

Hierarchical Linear Models

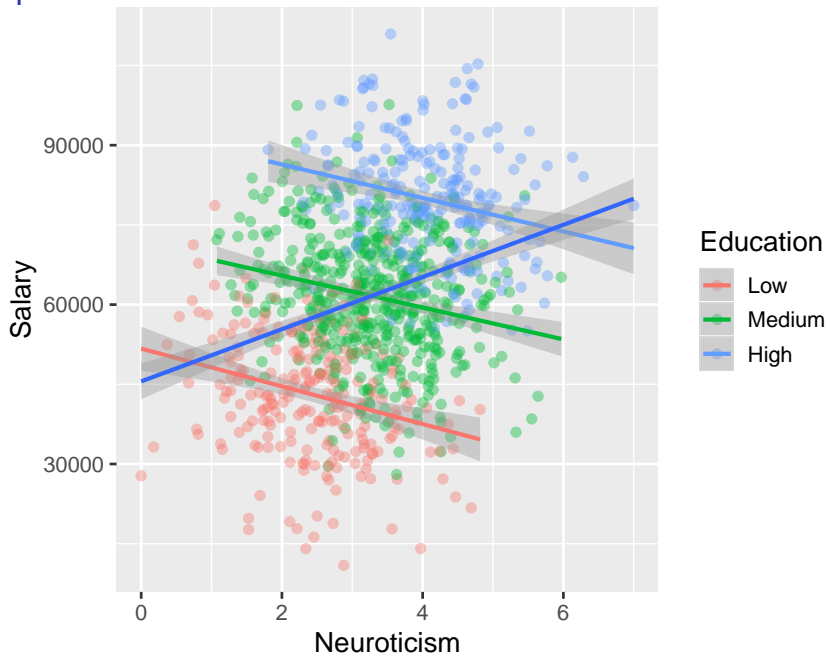
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A Common Challenge|Repeated measures

- Multiple measures from the same person who might also be part of a group
- Multiple exposures across people to the same stimuli
- Repeated-measures ANOVA accommodates correlated observations due to differences between people
- Multi-level general linear models go further and explicitly model general grouping effects (e.g., sampled stimuli)
- Addresses Simpson's paradox, where individual trends don't reflect the aggregate trend

Simpson's Paradox



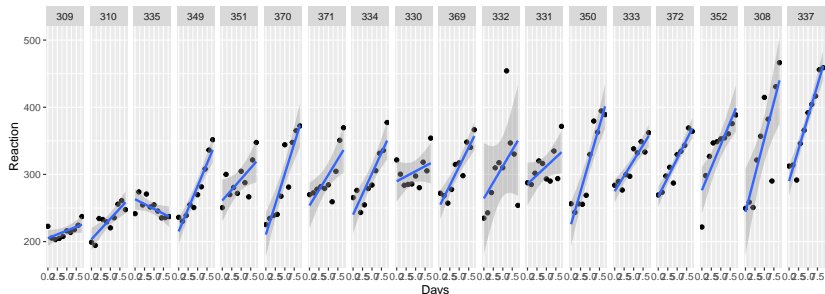
Useful packages

- `library(tidyverse)` for `ggplot2`, `dplyr`
- `library(skimr)` for `skim`, which provides a quick overview of data
- `library(lme4)` for mixed effect, multi-level general linear models
- `library(lmerTest)` for extracting p-values
- `library(psycho)` for extracting APA reportable sentences

Effect of Sleep Disruption

```
sleep.df = sleepstudy
```

```
ggplot(sleep.df, aes(Days, Reaction)) +  
  geom_point() +  
  geom_smooth(method = "lm") +  
  facet_grid(.~reorder(Subject, Reaction))
```



Uniform Effect of Sleep Disruption?

```
sleep.fit = lm(Reaction ~ Days, sleep.df)
```

```
sleep.df$prediction = predict(sleep.fit)
```

```
summary(sleep.fit)
```

```
##
```

```
## Call:
```

```
## lm(formula = Reaction ~ Days, data = sleep.df)
```

```
##
```

```
## Residuals:
```

##	Min	1Q	Median	3Q	Max
##	-110.848	-27.483	1.546	26.142	139.953

```
##
```

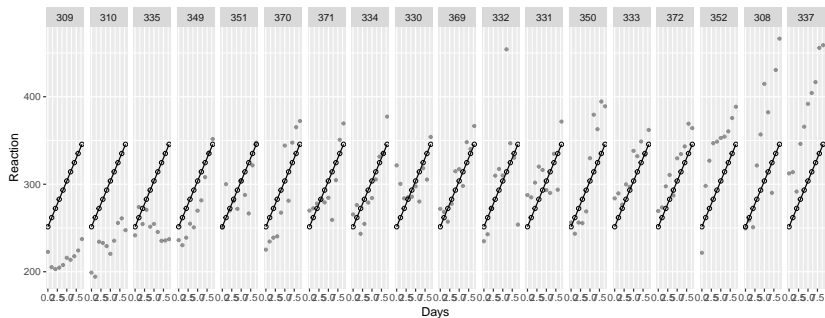
```
## Coefficients:
```

##	Estimate	Std. Error	t value	Pr(> t)
## (Intercept)	251.405	6.610	38.033	< 2e-16 ***
## Days	10.467	1.238	8.454	9.89e-15 ***

```
##
```

Uniform Effect of Sleep Disruption?

```
ggplot(sleep.df, aes(Days, Reaction)) +  
  geom_point(alpha = .4, size = 1) +  
  geom_point(aes(y=prediction), shape = 21) +  
  geom_line(aes(y=prediction)) +  
  facet_grid(.~reorder(Subject, Reaction))
```



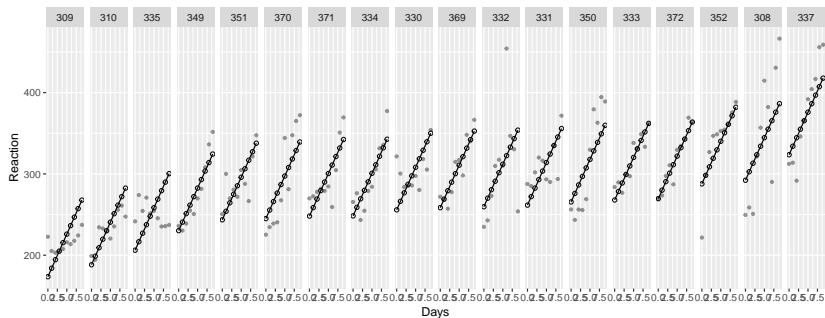
Random Effect for Intercept

```
sleep_int.fit = lmer(Reaction ~ Days + (1|Subject), sleep.c  
summary(sleep_int.fit)
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite  
## lmerModLmerTest]  
## Formula: Reaction ~ Days + (1 | Subject)  
##      Data: sleep.df  
##  
## REML criterion at convergence: 1786.5  
##  
## Scaled residuals:  
##      Min       1Q   Median       3Q      Max   
## -3.2257 -0.5529  0.0109  0.5188  4.2506   
##  
## Random effects:  
##   Groups    Name              Variance Std.Dev.  
##   Subject  (Intercept) 1378.2    37.12  
##   Residual                        960.5    30.99  
## Number of obs: 180, groups: Subject, 18
```

Random Effect for Intercept

```
ggplot(sleep.df, aes(Days, Reaction)) +  
  geom_point(alpha = .4, size = 1) +  
  geom_point(aes(y=prediction_int), shape = 21) +  
  geom_line(aes(y=prediction_int, group = Subject)) +  
  facet_grid(.~reorder(Subject, Reaction))
```



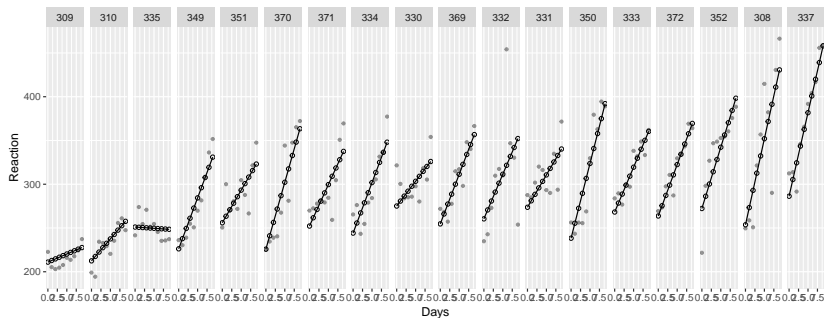
Random Effect for Intercept and Slope

```
sleep_int_slope.fit = lmer(Reaction ~ Days + (Days|Subject)  
summary(sleep_int_slope.fit)
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite  
## lmerModLmerTest]  
## Formula: Reaction ~ Days + (Days | Subject)  
##      Data: sleep.df  
##  
## REML criterion at convergence: 1743.6  
##  
## Scaled residuals:  
##      Min       1Q   Median       3Q      Max   
## -3.9536 -0.4634  0.0231  0.4633  5.1793   
##  
## Random effects:  
##   Groups    Name                Variance Std.Dev. Corr  
##   Subject  (Intercept)  611.90     24.737  
##           Days          35.08      5.923   0.07  
##   Residual                    654.04     25.560
```

Random Effects for Intercept and Slope

```
ggplot(sleep.df, aes(Days, Reaction)) +  
  geom_point(alpha = .4, size = 1) +  
  geom_line(aes(y=prediction_int_slope, group = Subject)) +  
  geom_point(aes(y=prediction_int_slope), shape = 21) +  
  facet_grid(.~reorder(Subject, Reaction))
```



Summary for Publication

- The “psycho” package provides functions for summarizing model output
- It creates a publication-ready summary table
- It can even create sentences describing the output
- Parameter estimates can be extracted for plotting

```
library(psycho)
analyze(sleep_int_slope.fit, CI = 95) %>%
  summary(results, round = 2) %>%
  mutate(p = format_p(p)) %>%
  kable()
```

Variable	Effect_Size	Coef	SE	t	df	p	
(Intercept)	NA	251.41	6.82	36.84	17.01	< .001***	
Days	medium	10.47	1.55	6.77	17.00	< .001***	

Summary for Publication|psycho can create sentences describing the output

```
analyze(sleep_int_slope.fit, CI = 95)
```

```
## The overall model predicting Reaction (formula = Reaction
```

```
## - The effect of Days is significant (beta = 10.47, SE
```

-The overall model predicting Reaction (formula = Reaction ~ Days + (Days | Subject)) has an total explanatory power (conditional R2) of 79.92%, in which the fixed effects explain 27.87% of the variance (marginal R2).

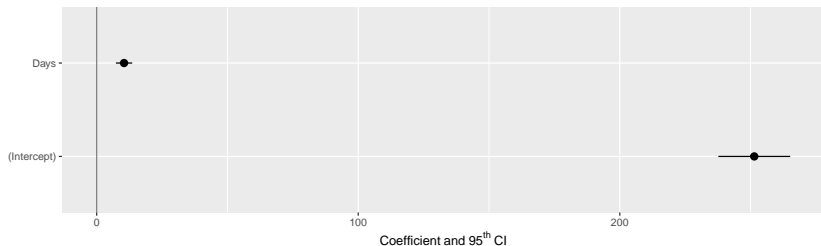
-The model's intercept is at 251.41 (SE = 6.82, 95% CI [237.68, 265.13]). Within this model:

-The effect of Days is significant (beta = 10.47, SE = 1.55, 95% CI [7.36, 13.58], $t(17) = 6.77$, $p < .001$) and can be considered as medium (std. beta = 0.54, std. SE = 0.079).

Summary for Publication|Parameter estimates can also be extracted for plotting

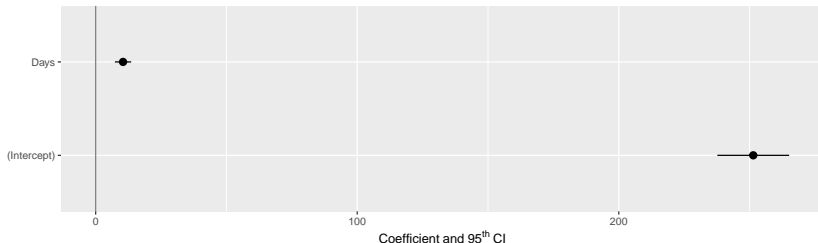
```
CI.lme.df = analyze(sleep_int_slope.fit)$summary
```

```
ggplot(CI.lme.df, aes(Variable)) +  
  geom_hline(yintercept = 0, alpha = .5)+  
  geom_pointrange(aes(y = Coef, ymin= CI_lower, ymax = CI_upper)) +  
  coord_flip() +  
  labs(x = "", y = expression(Coefficient~and~95^{th}~CI))
```



Exercise|Multi-level model fitting

- Fit a multi-level model with Subject as a random variable
- Use the `analyze` command from `psych` to extract an summary of effects
- Plot the effect estimates



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