

Methods 3: Multilevel Statistical Modeling and Machine Learning

Week 3: *Generalized linear mixed effects models*

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Messages

- Practical exercise due 23.59 tomorrow
- Make sure to add your GitHub repository – a few are still missing:
<https://cryptpad.fr/pad/#/2/pad/edit/U21qNTbLgfkRiGZU1bnmDE2o/>
- Remember, Class 2 (10-12) will be in 1453-116 tomorrow

RECAP on pooling

SLEEP STUDY EXAMPLE

<https://psyteachr.github.io/stat-models-v1/introducing-linear-mixed-effects-models.html>

Learning goals and outline –

Linear Mixed Effects Models (LMM)

1) Why can it be a good idea to do mixed effects modelling?

2) Understanding the basics of multilevel modelling

- also known as linear mixed effects modelling

3) Appreciating the difference between the different levels of effects

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= *or random* and *fixed* effects as they are also called

Pooling - summary

- Complete pooling

- ignoring a categorical predictor (e.g. *subject*)

- No pooling

- model each level of the categorical predictor separately

- Partial pooling

- we model both an average and each level of the categorical predictor (e.g. *subject*)

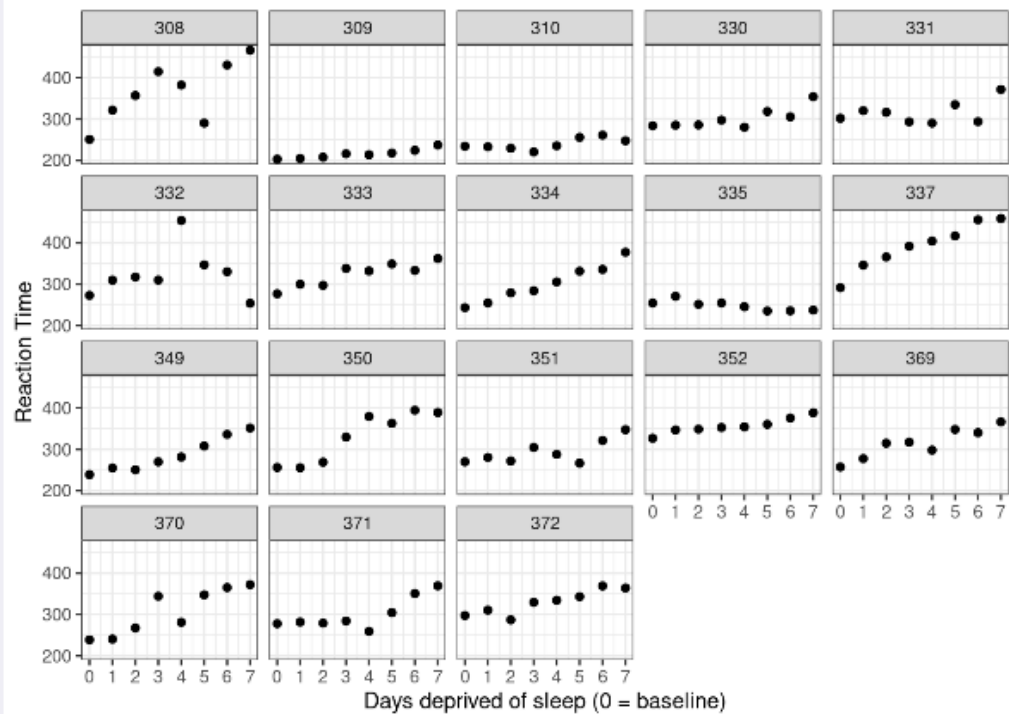


Figure 5.3: Data from Belenky et al. (2003), showing reaction time at baseline (0) and after each day of sleep deprivation.

COMPLETE POOLING

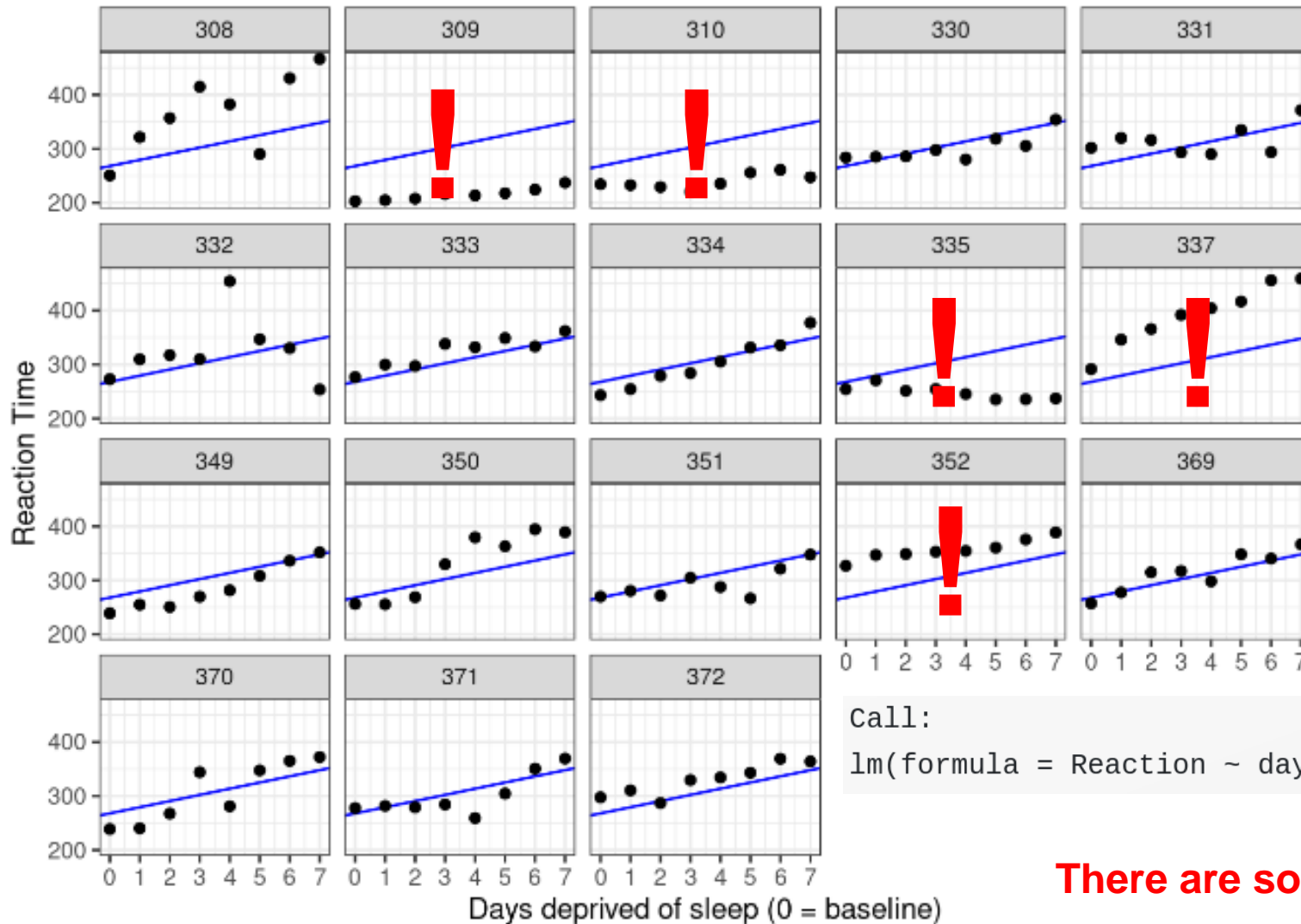
Coefficients:

	Estimate
(Intercept)	267.967
days_deprived	11.435

Call:

```
lm(formula = Reaction ~ days_deprived, data = sleep2)
```

There are some bad fits




```
lm(formula = Reaction ~ days_deprived + Subject + days_deprived:Subject,  
    data = sleep2)
```

```
## Coefficients:
```

##	Estimate
## (Intercept)	288.2175
## days_deprived	21.6905
## Subject309	-87.9262
## Subject310	-62.2856
## Subject330	-14.9533
## Subject331	9.9658
## Subject332	27.8157

... and the remaining 12 subjects

```
## days_deprived:Subject309 -17.3334  
## days_deprived:Subject310 -17.7915  
## days_deprived:Subject330 -13.6849  
## days_deprived:Subject331 -16.8231  
## days_deprived:Subject332 -19.2947  
## days_deprived:Subject333 -10.8151
```

... and the remaining 12 subjects

NO POOLING

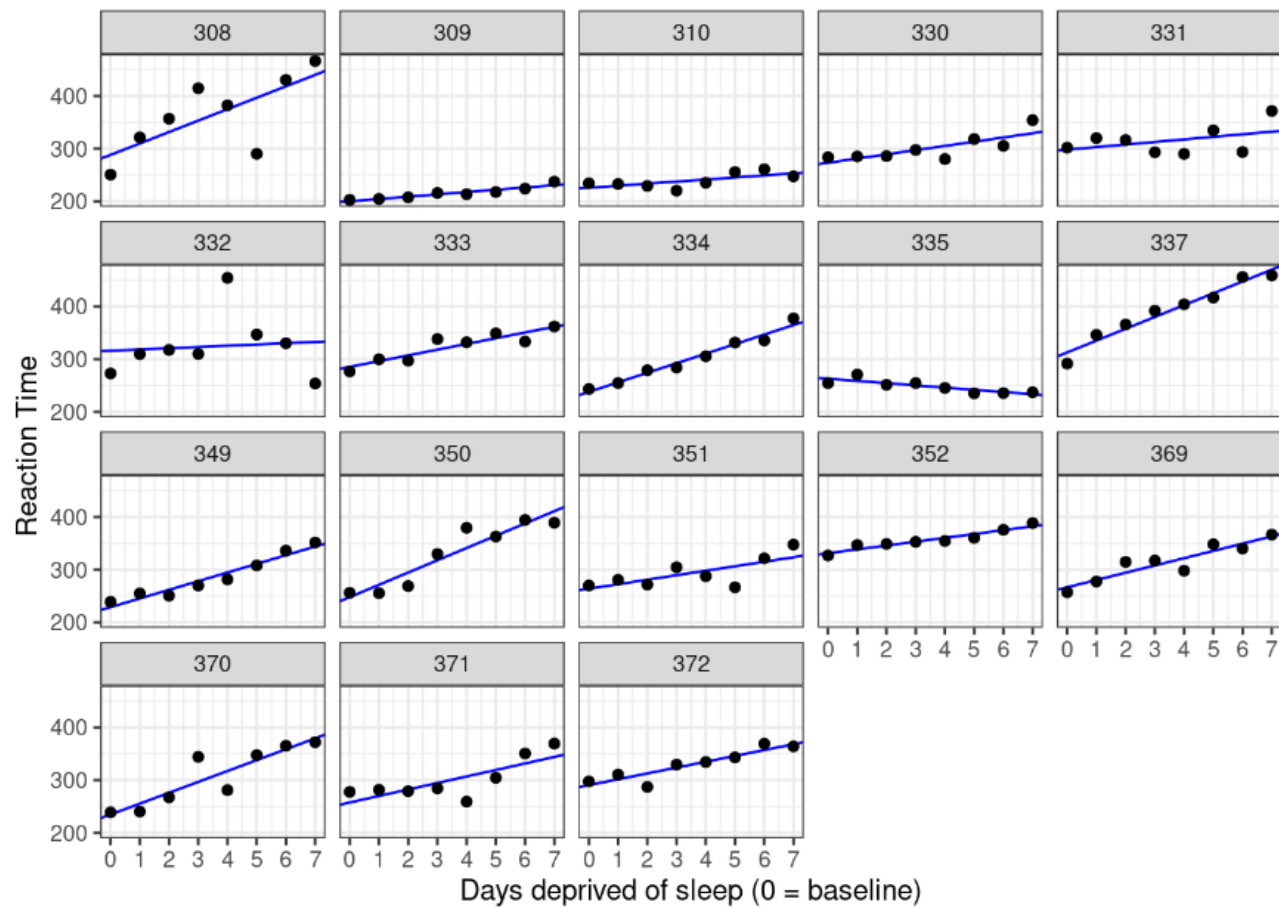


Figure 5.5: Data plotted against fits from the no-pooling approach.

NO POOLING

Good fits now:

What are the limits of this mo

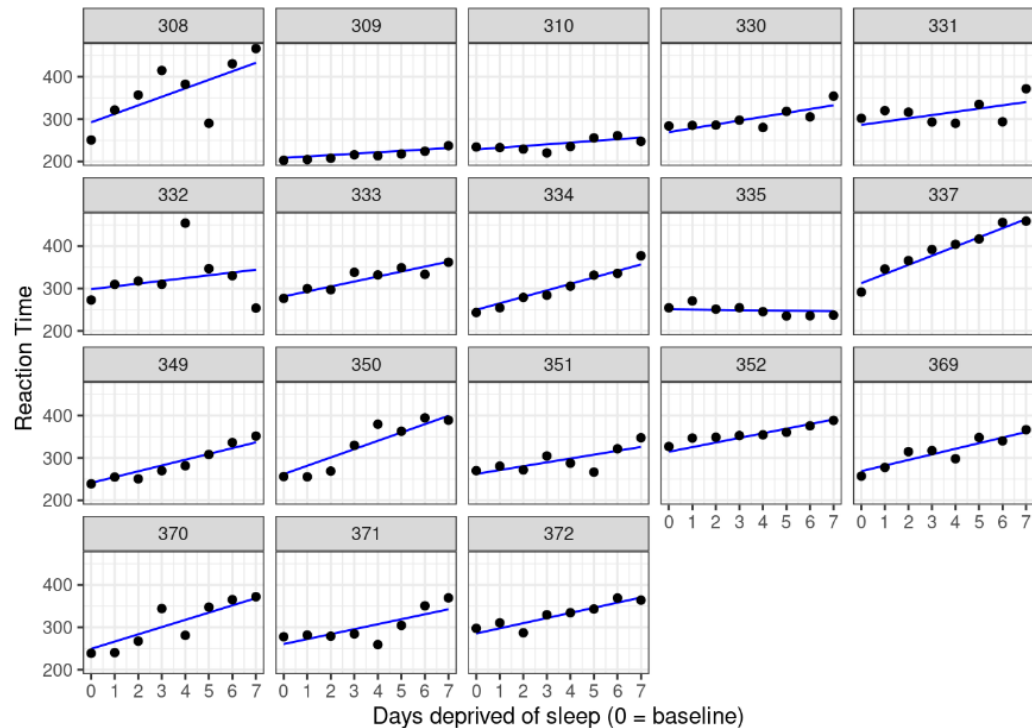


Figure 5.6: Data plotted against predictions from a partial pooling approach.

PARTIAL POOLING

Fixed effects:

	Estimate	Std. Error	t value
(Intercept)	267.967	8.266	32.418
days_deprived	11.435	1.845	6.197

```
ranef(pp_mod)[["Subject"]]
```

	(Intercept)	days_deprived
308	24.4992891	8.6020000
309	-59.3723102	-8.1277534
310	-39.4762764	-7.4292365
330	1.3500428	-2.3845976

Linear mixed model fit by REML ['lmerMod']

Formula: Reaction ~ days_deprived + (days_deprived | Subject)

Data: sleep2

No pooling vs partial pooling

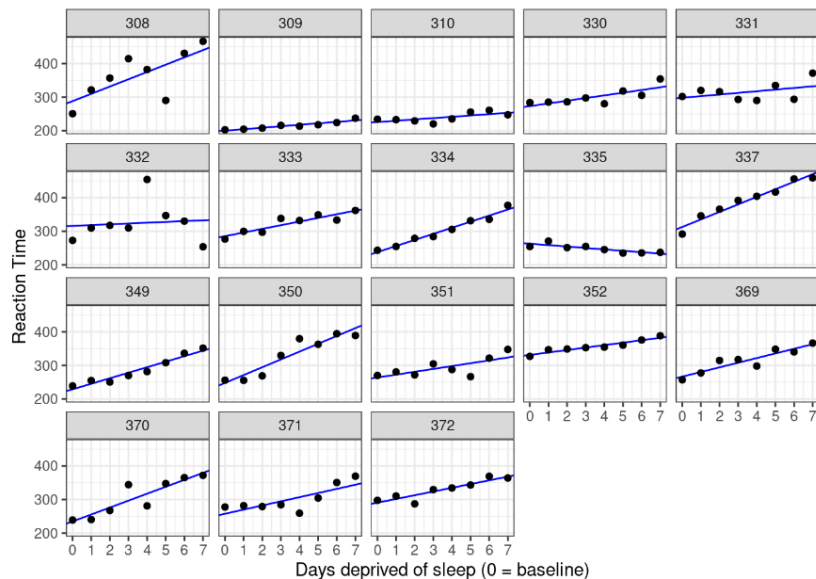


Figure 5.5: Data plotted against fits from the no-pooling approach.

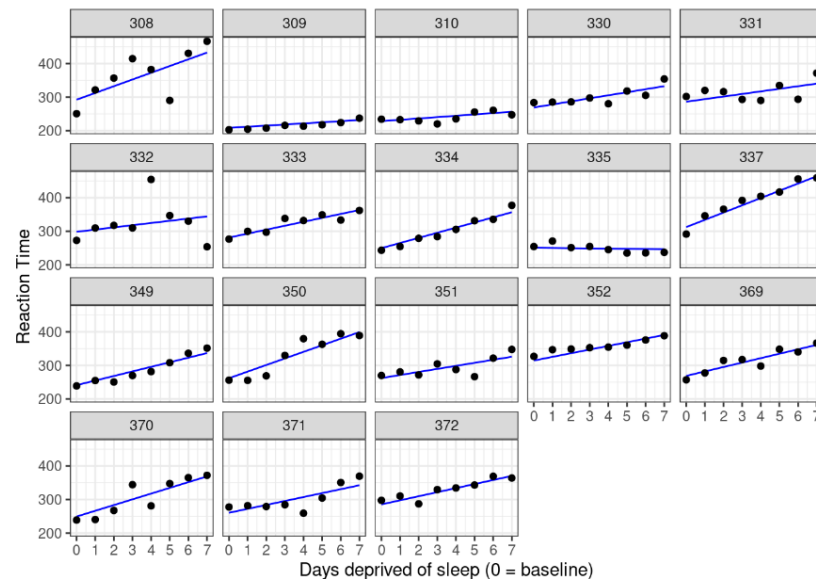


Figure 5.6: Data plotted against predictions from a partial pooling approach.

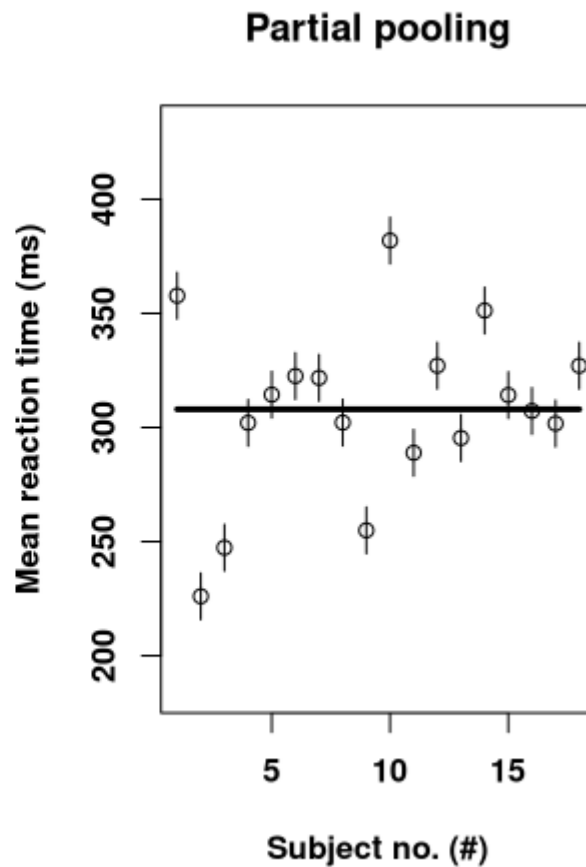
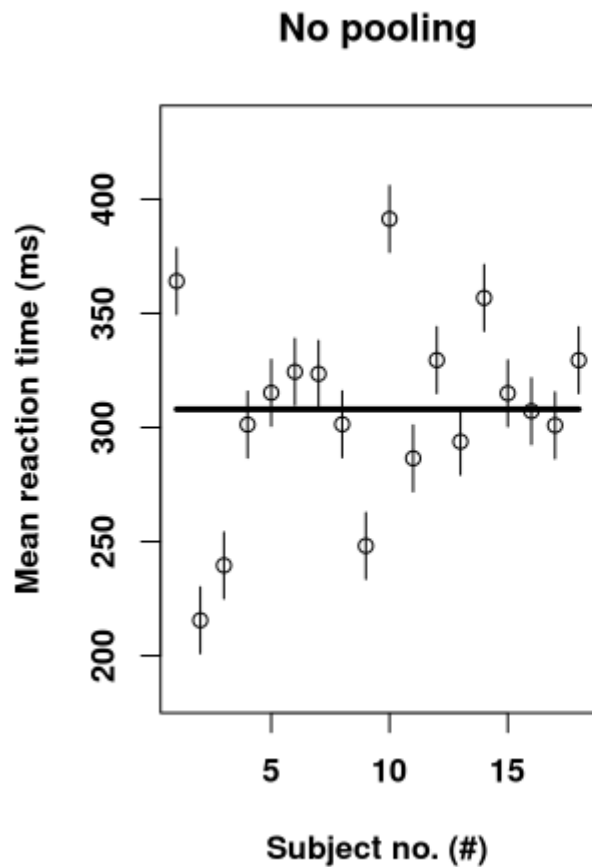
Both model the individual variance – but only one is generalisable outside the s

Partial pooling

(Gelman and Hill, 2006 (12.1))

$$\hat{\alpha}_j^{multilevel} \approx \frac{\frac{n_j}{\sigma_y^2} \bar{y}_j + \frac{1}{\sigma_\alpha^2} \bar{y}_{all}}{\frac{n_j}{\sigma_y^2} + \frac{1}{\sigma_\alpha^2}}$$

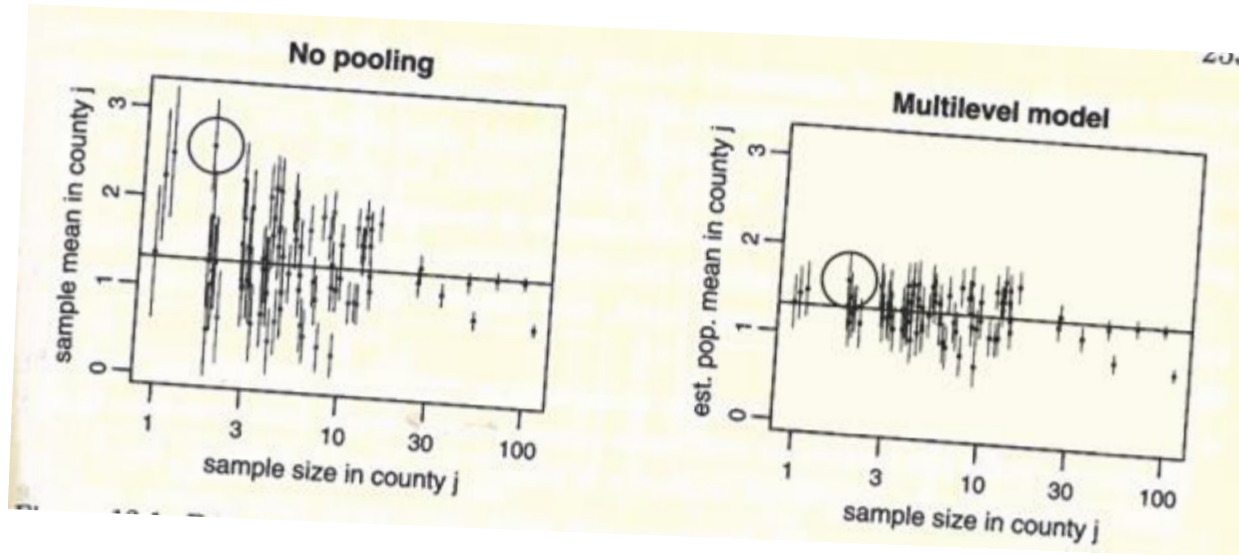
another scary looking th



What is the advantage

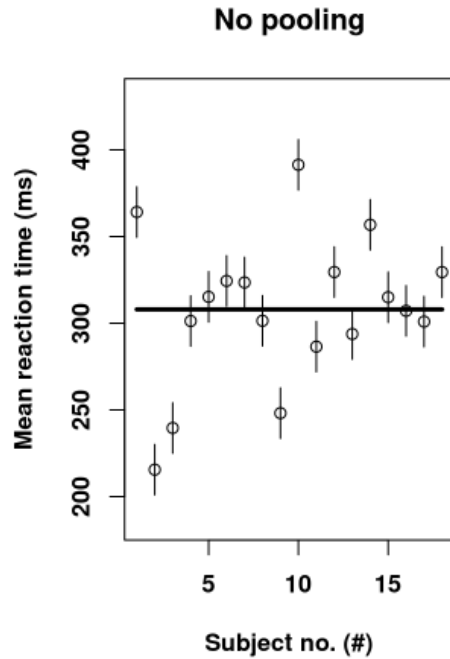
Now with different sample sizes

What is the advantage

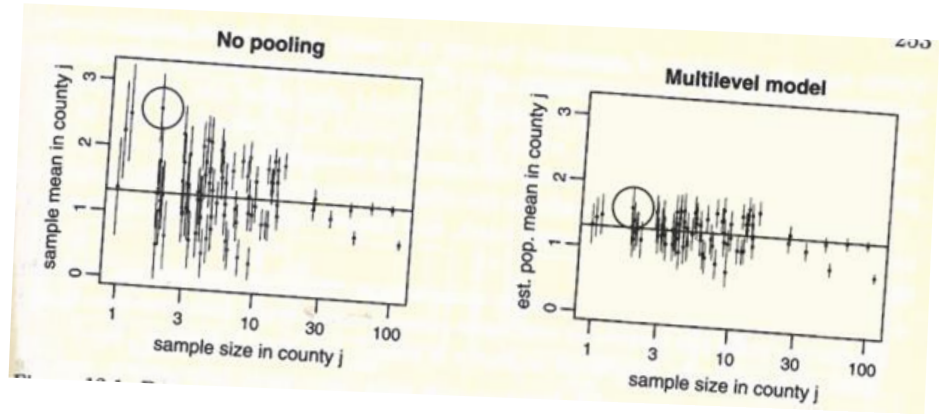
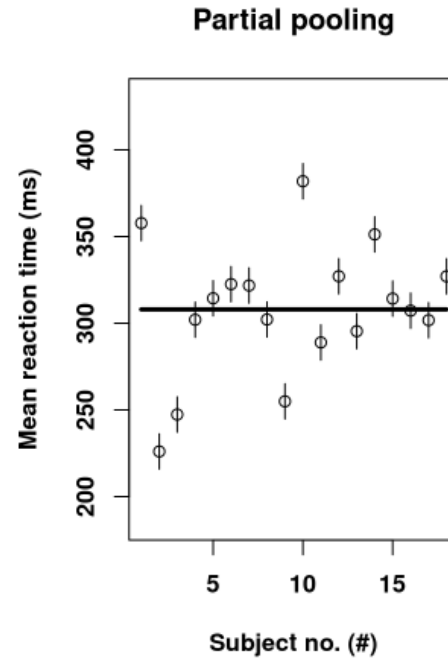


(Gelman and Hill, 2006)

Same n for each subject



Different n for each county



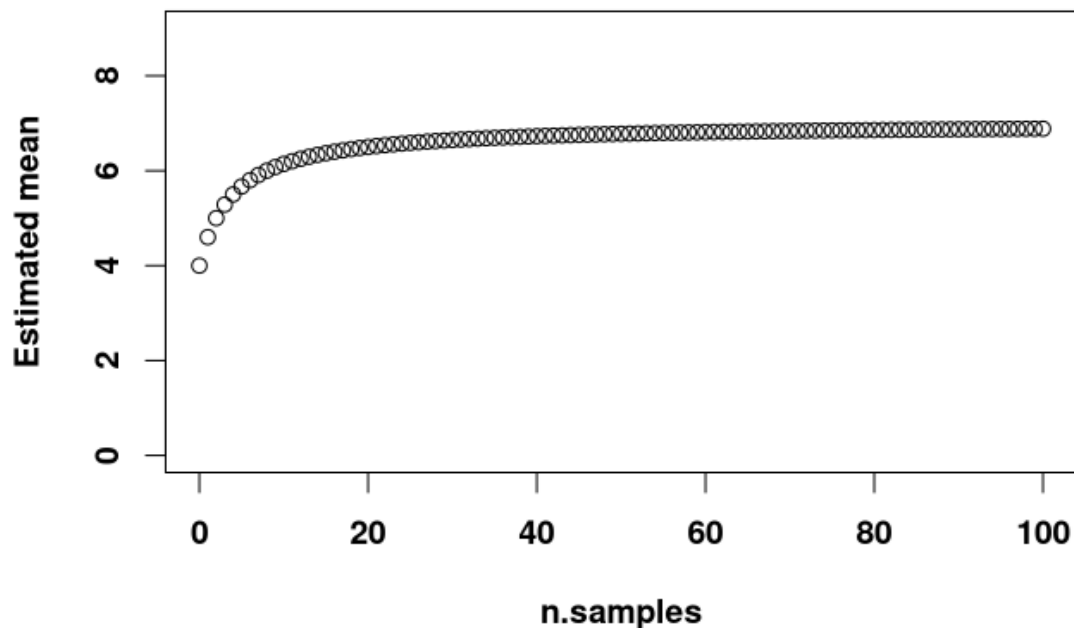
(Gelman and Hill, 2006)

```
estimate.multilevel.mean <- function(n.j, sigma.y, sigma.mean, y.j, y.all)
{
  alpha <- ((n.j / sigma.y^2) * y.j + (1 / sigma.mean^2) * y.all) /
    ((n.j / sigma.y^2) + (1 / sigma.mean^2))
  return(alpha)
}
```

"Baseline" plot

```
## "baseline"

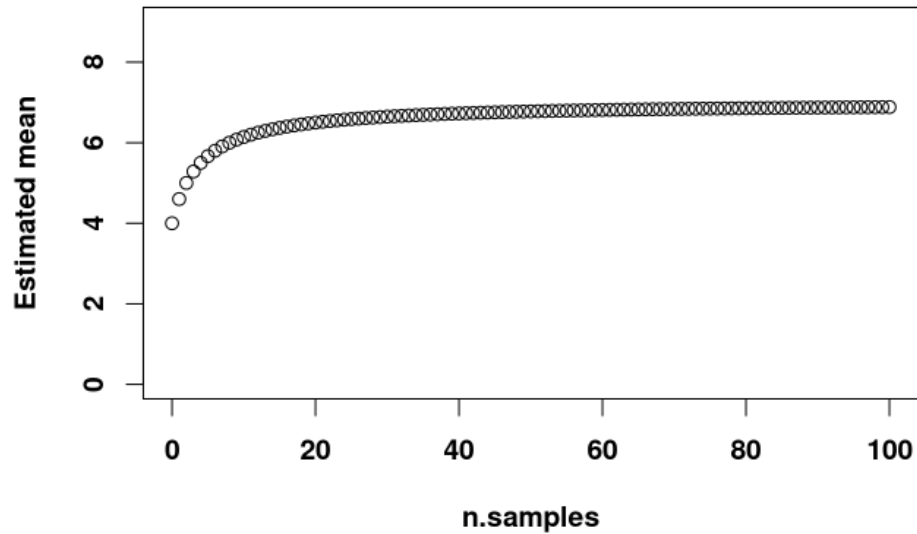
sigma.y <- 3
y.j <- 7
sigma.mean <- 1.5
y.all <- 4
ns <- 0:100
```



```
## "baseline"

sigma.y <- 3
y.j <- 7
sigma.mean <- 1.5
y.all <- 4
ns <- 0:100
```

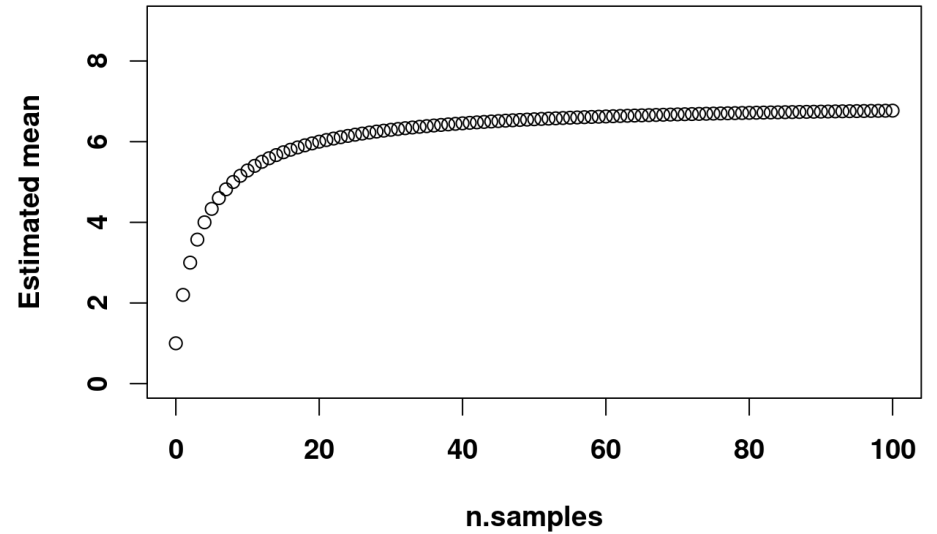
"Baseline" plot



```
## small group effect

sigma.y <- 3
y.j <- 7
sigma.mean <- 1.5
y.all <- 1
ns <- 0:100
```

Small group effect



```
## "baseline"
```

```
sigma.y <- 3
```

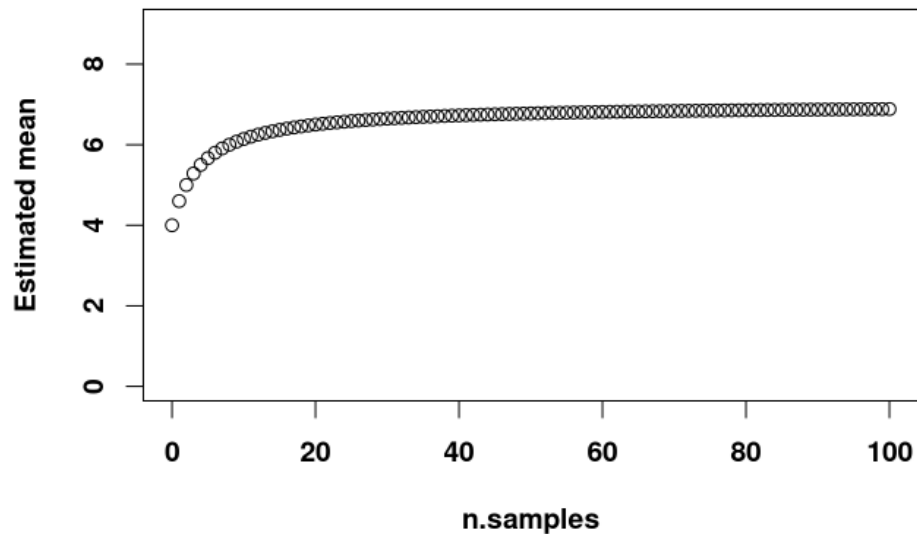
```
y.j <- 7
```

```
sigma.mean <- 1.5
```

```
y.all <- 4
```

```
ns <- 0:100
```

"Baseline" plot



```
## noisy individual effect
```

```
sigma.y <- 6
```

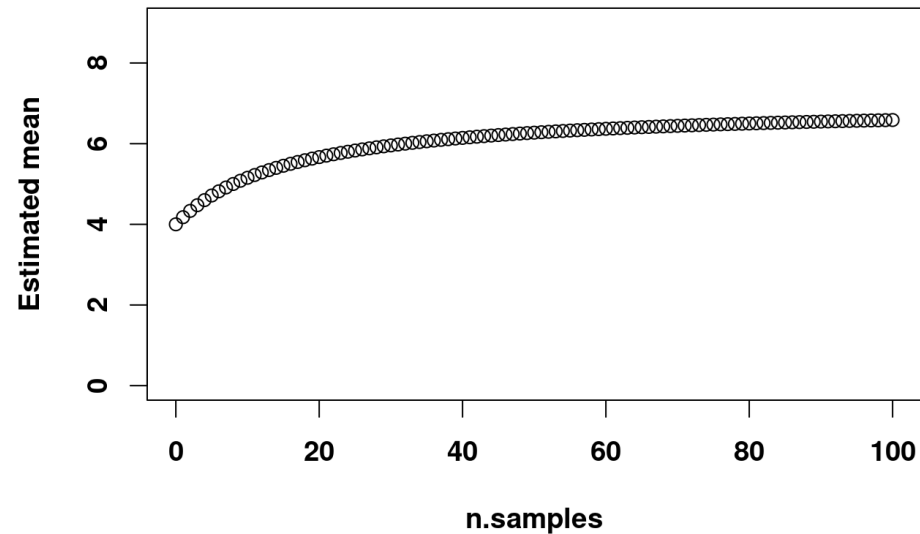
```
y.j <- 7
```

```
sigma.mean <- 1.5
```

```
y.all <- 4
```

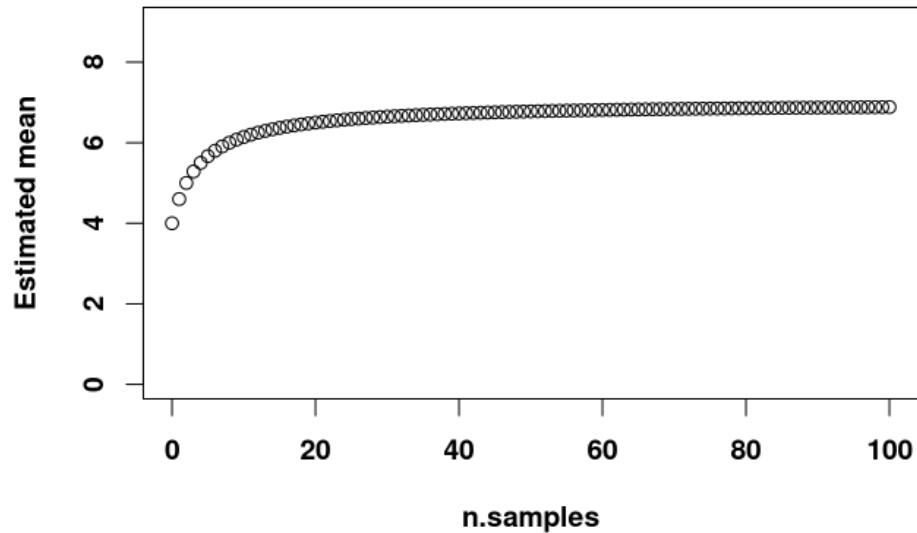
```
ns <- 0:100
```

Noisy individual effect



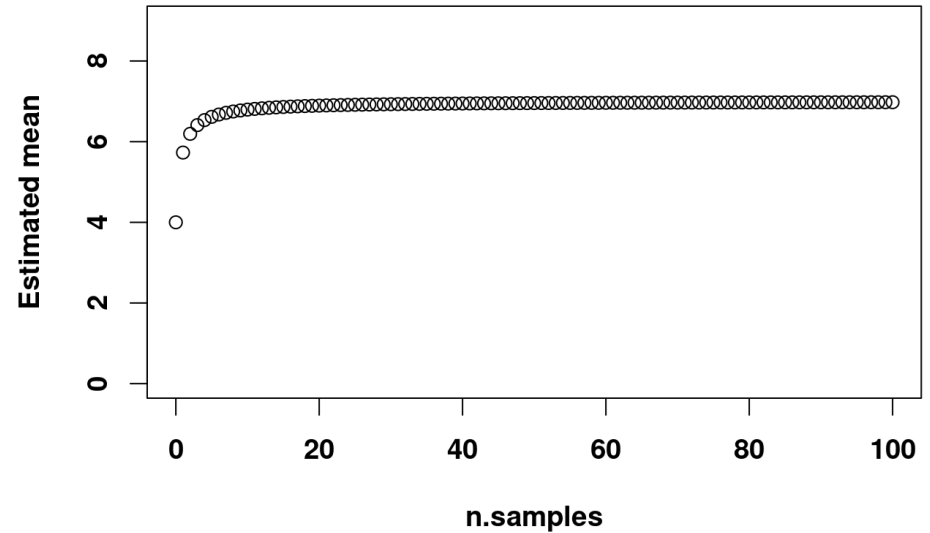
```
## "baseline"  
  
sigma.y <- 3  
y.j <- 7  
sigma.mean <- 1.5  
y.all <- 4  
ns <- 0:100
```

"Baseline" plot



```
## noisy group effect  
  
sigma.y <- 3  
y.j <- 7  
sigma.mean <- 3.5  
y.all <- 4  
ns <- 0:100
```

Noisy group effect



Motivation for multilevel modelling:

We want to use all the information in the data while fulfilling the assumptions necessary for the residuals

We can add:

Without letting small or uncertain samples unduly affect our group estimate

Now on to generalized linear mixed models ...

Did you learn?

Linear Mixed Effects Models (LMM)

1) Why can it be a good idea to do mixed effects modelling?

2) Understanding the basics of multilevel modelling

- also known as linear mixed effects modelling

3) Appreciating the difference between the different levels of effects

= *or random* and *fixed* effects as they are also called

**... but let's do a recap of the
generalized linear model first**

Learning goals

Generalized Linear Mixed Effects Models (GLMM)

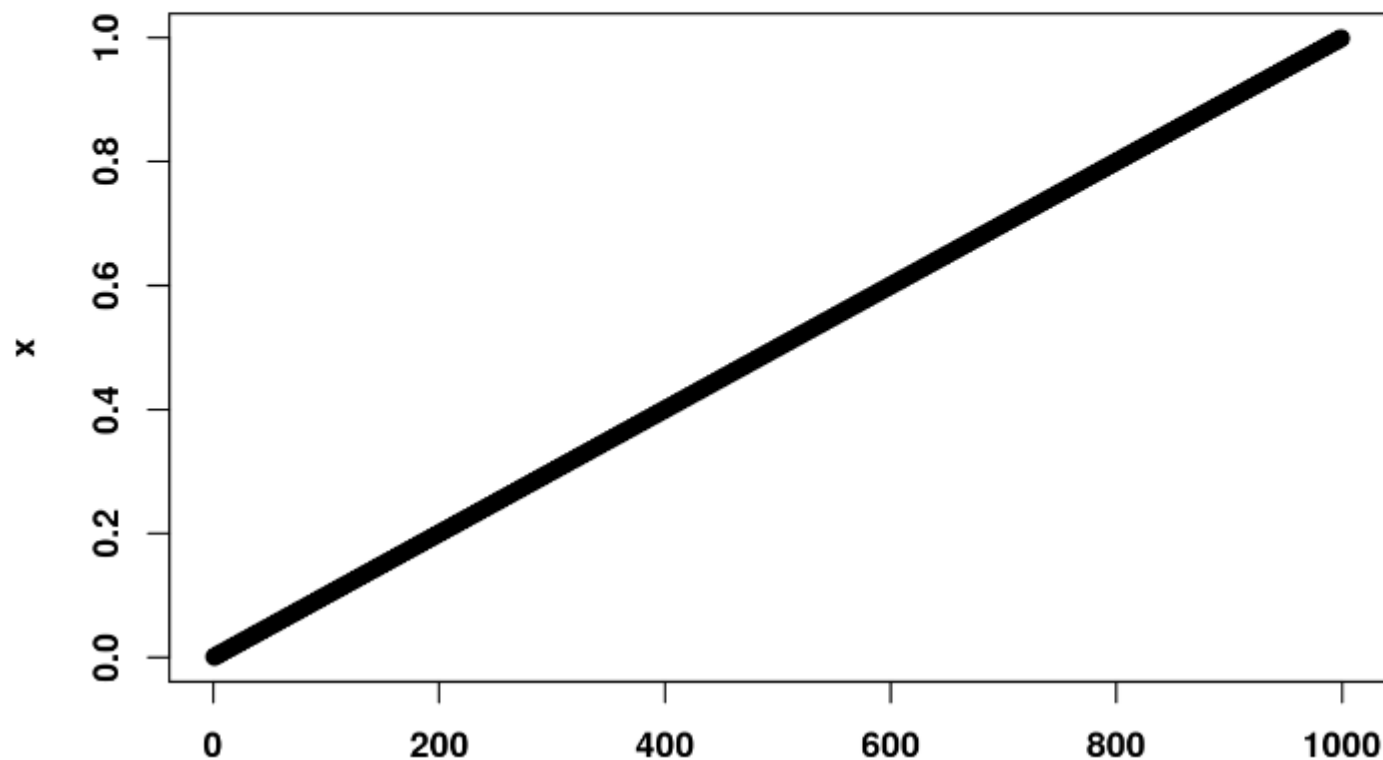
- 1) Understanding that we can extend the scope of our multilevel modelling by using appropriate link functions and data distributions
- 2) Understanding the multilevel equivalent of the GLM



Breaking all promises and goin

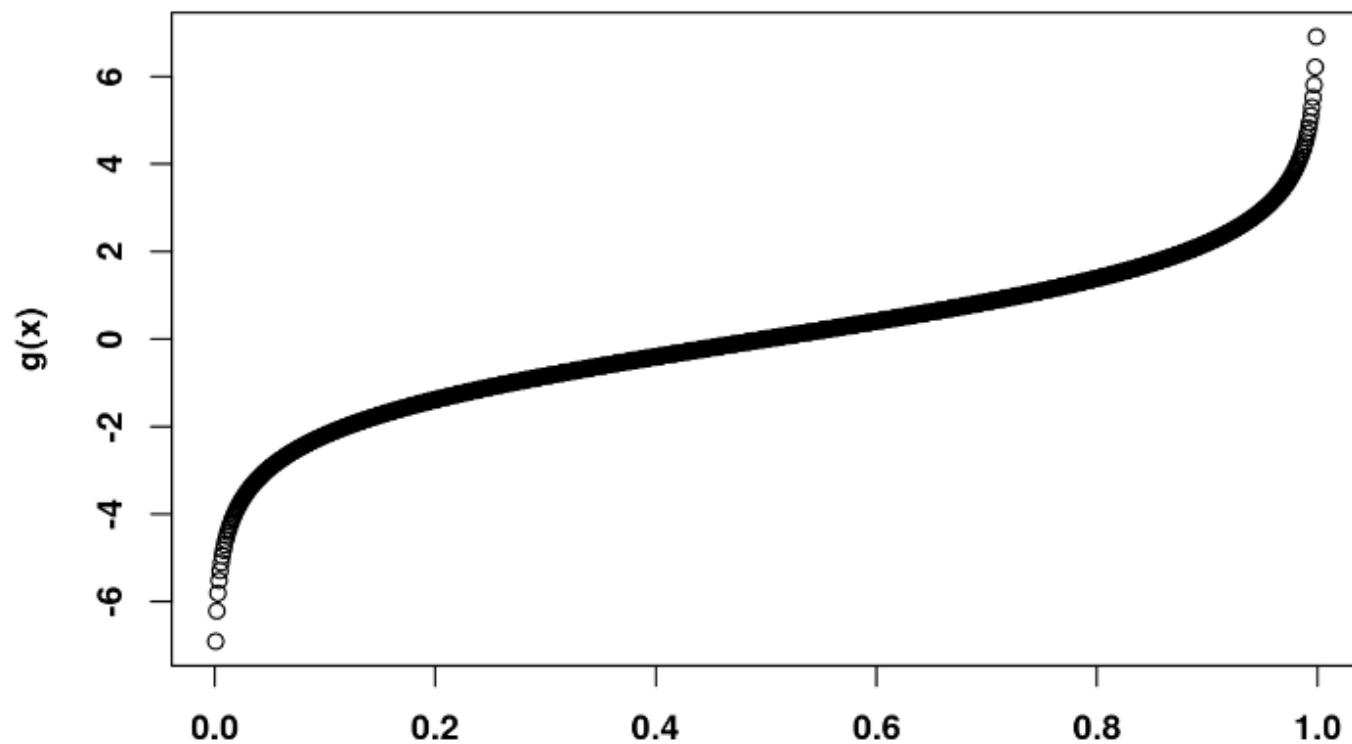

```
x <- seq(0.001, 0.999, 0.001)  
plot(x, main='Original probability data (on the range from 0-1)')
```

Original probability data (on the range from 0-1)

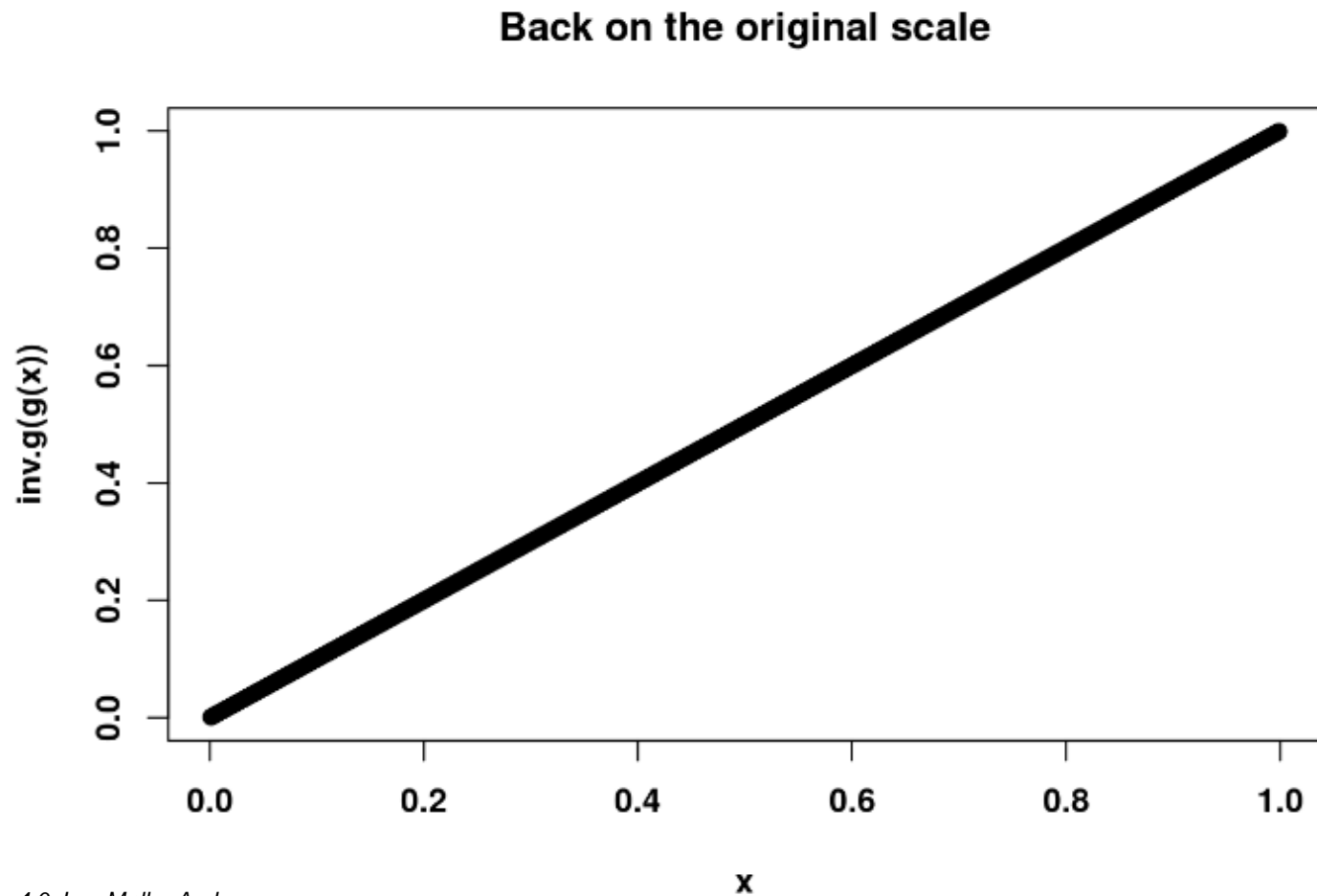


```
plot(x, g(x), main='Log-it transformed, on the range from -Inf to Inf')
```

Log-it transformed, on the range from -Inf to Inf



```
plot(x, inv.g(g(x)), main='Back on the original scale')
```



These are the fitted values

```
y.hat <- inv.g(X %*% beta.hat)
print(head(y.hat))
```

```
##           [,1]
## Mazda RX4    0.8172115
## Mazda RX4 Wag 0.6157283
## Datsun 710    0.9373069
## Hornet 4 Drive 0.2897304
## Hornet Sportabout 0.1415972
## Valiant      0.1320944
```

```
print(head(y.hat - logistic.model$fitted.values))
```

```
##           [,1]
## Mazda RX4      0
## Mazda RX4 Wag  0
## Datsun 710     0
## Hornet 4 Drive  0
## Hornet Sportabout 0
## Valiant        0
```

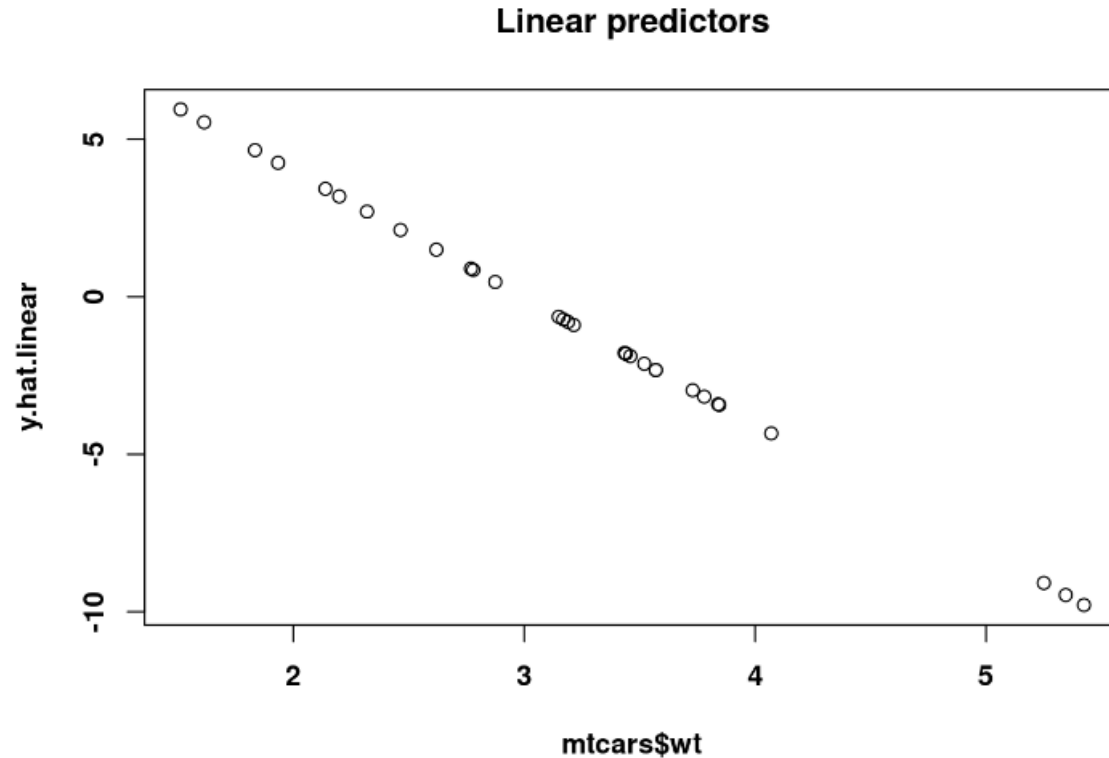
These are the linear predictors

```
y.hat.linear <- X %*% beta.hat
```

```
print(head(y.hat.linear - logistic.model$linear.predictors))
```

```
##           [,1]  
## Mazda RX4           0  
## Mazda RX4 Wag       0  
## Datsun 710           0  
## Hornet 4 Drive       0  
## Hornet Sportabout    0  
## Valiant              0
```

Looks like a “normal” linear regression



Some link functions

Usage

```
family(object, ...)
```

```
binomial(link = "logit")
```

```
gaussian(link = "identity")
```

```
Gamma(link = "inverse")
```

```
inverse.gaussian(link = "1/mu^2")
```

```
poisson(link = "log")
```

```
quasi(link = "identity", variance = "constant")
```

```
quasibinomial(link = "logit")
```

```
quasipoisson(link = "log")
```


Important difference from the general linear model

We also make maximum likelihood estimates for

Did you learn?

Generalized Linear Mixed Effects Models (GLMM)

- 1) Understanding that we can extend the scope of our multilevel modelling by using appropriate link functions and data distributions
- 2) Understanding the multilevel equivalent of the GLM

References

.Gelman, A., Hill, J., 2006. Data Analysis Using Regression and Multilevel/Hierarchical Models. Cambridge University Press.