2.72209	-31.1492	-77.9688	-51.4787
48:	11108.872:	3.20928	4.78158	2.72176	-31.9053	-80.0744	-52.9047
49:	11108.871:	3.24659	4.82278	2.71914	-32.9989	-83.1206	-54.9671
50:	11108.871:	3.28239	4.85920	2.71907	-34.0924	-86.1663	-57.0305
51:	11108.871:	3.31604	4.89126	2.71866	-35.1859	-89.2115	-59.0947
52:	11108.871:	3.34442	4.91913	2.72002	-36.1901	-92.0077	-60.9915
53:	11108.871:	3.36973	4.94546	2.72113	-37.0776	-94.4788	-62.6675
54:	11108.871:	3.39259	4.96998	2.72239	-37.9650	-96.9493	-64.3443
55:	11108.871:	3.41683	4.99579	2.72332	-38.8525	-99.4202	-66.0207
56:	11108.871:	3.44020	5.02127	2.72401	-39.7399	-101.891	-67.6972
57:	11108.871:	3.47088	5.05549	2.72465	-40.9619	-105.293	-70.0064
58:	11108.871:	3.50130	5.08730	2.72501	-42.1741	-108.668	-72.2971
59:	11108.870:	3.54083	5.12735	2.72505	-43.8354	-113.293	-75.4378
60:	11108.870:	3.58166	5.16897	2.72549	-45.4969	-117.919	-78.5780

61:	11108.870:	3.57103	5.15775	2.72541	-45.0932	-116.795	-77.8153
62:	11108.870:	3.58030	5.16713	2.72553	-45.4968	-117.919	-78.5785
63:	11108.870:	3.61731	5.20512	2.72598	-47.1114	-122.414	-81.6312
64:	11108.870:	3.63540	5.22371	2.72626	-47.9463	-124.738	-83.2102
65:	11108.870:	3.65738	5.24653	2.72665	-49.0452	-127.797	-85.2895
66:	11108.870:	3.68094	5.27097	2.72697	-50.1442	-130.856	-87.3681
67:	11108.870:	3.70419	5.29512	2.72724	-51.2432	-133.916	-89.4468
68:	11108.870:	3.70382	5.29489	2.72714	-51.2391	-133.904	-89.4392
69:	11108.870:	3.70372	5.29509	2.72680	-51.2499	-133.934	-89.4596
70:	11108.870:	3.72505	5.31734	2.72698	-52.3046	-136.870	-91.4551
71:	11108.870:	3.73523	5.32795	2.72702	-52.8152	-138.291	-92.4212
72:	11108.870:	3.75534	5.34876	2.72713	-53.8363	-141.134	-94.3533
73:	11108.870:	3.77507	5.36911	2.72724	-54.8574	-143.976	-96.2855
74:	11108.870:	3.79415	5.38864	2.72735	-55.8785	-146.819	-98.2178
75:	11108.870:	3.81913	5.41386	2.72772	-57.2752	-150.706	-100.861
76:	11108.870:	3.84466	5.43989	2.72809	-58.6719	-154.594	-103.505
77:	11108.870:	3.85358	5.44893	2.72833	-59.2238	-156.130	-104.549
78:	11108.870:	3.89183	5.48813	2.72887	-61.4314	-162.275	-108.728
79:	11108.870:	3.89233	5.48890	2.72878	-61.5124	-162.500	-108.882
80:	11108.870:	3.91404	5.51138	2.72888	-62.8092	-166.109	-111.337
81:	11108.870:	3.92430	5.52201	2.72888	-63.4305	-167.839	-112.513

82:	11108.870:	3.94457	5.54292	2.72886	-64.6732	-171.297	-114.865
83:	11108.870:	3.96442	5.56338	2.72879	-65.9159	-174.756	-117.218
84:	11108.870:	3.99363	5.59335	2.72864	-67.8061	-180.017	-120.797
85:	11108.870:	4.02246	5.62257	2.72898	-69.6964	-185.278	-124.376
86:	11108.870:	4.04654	5.64655	2.72954	-71.4205	-190.076	-127.642
87:	11108.870:	4.07169	5.67218	2.72980	-73.1446	-194.874	-130.907
88:	11108.870:	4.06953	5.67031	2.72970	-73.0381	-194.578	-130.705
89:	11108.870:	4.08436	5.68574	2.72974	-74.1011	-197.536	-132.719
90:	11108.870:	4.09548	5.69720	2.72976	-74.8991	-199.757	-134.230
91:	11108.870:	4.11748	5.71981	2.72981	-76.4953	-204.199	-137.253
92:	11108.870:	4.13498	5.73779	2.72978	-77.8038	-207.841	-139.732
93:	11108.870:	4.15109	5.75413	2.72975	-79.0754	-211.380	-142.141
94:	11108.870:	4.16779	5.77108	2.72982	-80.3471	-214.919	-144.549
95:	11108.870:	4.18428	5.78781	2.72992	-81.6187	-218.457	-146.958
96:	11108.870:	4.18484	5.78823	2.73003	-81.6855	-218.643	-147.085
97:	11108.870:	4.18716	5.79064	2.72995	-81.8857	-219.200	-147.464
98:	11108.870:	4.20695	5.81048	2.73027	-83.4878	-223.659	-150.499
99:	11108.870:	4.21345	5.81712	2.73028	-84.0168	-225.131	-151.502
100:	11108.870:	4.22635	5.83037	2.73025	-85.0748	-228.075	-153.506
101:	11108.870:	4.25192	5.85664	2.73020	-87.1908	-233.963	-157.515
102:	11108.870:	4.27673	5.88252	2.72986	-89.3067	-239.852	-161.524

103:	11108.870:	4.28524	5.89112	2.72998	-90.1149	-242.100	-163.055
104:	11108.870:	4.31403	5.92022	2.73029	-92.6455	-249.142	-167.850
105:	11108.870:	4.32456	5.93065	2.73053	-93.6039	-251.809	-169.666
106:	11108.870:	4.33436	5.93987	2.73109	-94.5623	-254.476	-171.483
107:	11108.870:	4.35277	5.95848	2.73121	-96.2805	-259.257	-174.739
108:	11108.870:	4.37086	5.97729	2.73092	-97.9988	-264.039	-177.995
109:	11108.870:	4.38919	5.99615	2.73079	-99.7171	-268.820	-181.251
110:	11108.870:	4.39342	6.00062	2.73068	-100.139	-269.994	-182.050
111:	11108.870:	4.42375	6.03185	2.73045	-103.092	-278.210	-187.646
112:	11108.870:	4.42003	6.02787	2.73060	-102.791	-277.374	-187.077
113:	11108.870:	4.43195	6.03990	2.73068	-103.993	-280.718	-189.355
114:	11108.870:	4.43902	6.04707	2.73071	-104.711	-282.716	-190.716
115:	11108.870:	4.45305	6.06125	2.73079	-106.147	-286.711	-193.437
116:	11108.870:	4.46681	6.07505	2.73093	-107.583	-290.706	-196.158
117:	11108.870:	4.47692	6.08503	2.73113	-108.687	-293.779	-198.251
118:	11108.870:	4.49525	6.10365	2.73116	-110.647	-299.232	-201.966
119:	11108.870:	4.51301	6.12197	2.73102	-112.606	-304.684	-205.680
120:	11108.870:	4.53107	6.14042	2.73098	-114.566	-310.137	-209.394
121:	11108.870:	4.53070	6.14001	2.73098	-114.554	-310.105	-209.373
122:	11108.870:	4.53074	6.14011	2.73092	-114.566	-310.137	-209.395
123:	11108.870:	4.54388	6.15342	2.73096	-116.044	-314.248	-212.195

124:	11108.870:	4.54910	6.15872	2.73095	-116.635	-315.893	-213.316
125:	11108.870:	4.55943	6.16922	2.73092	-117.817	-319.182	-215.556
126:	11108.870:	4.57993	6.19001	2.73092	-120.181	-325.760	-220.038
127:	11108.870:	4.59575	6.20608	2.73083	-122.064	-331.000	-223.608
128:	11108.870:	4.60914	6.21952	2.73098	-123.696	-335.540	-226.701
129:	11108.870:	4.62805	6.23841	2.73134	-126.095	-342.214	-231.249
130:	11108.870:	4.64748	6.25788	2.73166	-128.494	-348.890	-235.798
131:	11108.870:	4.65512	6.26588	2.73141	-129.446	-351.538	-237.602
132:	11108.870:	4.67425	6.28550	2.73119	-131.846	-358.215	-242.152
133:	11108.870:	4.69170	6.30388	2.73035	-134.085	-364.445	-246.396
134:	11108.870:	4.70907	6.32157	2.73021	-136.324	-370.674	-250.641
135:	11108.870:	4.70524	6.31710	2.73082	-135.913	-369.533	-249.863
136:	11108.870:	4.71413	6.32547	2.73148	-137.181	-373.060	-252.267
137:	11108.870:	4.72446	6.33566	2.73181	-138.605	-377.022	-254.968
138:	11108.870:	4.73464	6.34583	2.73208	-140.029	-380.985	-257.668
139:	11108.870:	4.74476	6.35618	2.73214	-141.454	-384.947	-260.368
140:	11108.870:	4.76631	6.37842	2.73178	-144.393	-393.126	-265.942
141:	11108.869:	4.76523	6.37779	2.73120	-144.227	-392.664	-265.627
142:	11108.869:	4.76643	6.37901	2.73119	-144.393	-393.126	-265.942

summary(fit.ml)

Linear mixed-effects model fit by maximum likelihood

Data: dd

	AIC	BIC	logLik
	12848.11	12915.18	-6412.053

Random effects:

Formula: ~1 + ses + Sex | id

Structure: General positive-definite, Log-Cholesky parametrization

	StdDev	Corr
(Intercept)	1.5507024	(Intr) ses
ses	0.1796016	0.998
SexMale	0.3973486	0.999 0.998
Residual	6.0997280	

Fixed effects: mathach ~ ses * Sector + Sex

	Value	Std.Error	DF	t-value	p-value
(Intercept)	12.951009	0.4218649	1934	30.699421	0.0000
ses	1.830092	0.2807716	1934	6.518079	0.0000
SectorPublic	-1.618970	0.6103963	38	-2.652326	0.0116
SexMale	1.173682	0.3321072	1934	3.534045	0.0004
ses:SectorPublic	1.344947	0.4212879	1934	3.192466	0.0014

Correlation:

	(Intr)	ses	SctrPb	SexMal
ses		0.079		
SectorPublic		-0.655	-0.058	
SexMale		-0.220	0.019	-0.012
ses:SectorPublic		-0.048	-0.667	0.169 -0.032

Standardized Within-Group Residuals:

	Min	Q1	Med	Q3	Max
	-3.06009142	-0.75212643	0.04604295	0.79450300	2.66713374

Number of Observations: 1977

Number of Groups: 40

Notes on testing: ML vs REML —

The default for lme is fitting with ‘Residual Maximum Likelihood’ (REML) – it has other interpretations but everyone agrees on ‘REML’ – With REML, maximum likelihood is applied ONLY to the parameters for random effects, i.e. the G matrix and sigma (or R). The Betas play no direct role, they are only estimated with Generalized Least Squares as an afterthought. This means that the likelihood with REML is only informative for the the RE model: the G

and R matrices. You can't compare two models that have different FE models.

Full maximum likelihood (ML) does look at Betas, G and R together. So the likelihood with ML can be used to test two models with nested FE models.

The consequence is that:

If two models have been fitted with REML, they must have identical FE models to compare them with 'anova': p-values, AIC, BIC.

If two models have been fitted with ML, you can use 'anova' regardless of the FE or the RE models — PROVIDED they have the same response variable. This includes requiring that they have the same set of observations, hence the same omitted rows due to missing values.

Wald tests can be used for both ML or REML fits. They might be more accurate with REML fits. However, Wald tests tend to be useful mainly for the FE part of the model. 'wald' in 'spida2' only works for the FE model. To compare two RE models (with identical FE models) LRT is generally better than Wald. However, classic p-values tend to be too large in many cases and can be adjusted through simulation.

Summary:

- Wald is ok for the FE model whether ML or REML
- anova (with simulation adjustment) is ok for the RE model provided the FE models are

identical

- anova can be used to test different FE models or models that differ in both FEs and REs ONLY IF THE MODELS ARE fitted with ML.

Diagnostics —

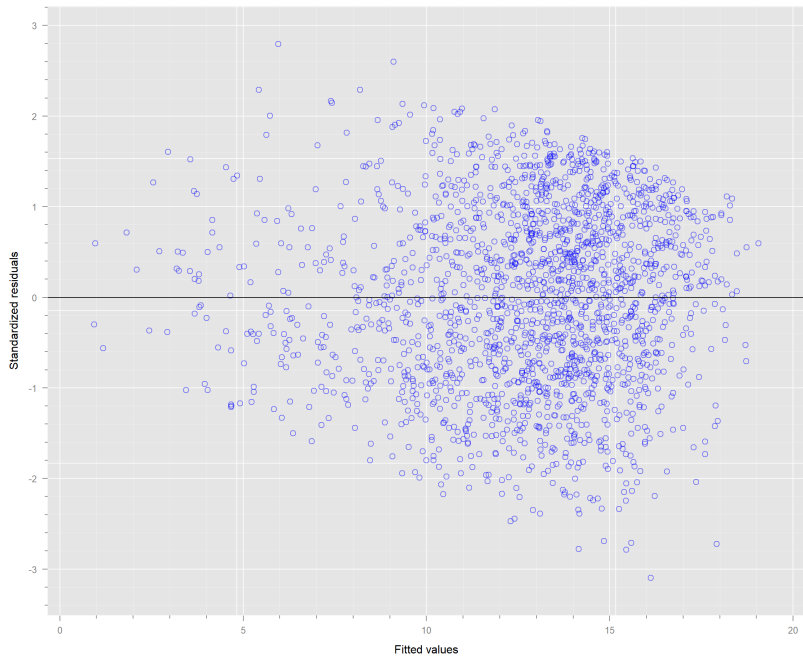
The ‘car’ package has methods to conduct influence diagnostics with a wide range of models include ‘lme’ models. I plan to incorporate a discussion of these methods soon.

Diagnostics with Level 1 residuals —

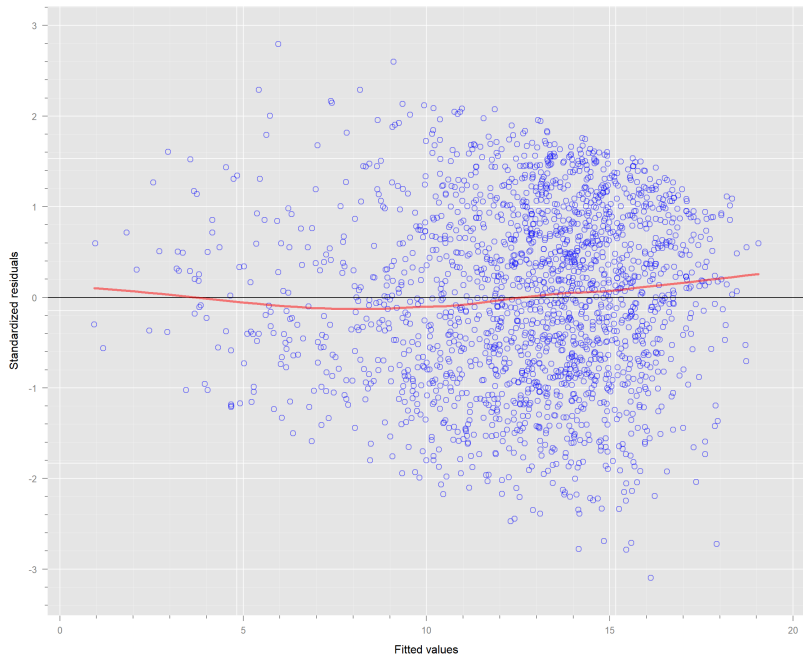
Level 1 residuals

Level 1 standardized residuals versus yhat:

```
gd(pch=1, alpha = .5)  
plot(fitc)
```



```
plot(fitc) + layer(panel.loess(x,y, lwd=3,col='red'))
```



Not as generous as plot for 'lm'.

In contrast with least-squares, it is possible to have a non-flat least-squares regression here.

```
diag.df <- data.frame( resid = resid(fitc), fitted = fitted(fitc))  
summary( lm( resid ~ fitted, diag.df))
```

Call:

```
lm(formula = resid ~ fitted, data = diag.df)
```

Residuals:

Min	1Q	Median	3Q	Max
-19.1659	-4.5557	0.1715	4.7476	17.5492

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	-0.84287	0.60009	-1.405	0.160
fitted	0.06594	0.04572	1.442	0.149

Residual standard error: 6.056 on 1975 degrees of freedom

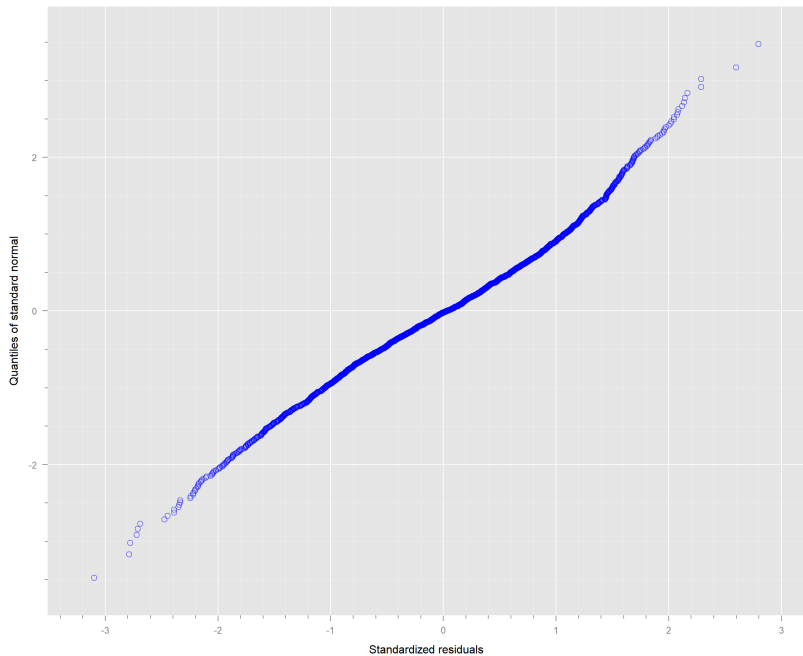
Multiple R-squared: 0.001052, Adjusted R-squared: 0.0005463

F-statistic: 2.08 on 1 and 1975 DF, p-value: 0.1494

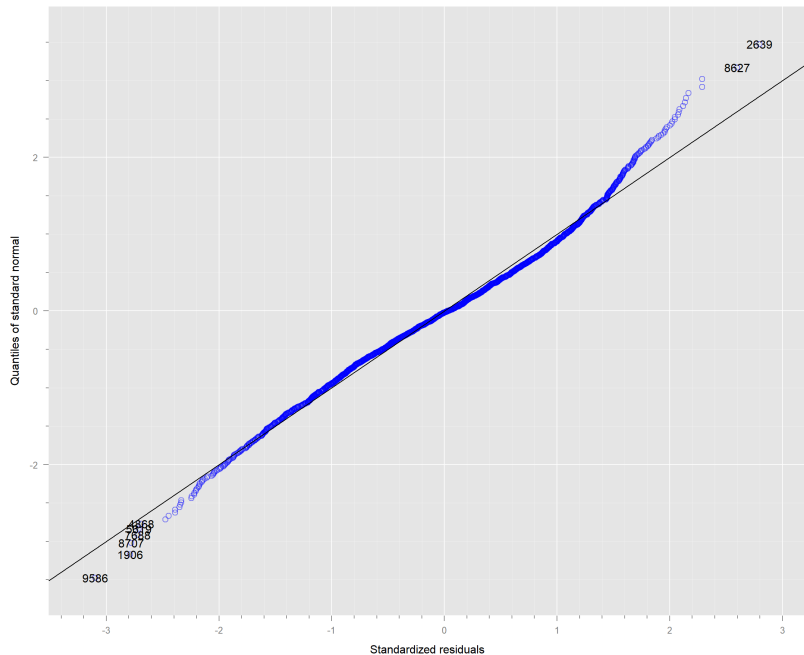
It could be a symptom of lack of independence between school-level random effects and level 1 errors or between school-level random effects and other predictors. One possible cure is adding a contextual effect if there isn't one.

Normality:

```
qqnorm( fitc )
```



```
qqnorm( fitc , abline = c(0,1), id = .01 )
```

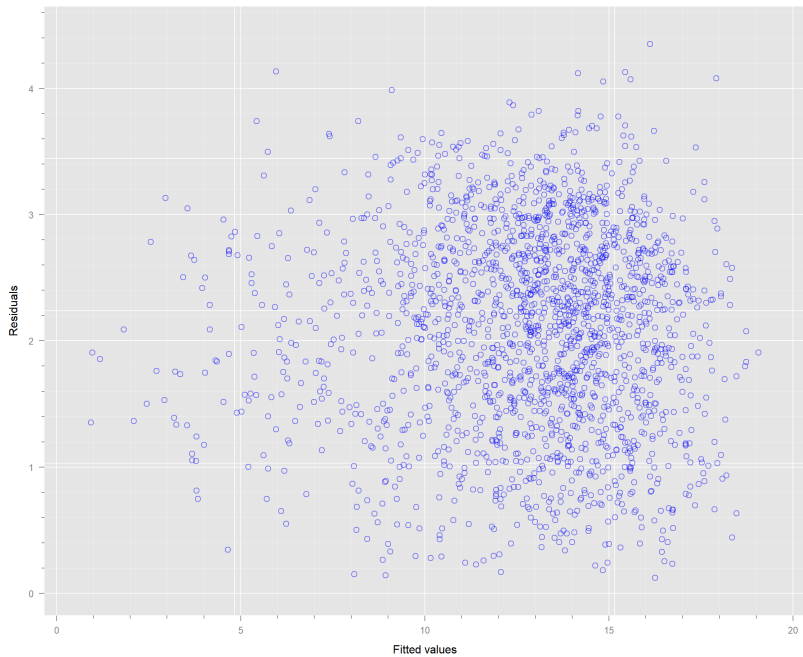


BEWARE: data on horizontal axis SO distribution is slightly platykurtic. Variance might be overestimated with extreme values thereby overadjusting.

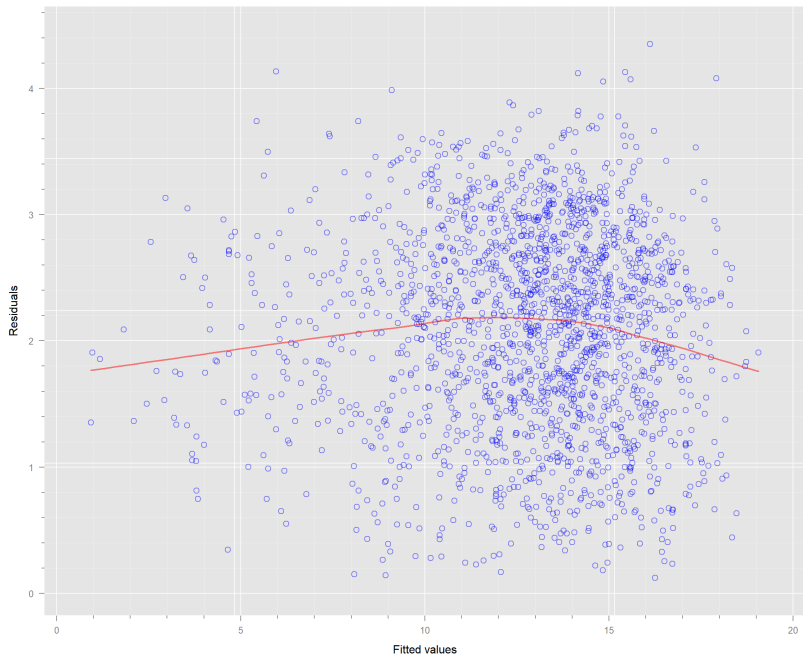
Scale - location plot: —

Using `sqrt(abs(resid))` produces an approximate normal if everything is ok

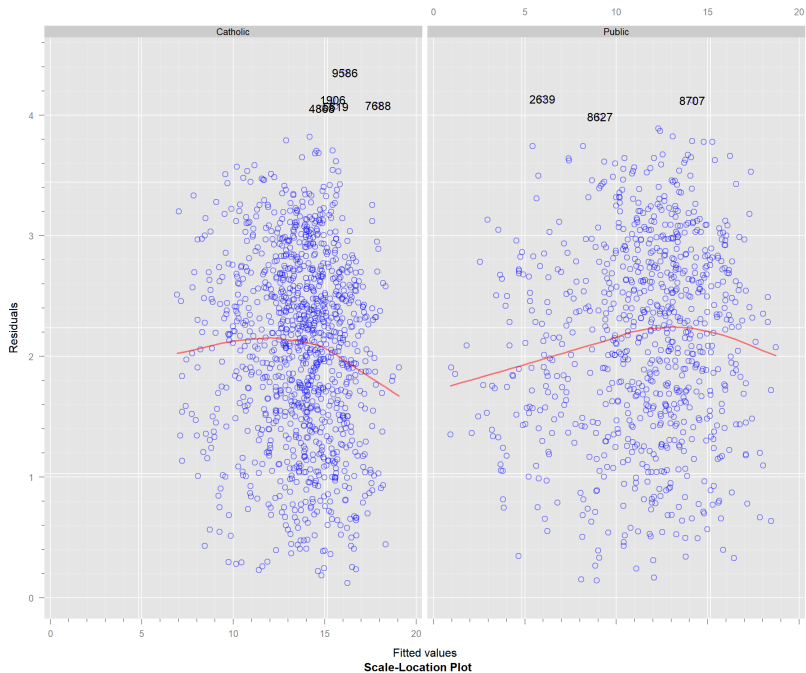
```
plot( fitc , sqrt( abs( resid(.)) ) ~ fitted(.))
```



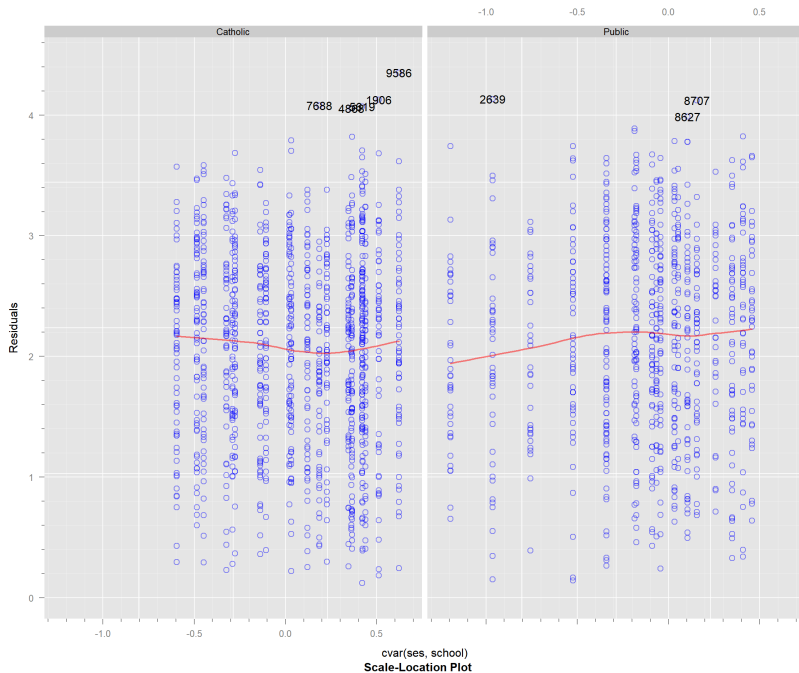
```
plot( fitc , sqrt( abs( resid(.))) ~ fitted(.)) +  
  layer(panel.loess(x, y, lwd = 2, col = 'red'))
```



```
plot( fitc ,  
      sqrt( abs( resid(.))) ~ fitted(.)| Sector,  
      id = .01, sub = "Scale-Location Plot") +  
layer(panel.loess(x, y, lwd = 2, col = 'red'))
```

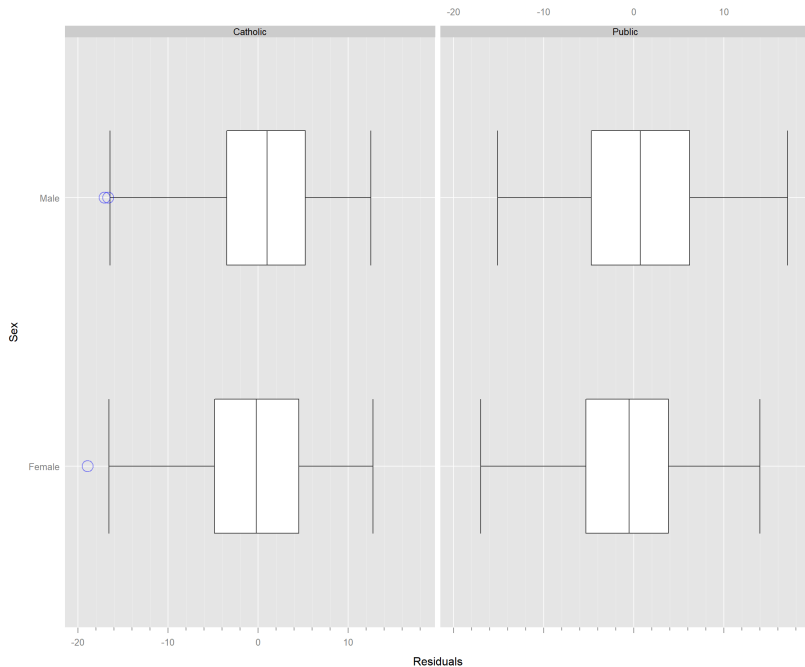


```
plot( fitc , sqrt( abs( resid(.))) ~ cvar(ses,school)| Sector,  
      id = .01, sub = "Scale-Location Plot") +  
  layer(panel.loess(x, y, lwd = 2, col = 'red'))
```

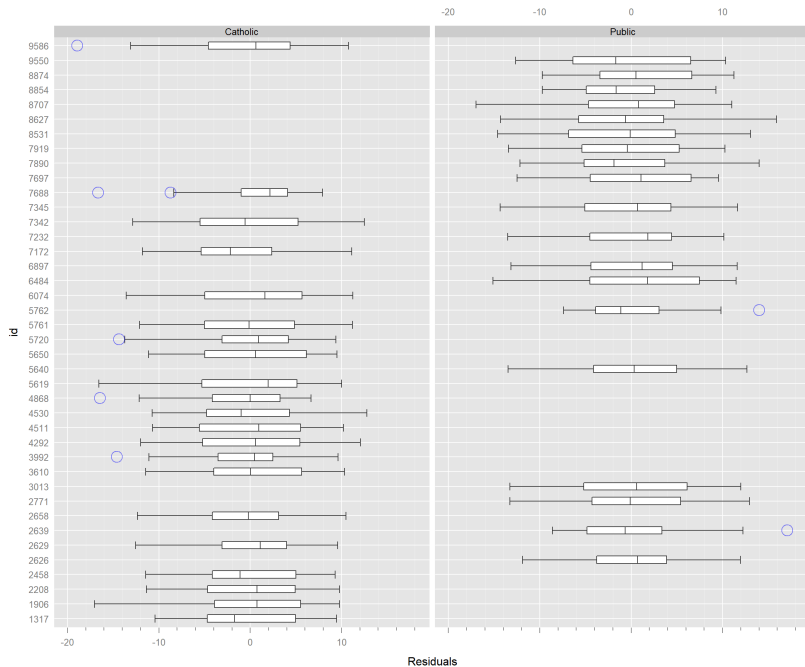


other views of residuals:

```
plot( fitc, Sex ~ resid(.) | Sector)
```

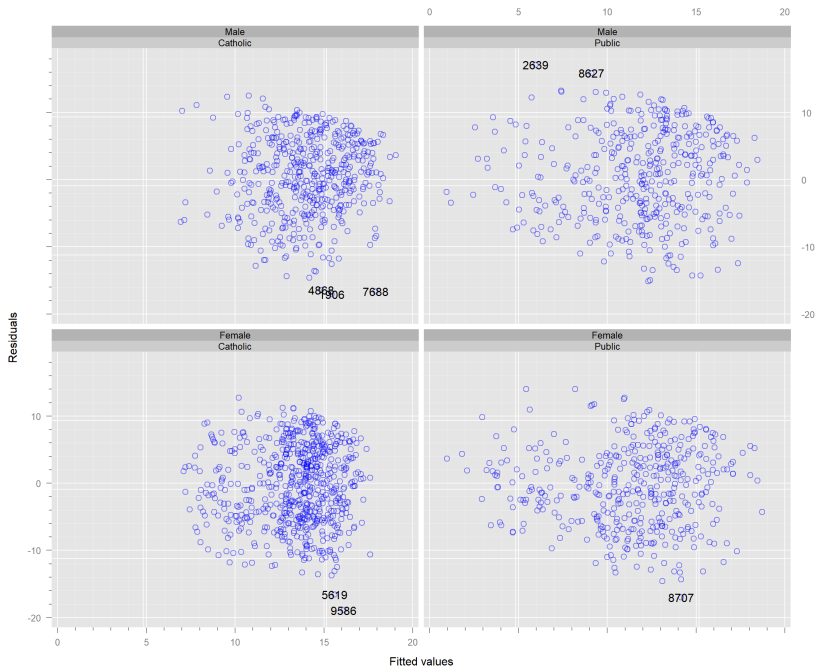


```
plot( fitc, id ~ resid(.) | Sector)
```



Omitted variables:

```
plot( fitc , resid(.) ~ fitted(.) | Sector*Sex, id = .01)
```



Why MM residual plots don't look like OLS residual plots:

Question: Performing OLS regression diagnostics, you find that the residuals are correlated with the predicted values. What would this signify and what action should you take?

Answer: at end of script.

With MM residuals:

Why are residuals not uncorrelated with predicted values (a canon of OLS)?

Reason: if we were using OLS on the pooled data, residuals would be uncorrelated.

But MMs is roughly like doing OLS on pooled data, then adding an additional fit at the cluster level.

If unaccounted between cluster effects are in the same direction as within cluster effect then higher residuals with higher fitted values will get a second fit, This shrinks the high positive OLS residuals and stretches the corresponding fitted values creating a negative slope in the residual vs fitted plot.

Diagnostics with Level 2 residuals —

Make `cvar(ses,id)` a variable in `dd`

```
dd$ses.m <- with( dd, cvar( ses, id))
fitc <- lme( mathach ~ (ses + ses.m) * Sector, dd,
            random = ~ 1 + ses | id )
```

- 'ranef(fitc)' produces Level 2 errors: u in each school
- 'coef(fitc)' produces Level 2 BLUPS: $\beta = \gamma + u$

```
some( coef ( fitc ) )      # BLUP in each cluster
```

	(Intercept)	ses	ses.m	SectorPublic	ses:SectorPublic
2208	13.53819	1.817657	2.407785	-0.934345	1.173092
2626	14.30123	1.714866	2.407785	-0.934345	1.173092
3610	14.49506	1.824884	2.407785	-0.934345	1.173092
3992	13.25713	1.564306	2.407785	-0.934345	1.173092
4292	14.42899	1.294163	2.407785	-0.934345	1.173092
5762	13.17761	1.514336	2.407785	-0.934345	1.173092
6074	14.52082	1.565613	2.407785	-0.934345	1.173092
6484	14.51768	1.368009	2.407785	-0.934345	1.173092
7890	12.86207	1.393502	2.407785	-0.934345	1.173092
9586	12.56447	1.694839	2.407785	-0.934345	1.173092
ses.m:SectorPublic					
2208		1.414896			
2626		1.414896			

3610	1.414896
3992	1.414896
4292	1.414896
5762	1.414896
6074	1.414896
6484	1.414896
7890	1.414896
9586	1.414896

```
some( ranef ( fitc ) )      # RE in each cluster
```

	(Intercept)	ses
1906	0.36252477	0.050644237
2208	0.18213282	0.118259831
3013	0.35388133	0.046187778
3610	1.13900591	0.125486781
4868	-1.63966343	0.004294988
5619	0.02874504	0.502254674
5761	-0.63626758	0.270788614
6484	1.16162333	-0.331388396
7172	-2.88069356	0.202229471
8531	-1.15715031	0.079146909

Note: `coef( fitc ) = fixed.effect + random.effect pairs( cbind( coef(fitc), ranef( fitc)) )` # can you interpret what this says?

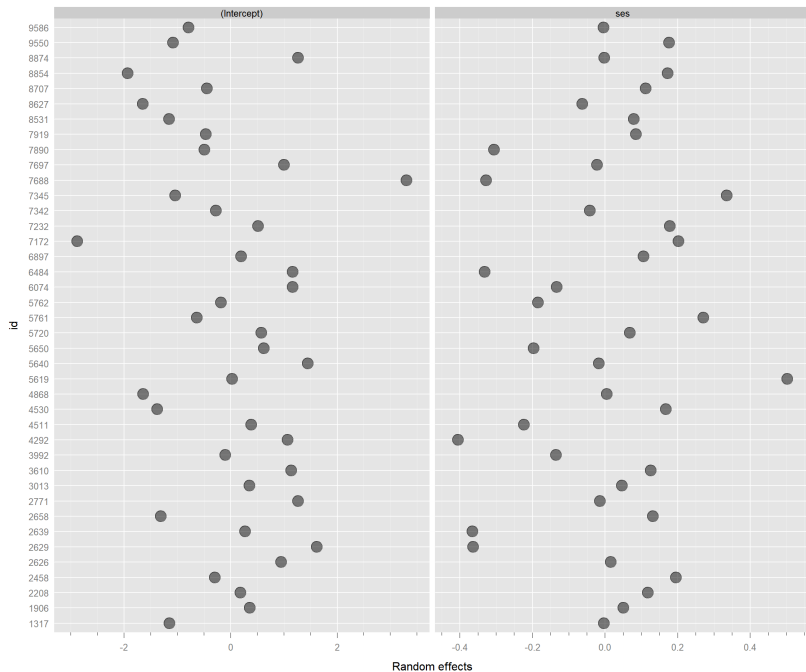
```
re <- ranef( fitc, aug = T)    # creates data frame with up( dd, ~ school) variables
some( re )
```

	(Intercept)	ses	school	mathach	Sex	Minority	Size
1317	-1.1509678	-0.003523084	1317	13.177687	Female	Yes	455
2658	-1.3121006	0.131984540	2658	13.396156	Female	No	780
4292	1.0729371	-0.405233961	4292	12.864354	Male	Yes	1328
6074	1.1647641	-0.133784197	6074	13.779089	Female	No	2051
6484	1.1616233	-0.331388396	6484	12.912400	Female	No	726
6897	0.1989012	0.106243527	6897	15.097633	Female	No	1415
7172	-2.8806936	0.202229471	7172	8.066818	Female	Yes	280
7342	-0.2737076	-0.042255077	7342	11.166414	Male	No	1220
8707	-0.4475834	0.112155798	8707	12.883938	Female	No	1133
9550	-1.0779843	0.176773458	9550	11.089138	Female	No	1532

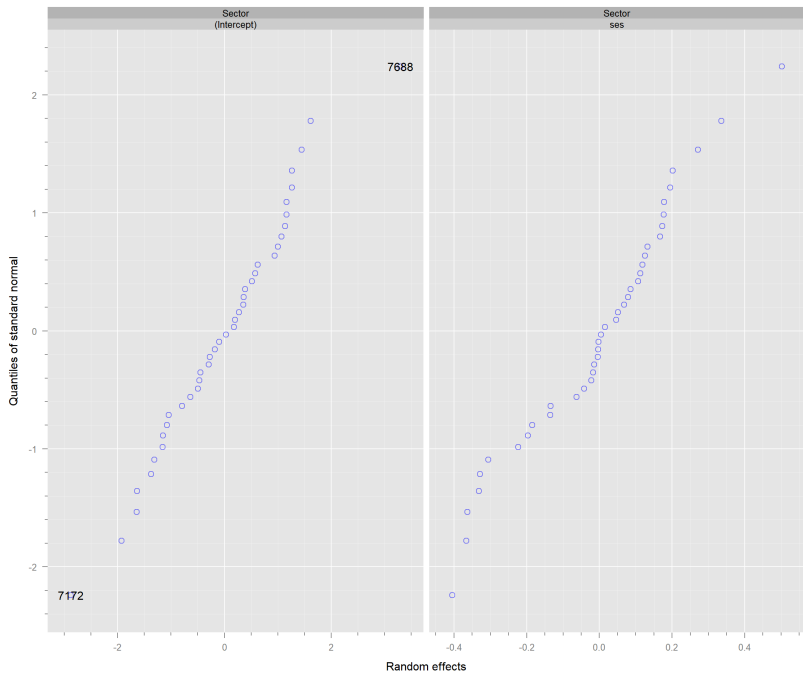
	Sector	PRACAD	DISCLIM	n	ses.m	ses.cwg	ses.iqr
1317	Catholic	0.95	-1.694	48	0.34533333	1.502975e-17	0.7825
2658	Catholic	0.79	-0.961	45	0.43844444	-9.250654e-18	0.9100
4292	Catholic	0.76	-0.674	65	-0.48615385	-5.096167e-18	0.8600
6074	Catholic	0.32	-1.018	56	-0.27675000	-1.487390e-18	0.8175

6484	Public	0.19	0.218	35	-0.18514286	-2.762392e-18	0.7400
6897	Public	0.55	-0.361	49	0.34955102	-1.359125e-17	1.0700
7172	Catholic	0.05	1.013	44	-0.28981818	-3.027573e-17	1.0275
7342	Catholic	0.46	0.380	58	-0.44782759	-4.974128e-19	0.7125
8707	Public	0.48	1.542	48	0.15512500	8.673617e-19	1.1350
9550	Public	0.45	0.791	29	0.05303448	9.077389e-18	1.0200
	ses.trange		ses.sd	ses.rank	minority.diff	id.sec	ses.cvar
1317	1.166	0.5561583		24.5	-0.2676044	C1317	0.34533333
2658	1.698	0.6402846		23.0	-0.1830556	C2658	0.43844444
4292	1.708	0.6511382		33.0	-0.0150000	C4292	-0.48615385
6074	1.440	0.6271235		28.5	-0.3279630	C6074	-0.27675000
6484	1.932	0.6958345		18.0	-0.5227273	P6484	-0.18514286
6897	2.090	0.7445231		25.0	-0.7075000	P6897	0.34955102
7172	1.686	0.6764417		22.5	0.3915447	C7172	-0.28981818
7342	1.323	0.5648459		29.5	0.1027413	C7342	-0.44782759
8707	2.169	0.8042577		24.5	-0.5873333	P8707	0.15512500
9550	1.824	0.7847035		15.0	-0.5623913	P9550	0.05303448

```
plot( re )      # special plot method
```



```
qqnorm( fitc, ~ ranef(.) | Sector, id = .05)
```



```

if(interactive) {
  Init3d()
  Plot3d( '(Intercept)' ~ ses + ses.m | Sector, re)  # note funny names in backg
  Ell3d()
}

```

EXERCISE:

- Plot random effects against variables you feel might reveal a pattern
- Experiment with the alternative RE model.

Influence diagnostics – drop one row or one cluster at a time —

Here's a quick dropone function

```

dropone <- function(fit) {
  data <- getData(fit)
  index <- 1:nrow(data)
  lapply(index, function(i) fixef(update(fit, data=data[-i,])))
}

```

Here's a fancier one that will drop clusters and return NA if there's an error in fitting

```

dropone <- function(fit, form = NULL, FUN = fixef, data = getData(fit),...) {
  if(is.null(form)) {
    by <- factor(1:nrow(data))
    data$by <- by
    dframe <- data
  } else {
    by <- model.frame(form, data)
    by <- do.call(paste, c(by, sep='/'))
    data$by <- factor(by) # bad if by already exists in data
    dframe <- up(data, form)
  }
  values <- dframe$by
  names(values) <- values
  ret <- lapply(values, function(v) {
    ret <- try(update(fit, data = data[by != v, ,drop=FALSE],...))
    if(class(ret) == 'try-error') NA else FUN(ret)
  })
  ret <- do.call(rbind,ret)
  colnames(ret) <- paste0('b_',colnames(ret))
  cbind(dframe, ret)
}

```

Ideas for improvement:

- optionally standardize relative to SEs

The following takes about 5 minutes so it's been precooked so we can take it out of the oven.

```
# system.time(  
#   fits.dropone <- dropone(fitc)  
# )
```

Take fits.dropone out of the oven:

```
con <- url('http://blackwell.math.yorku.ca/ICPSR/fits.dropone.rda')  
load(con, verbose = T)
```

Loading objects:

fits.dropone

```
close(con)
```

```
str(fits.dropone)
```

```
'data.frame':    1977 obs. of  26 variables:  
 $ school      : int  1317 1317 1317 1317 1317 1317 1317 1317 1317 1317 13  
 $ mathach     : num  12.86 8.96 4.76 21.41 20.75 ...  
 $ ses        : num  0.882 0.932 -0.158 0.362 1.372 ...
```

```

$ Sex          : Factor w/ 2 levels "Female","Male": 1 1 1 1 1 1 1 1 1
$ Minority     : Factor w/ 2 levels "No","Yes": 1 2 2 2 1 2 1 2 2 2 .
$ Size        : int   455 455 455 455 455 455 455 455 455 455 ...
$ Sector       : Factor w/ 2 levels "Catholic","Public": 1 1 1 1 1 1
$ PRACAD       : num   0.95 0.95 0.95 0.95 0.95 0.95 0.95 0.95 0.95 0.
$ DISCLIM     : num  -1.69 -1.69 -1.69 -1.69 -1.69 ...
$ n           : int   48 48 48 48 48 48 48 48 48 48 ...
$ ses.m       : num   0.345 0.345 0.345 0.345 0.345 ...
$ ses.cwg     : num   0.5367 0.5867 -0.5033 0.0167 1.0267 ...
$ ses.iqr     : num   0.782 0.782 0.782 0.782 0.782 ...
$ ses.trange  : num   1.17 1.17 1.17 1.17 1.17 ...
$ ses.sd      : num   0.556 0.556 0.556 0.556 0.556 ...
$ ses.rank    : num   39 40 6 25 47 20 20 14.5 8 33 ...
$ minority.diff : num  -0.268 -0.268 -0.268 -0.268 -0.268 ...
$ id          : int  1317 1317 1317 1317 1317 1317 1317 1317 1317 13
$ id.sec      : Factor w/ 40 levels "P5762","P2639",...: 33 33 33 33
..- attr(*, "scores")= num [1:40(1d)] -1.194 -0.964 -0.757 999.403 -0.523 .
.. ..- attr(*, "dimnames")=List of 1
.. .. ..$ : chr "P5762" "P2639" "P8854" "C4530" ...
$ by          : Factor w/ 1977 levels "1","2","3","4",...: 1 2 3 4 5
$ b_(Intercept) : num  13.4 13.4 13.4 13.3 13.4 ...

```

```
$ b_ses           : num  1.7 1.71 1.69 1.7 1.69 ...
$ b_ses.m         : num  2.41 2.42 2.43 2.39 2.4 ...
$ b_SectorPublic  : num  -0.935 -0.937 -0.942 -0.928 -0.933 ...
$ b_ses:SectorPublic : num  1.17 1.16 1.18 1.17 1.19 ...
$ b_ses.m:SectorPublic: num  1.41 1.41 1.39 1.43 1.42 ...
```

```
# scatterplotMatrix(fits.dropone[,grep('~b_',names(fits.dropone))], id = TRUE)
```

Dropping one school at a time

```
system.time(
  fits.dropschool <- dropone(fitc, ~ id) # only 6 seconds
)
```

```
user  system elapsed
4.67   0.06   5.16
```

- If some models don't converge, rerun fitc with larger msMaxIter, returnObject = T, sing.tol = 10e-20 to see if all drops will converge.

```
fitc$call
```

```
lme.formula(fixed = mathach ~ (ses + ses.m) * Sector, data = dd,
  random = ~1 + ses | id)
```

```
fitc <- update(fitc, control = list(returnObject = TRUE, msMaxIter = 1000) )
fitc$call
```

```
lme.formula(fixed = mathach ~ (ses + ses.m) * Sector, data = dd,
  random = ~1 + ses | id, control = list(returnObject = TRUE,
    msMaxIter = 1000))
```

```
system.time(
  fits.dropschool <- dropone(fitc, ~ id) # only 6 seconds
)
```

```
user  system elapsed
4.06   0.00   4.30
```

```
str(fits.dropschool)
```

```
'data.frame':    40 obs. of  21 variables:
 $ school      : int  1317 1906 2208 2458 2626 2629 2639 2658 2771 30...
 $ Size        : int  455 400 1061 545 2142 1314 2713 780 415 760 ...
 $ Sector      : Factor w/ 2 levels "Catholic","Public": 1 1 1 1 2 1 ...
 $ PRACAD      : num  0.95 0.87 0.68 0.89 0.4 0.81 0.14 0.79 0.24 0.5...
 $ DISCLIM     : num  -1.694 -0.939 -0.864 -1.484 0.142 ...
 $ n           : int  48 53 60 57 38 57 42 45 55 53 ...
 $ ses.m       : num  0.3453 0.5116 0.4232 0.2278 -0.0648 ...
```

```

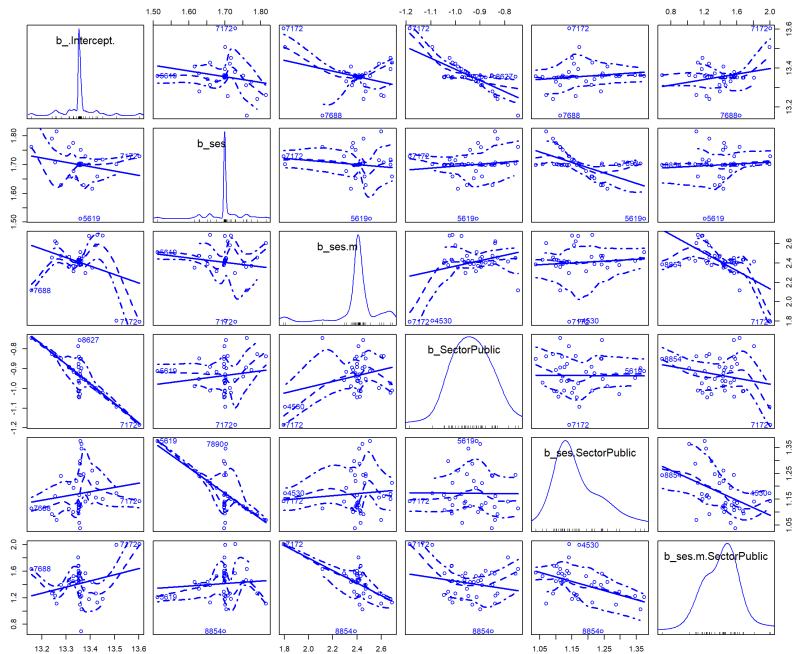
$ ses.iqr           : num  0.782 1.01 0.983 0.79 0.597 ...
$ ses.trange        : num  1.17 1.59 1.5 1.95 1.38 ...
$ ses.sd            : num  0.556 0.614 0.598 0.658 0.56 ...
$ minority.diff      : num  -0.268 -0.366 NaN -0.582 NaN ...
$ id                : Factor w/ 40 levels "1317","1906",...: 1 2 3 4 5 6 7
$ id.sec            : Factor w/ 40 levels "P5762","P2639",...: 33 39 37 32
$ ses.cvar          : num  0.3453 0.5116 0.4232 0.2278 -0.0648 ...
$ by                : Factor w/ 40 levels "1317","1906",...: 1 2 3 4 5 6 7
$ b_(Intercept)     : num  13.4 13.3 13.3 13.4 13.4 ...
$ b_ses             : num  1.71 1.68 1.65 1.63 1.7 ...
$ b_ses.m           : num  2.58 2.31 2.39 2.47 2.42 ...
$ b_SectorPublic     : num  -0.999 -0.917 -0.925 -0.954 -1.012 ...
$ b_ses:SectorPublic : num  1.16 1.19 1.22 1.25 1.13 ...
$ b_ses.m:SectorPublic: num  1.26 1.52 1.44 1.33 1.45 ...

```

```

scatterplotMatrix(fits.dropschool[,grep('^b_',names(fits.dropschool))], id = TRUE)

```



What does this show?

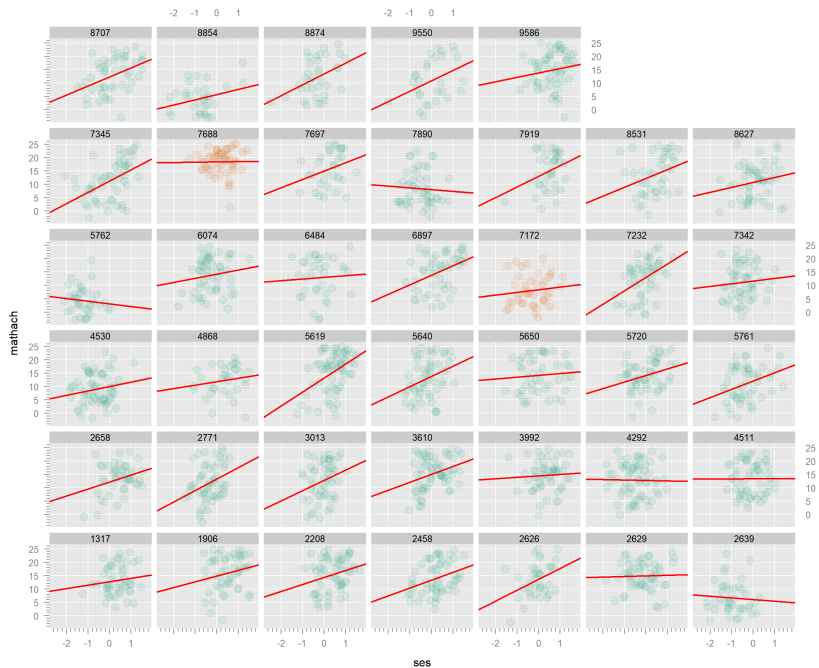
Which points represent Catholic schools and which Public schools?

```
if(interactive) {  
  names(fits.dropschool)  
  Init3d()  
  Plot3d('b_(Intercept)' ~ b_ses + 'b_ses:SectorPublic' | Sector,  
         fits.dropschool, sphere.size = 1.5)  
  Id3d() # to identify schools  
}  
id_miss <- is.na(fits.dropschool$b_ses)  
categ <- ifelse(dd$id %in% c('7688','7172'), 'out', 'ok')  
subset( fits.dropschool, id %in% c('7688','7172'))
```

	school	Size	Sector	PRACAD	DISCLIM	n	ses.m	ses.iqr	ses.trange
7172	7172	280	Catholic	0.05	1.013	44	-0.2898182	1.0275	1.686
7688	7688	1410	Catholic	0.65	-0.575	54	0.1858889	0.8275	1.378
	ses.sd	minority.diff	id	id.sec	ses.cvar	by	b_(Intercept)		
7172	0.6764417	0.3915447	7172	C7172	-0.2898182	7172		13.60385	
7688	0.5644347	-0.1387347	7688	C7688	0.1858889	7688		13.15648	
	b_ses	b_ses.m	b_SectorPublic	b_ses:SectorPublic					
7172	1.729063	1.792919	-1.185338				1.141840		

7688	1.761263	2.115580	-0.745562	1.114109
	b_ses.m:SectorPublic			
7172		2.005783		
7688		1.635031		

```
gd(alpha=c(.1,.1))
xyplot(mathach ~ ses | paste(id), dd, groups = categ) +
  glayer(panel.lmline(x,y, lwd = 2, col = 'red'))
```



Not surprising!

Exercises —

Choose the problem that seems most appropriate:

#

*# 1) Add the variable 'Sex' to analysis above. Note that Sex has
a contextual variable: Sex composition, a variable with
interestingly different features in Catholic vs Public school.*

#

*# If you wish to gain some experience with convergence problems
use the full Level 1 model (~ ses*Sex | id) for your RE model.
If this fails to converge, try to use the advice given above to
correct the problem. If you prefer not to bother for now,
just use the additive model:*

#

random = ~1 + ses + Sector |id

#

*# Try to estimate the Gender Gap under various circumstances:
ses and Sector if the effect of Sex interacts with these variables.*

#

```
# Which circumstances are associated with large gender gaps,  
# with small gender gaps?  
#  
# Make graphs of the gender gap as a function of its moderators,  
# preferably with confidence bounds.  
#  
# 2) Analyze the data in 'bdf' in 'nlme':  
# > data (bdf)  
# > ?bdf  
# on Language Scores of 8-Graders in The Netherlands  
#  
# This is similar to the 'hs' dataset but different enough  
# to be interesting.  
#  
# The response of interest is 'langPOST' a measure of language  
# achievement at the end of the school year.  
#  
# Potential predictors include: ses, sex, denominat (Sector),  
# Minority, IQ.verb as well as a pretest: langPRE.  
#  
# This raises the interesting question of the best use for the
```

```
# pretest. Should it be used as a covariate or should you analyze
# the gain score: langPOST - langPRE as the response.
#
# As usual, the best course of action may depend on the questions
# and intended interpretation of the results.
#
# 3) Study the role of 'Minority' in the 'hs' data following the
# script below
#
# 4) Validate the model above with the other half of the data 'hs2'
# What differences do you note?
#
```