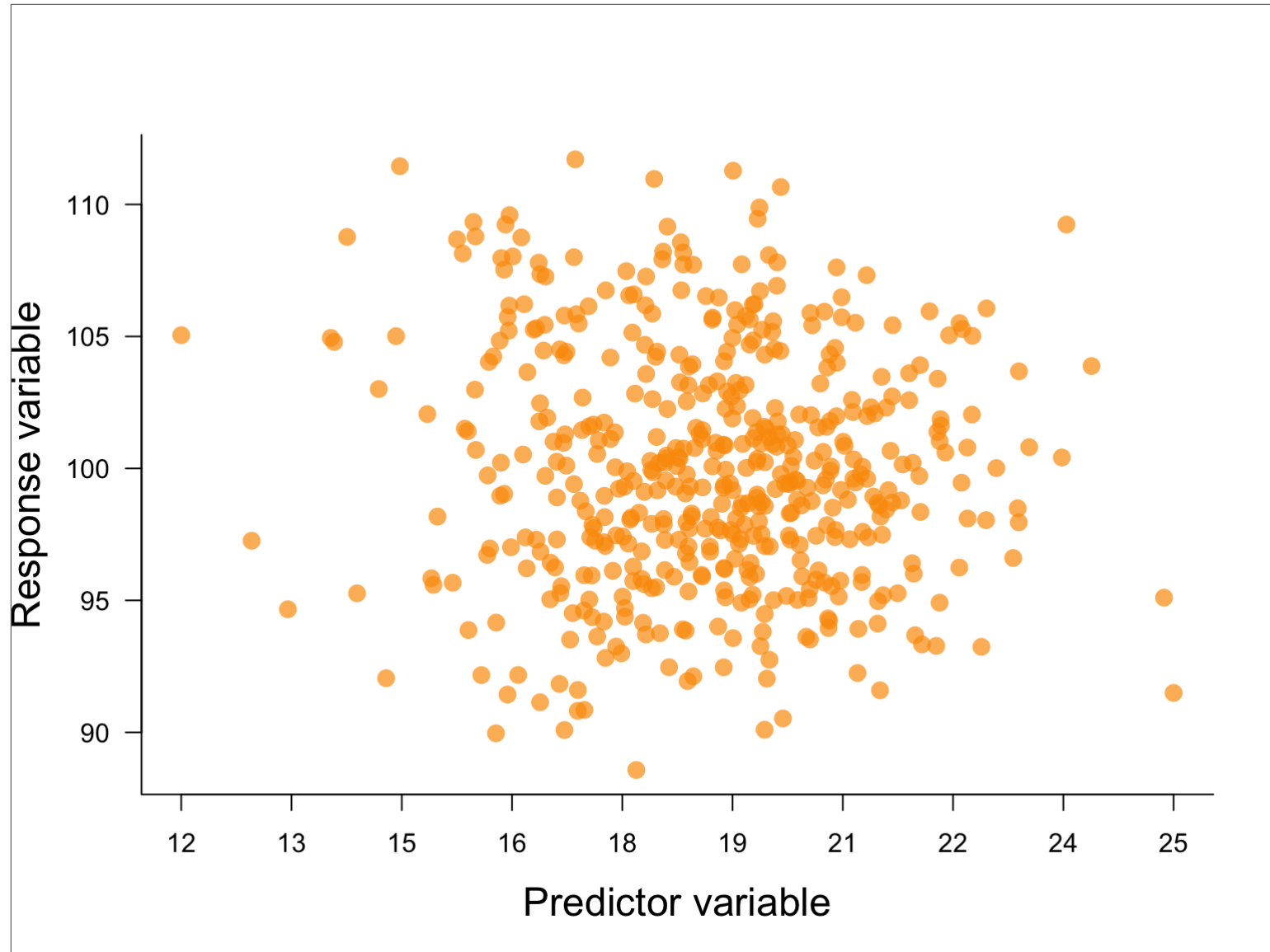


# **MIXED EFFECTS MODELS**

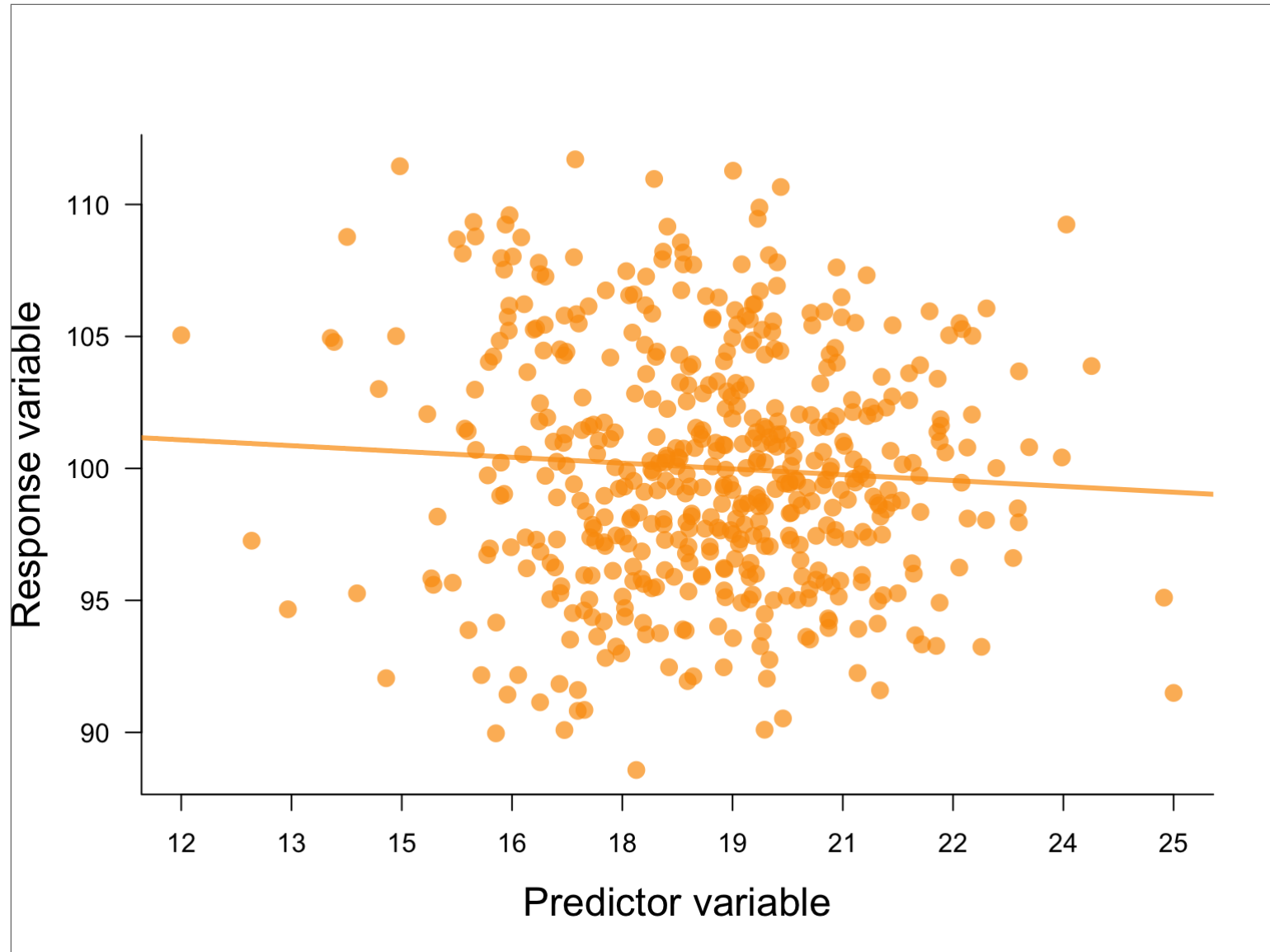
**2 NOVEMBER 2018**

## AN EXAMPLE



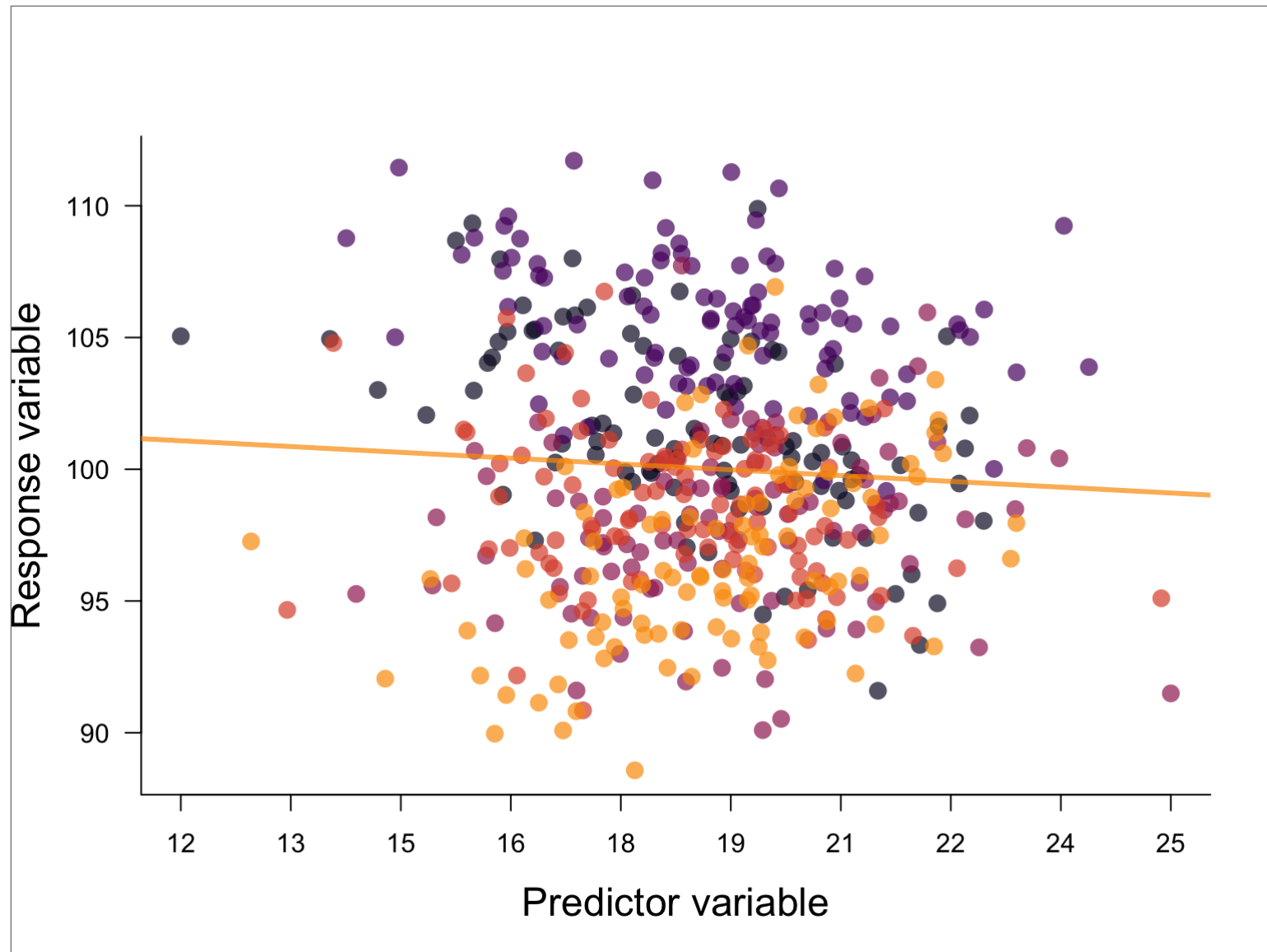


# SINGLE REGRESSION LINE



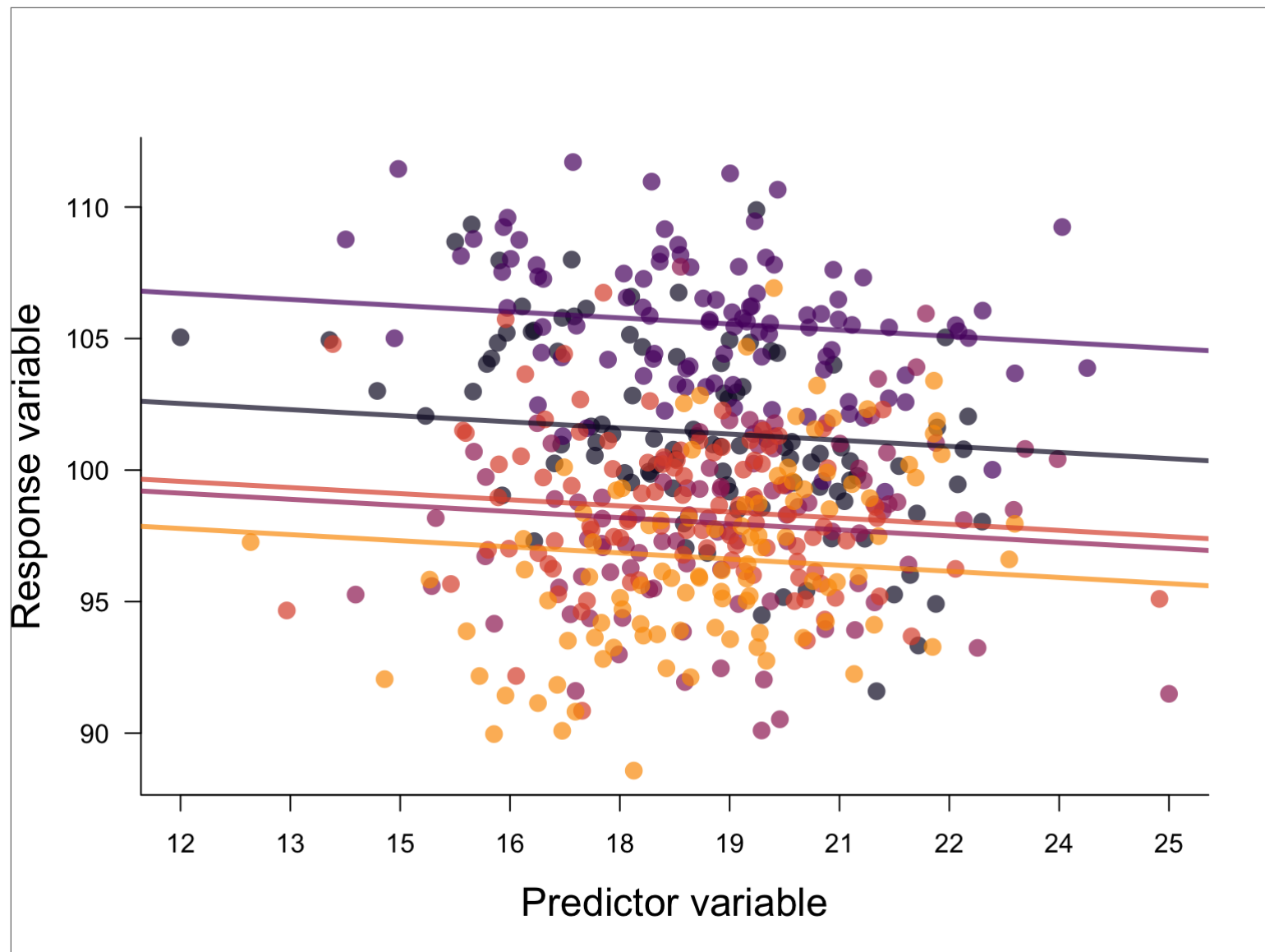


# KNOWN SOURCES OF VARIATION





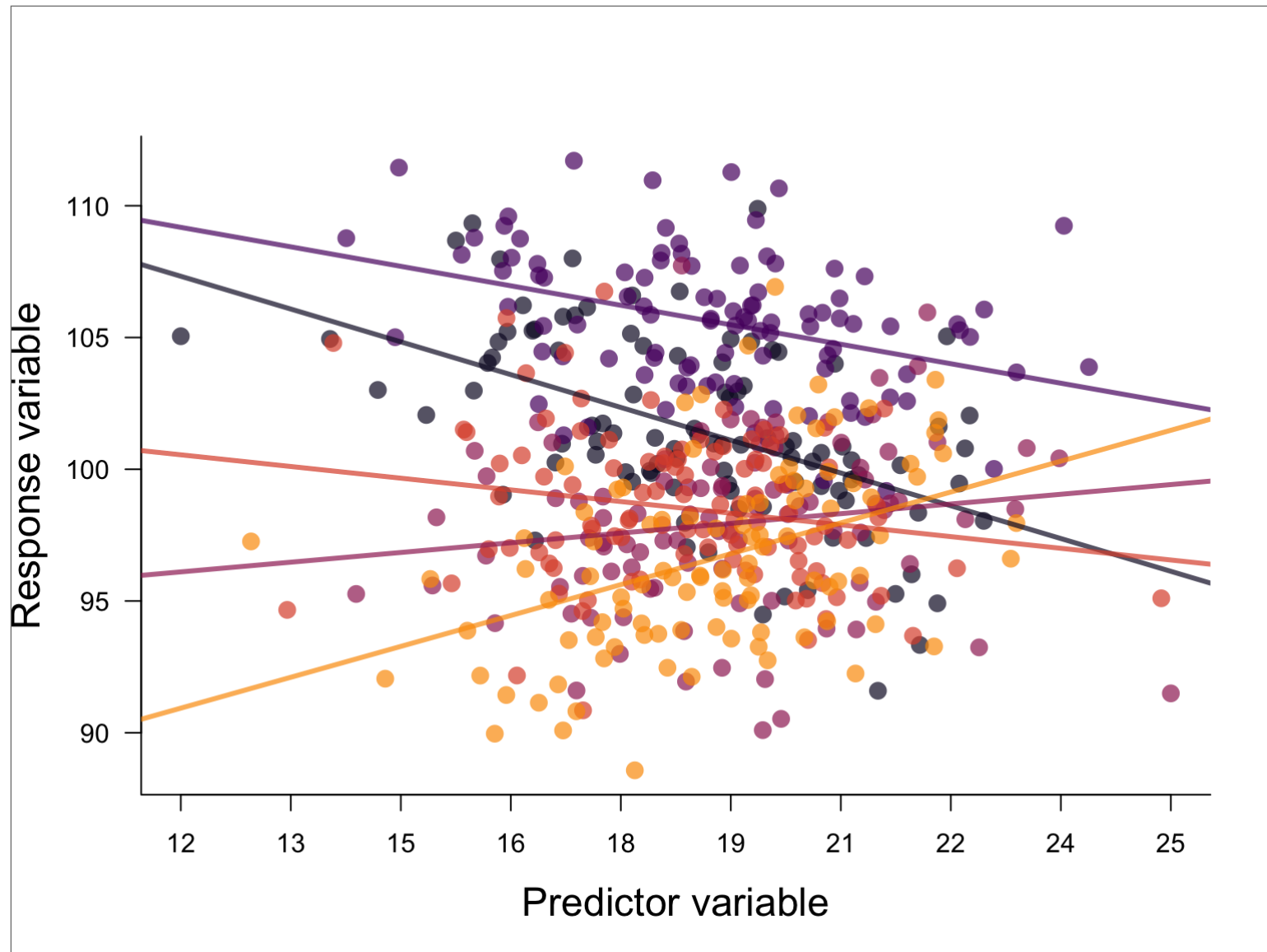
# VARIATION IN INTERCEPTS







# VARIATION IN INTERCEPTS AND SLOPES





## **VARIATION IN PARAMETERS**

- in some cases, variation is directly of interest
- in other cases, it's a nuisance
  - can break independence assumptions
  - can introduce extra noise
- mixed models can help

## MIXED MODELS

- mix of *fixed* and *random* effects
- these terms are not consistently defined
- in this context, really only matters for factors (categorical variables)

## **FIXED EFFECTS**

- these are what we've used in general linear models
  - intercepts
  - slopes
  - interactions
- my definition: categories are independent

# **RANDOM EFFECTS**

- parameters differ among categories but categories aren't fully independent
- some definitions:
  - random if we haven't sampled the entire population
  - random if we “don't care” about the factor
  - random if there is some form of shrinkage

# **RANDOM EFFECTS**

- examples: repeated measures, spatial blocks
  - can be a really good way to account for non-independent observations
- caveat: lme4 methods often require >5 levels for random effects models to work
- be pragmatic (and check model fit!)



## **MIXED MODELS**

- assumptions: much the same as a general linear model
- residuals are independent
- normally distributed residuals
- constant variance of residuals

## **MIXED MODELS IN R**

- use a formula interface to define models

# MIXED MODELS IN R

```
# load lme4 package
library(lme4)

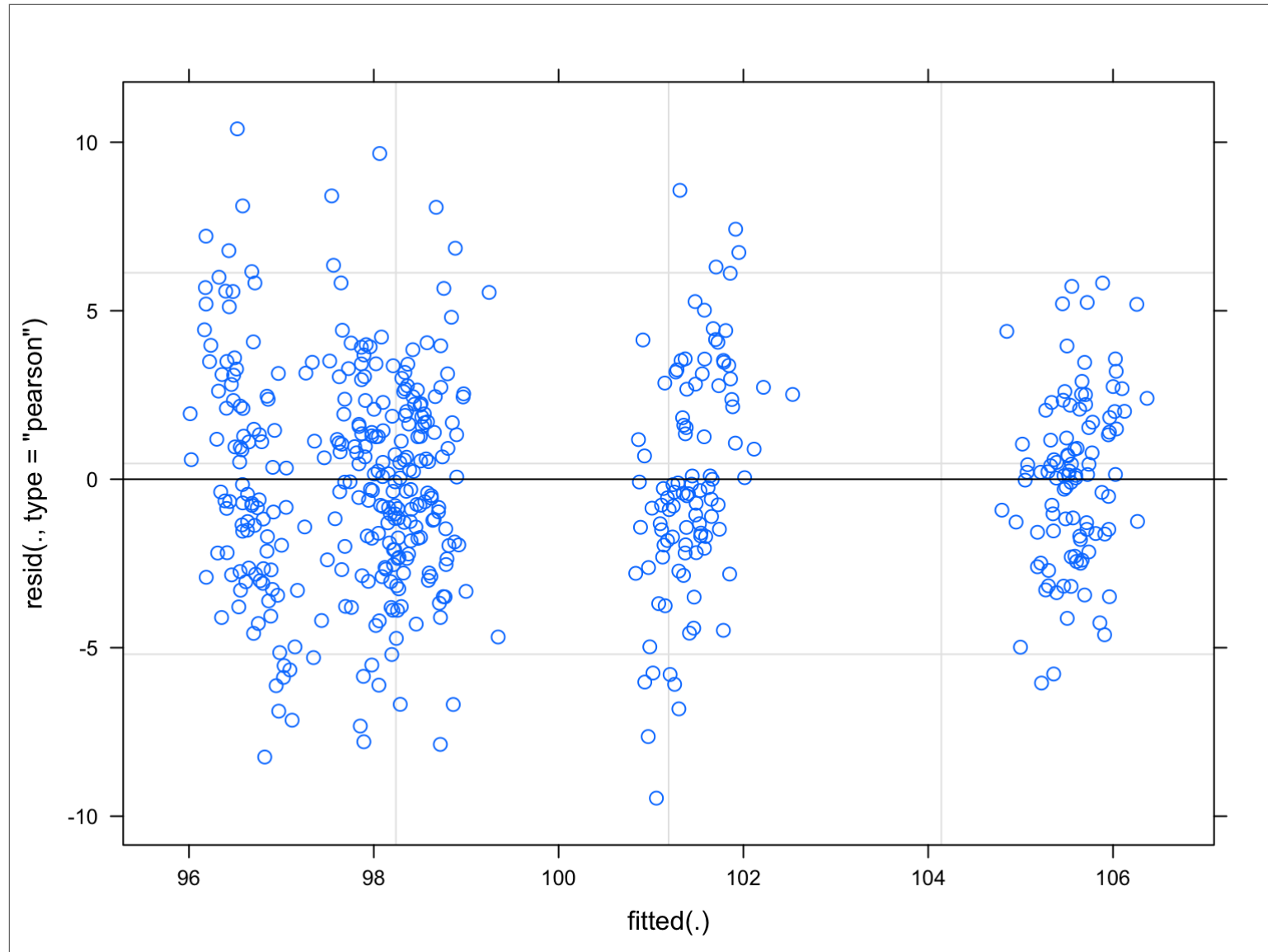
# fit model with single intercept and slope
mod_lm <- lm(response ~ predictor)

# fit model with random intercepts
mod_int <- lmer(response ~ predictor + (1 | block))

# fit model with random intercepts and slopes
mod_slope <- lmer(response ~ predictor + (1 + predictor | block))

# fit model with nested random intercepts
mod_slope <- lmer(response ~ predictor + (1 + predictor | block / nested_block))
```

# PLOT FUNCTION





# SUMMARY FUNCTION

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: y ~ x + (1 | z)
##
## REML criterion at convergence: 2593.1
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.9891 -0.6603 -0.0267  0.6876  3.2835
##
## Random effects:
##   Groups      Name             Variance Std.Dev.
##   z          (Intercept) 12.82      3.580
##   Residual                10.03      3.166
## Number of obs: 500, groups:  z, 5
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept) 100.0136    1.6074  62.222
## x            -0.3119    0.1450  -2.151
##
## Correlation of Fixed Effects:
##   (Intr)
## x 0.004
```

# MIXED MODELS IN R

```
# print the fixed effects  
fixef(mod_int)
```

```
## (Intercept)          x  
## 100.0135534  -0.3118507
```

```
# print the random effects  
ranef(mod_int)
```

```
## $z  
## (Intercept)  
## 1    1.387692  
## 2    5.573009  
## 3   -2.021700  
## 4   -1.574244  
## 5   -3.364757
```





# MIXED MODELS IN R

```
# print the fixed effects  
fixef(mod_slope)
```

```
## (Intercept)          x  
## 99.9562825 -0.2379978
```

```
# print the random effects  
ranef(mod_slope)
```

```
## $z  
## (Intercept)          x  
## 1 1.317356 -1.4299245  
## 2 5.629131 -0.7534335  
## 3 -2.060671 0.7314980  
## 4 -1.563628 -0.3570966  
## 5 -3.322187 1.8089566
```



# INTERPRETING RANDOM EFFECTS

- in short: don't!
- if you care about it, it might be better as a fixed effect
- however, can still look at “variance components”
  - technical term: variance partitioning
- `VarCorr(mod)` is useful for this (but so is `summary(mod)`)

# MODEL ASSESSMENT AND MODEL SELECTION

- many different approaches (see Worksheet 1)
- start by assessing model fit
- but also need to assess model fit *for purpose*
- which model is “best”?
- my approach: often decide on random effects *a priori* and don’t “select” these

