

Introduction

Peter Hoff
Duke STA 610

Multilevel data
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Subpopulation inferences
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Population inferences
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Cross-level inferences
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Multilevel data

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Cross-level inferences

Multilevel data

Multilevel data: Data for which there are

- multiple nested levels of sampling, and/or
- multiple nested sources of variability.

Such data are also often called *hierarchical data* or *clustered data*.

Examples:

Educational testing: students nested within classes;

Small area estimation: households nested within counties;

Agricultural experiments: subplots nested within whole plots;

Clinical trials: measurements nested within patients, patients within hospitals.

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Terminology

observational unit: an object or condition for which data are measured.

macro-level unit: a unit within which other units are nested.

micro-level unit: a unit nested within another unit.

Synonyms:

- macro-level unit, top-level unit, clusters, groups;
- micro-level unit, bottom-level unit, units.

If there are only two levels, we will say *units* are nested within *groups*.

Notation: $y_{i,j}$ = measurement of *i*th unit in *j*th group.

Populations:

- The population: all possible units from all possible groups;
- A subpopulation: all possible units from a single group group.

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Types of multilevel inference

Subpopulation inferences: Group-specific features are of primary interest.

- What is the mean within each group, based on a sample from each group?
- What is the treatment effect for each group?
- Do the groups differ? If so, how do they differ?

Population inferences: Across-group averages are of primary interest.

- What is the population mean, based on cluster sample?
- What is the population treatment effect?

Cross-level inferences: Both types of features are important.

- What is the average treatment effect, adjusting for group differences?

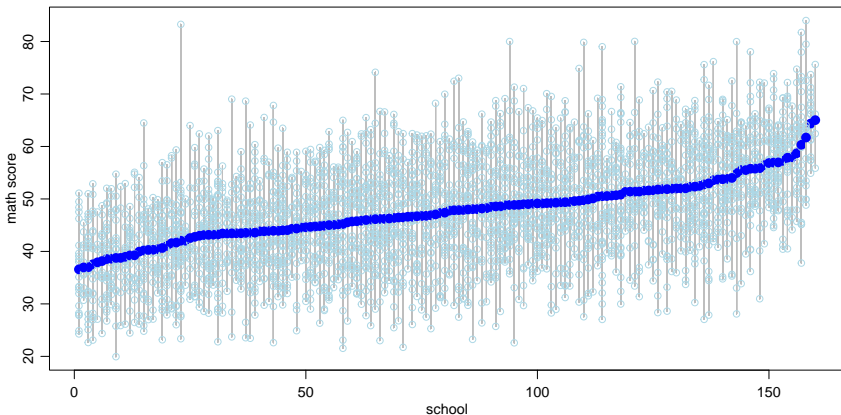
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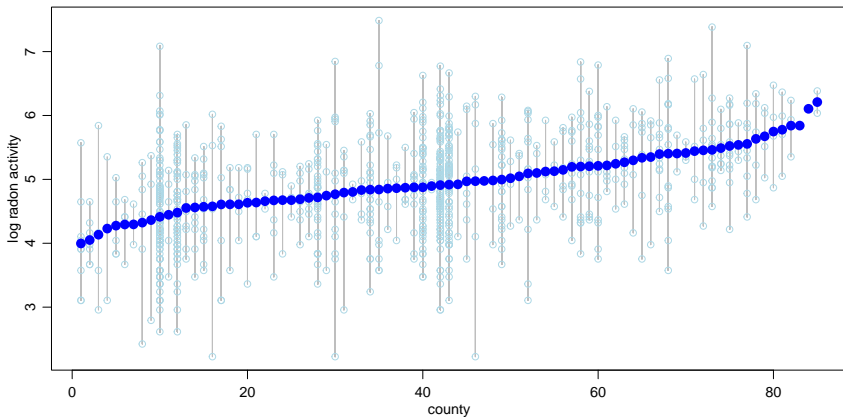
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Example: Educational testing data



Exercise: Identify the population and subpopulations.

Example: Environmental monitoring data



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Group-specific inferences

Targets of inference: Subpopulation means $\theta_1, \dots, \theta_m$.

Data: Subpopulation samples $\{y_{1,1}, \dots, y_{1,n_1}\}, \dots, \{y_{1,p}, \dots, y_{1,n_m}\}$.

Statistical methods:

- Variance tests and estimation: What is $\text{Var}[\theta_1, \dots, \theta_m]$? Is it zero?
- Estimates of θ_j : $\hat{\theta}_j = \bar{y}_{\cdot j}$ or $\hat{\theta}_j = w\bar{y}_{\cdot j} + (1 - w)\bar{y}_{\cdot \cdot}$;
- Confidence intervals: $\Pr(\theta_j \in C(\mathbf{y}) | \theta_j) = 1 - \alpha$, or $\Pr(\theta^* \in C(\mathbf{y})) = 1 - \alpha$;

Cluster sampling

Survey design: Consider the costs of obtaining soil samples from

- 100 randomly sampled locations in a city, versus
- 10 randomly sampled locations from 10 randomly sampled neighborhoods.

Cluster sampling:

The second sampling scheme is called *cluster sampling* or *two-stage sampling*.

Cluster sampling

- is often cheaper per sampled unit;
- often gives less reliable estimates of population means.

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Estimation of a population mean

Task: Estimate the population mean μ from sample data.

Questions:

- How do cluster sampling and SRS compare?
- How do you infer μ from cluster sample data?

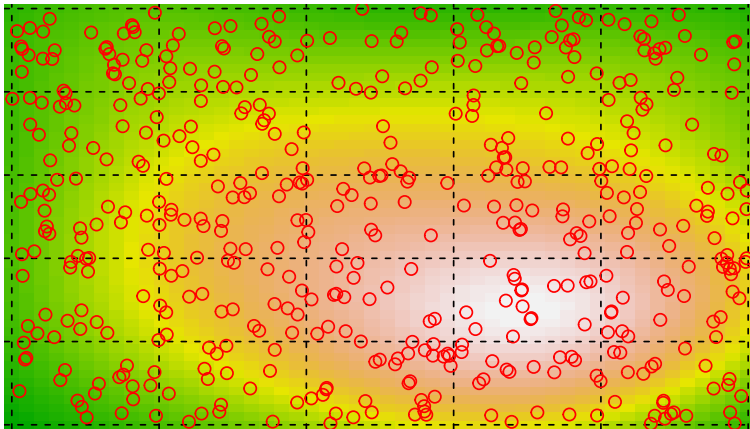
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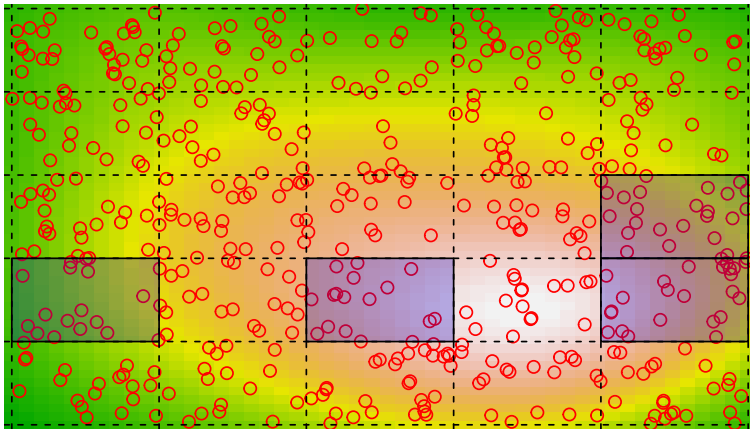
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Two-stage sampling



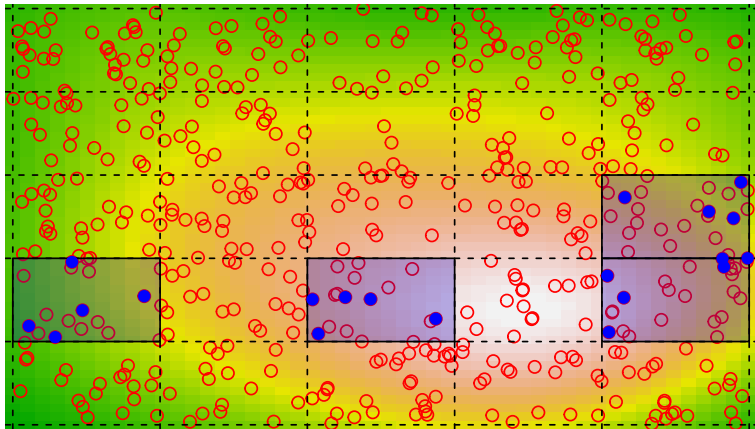
$$\mu=2.0494009$$

Two-stage sampling



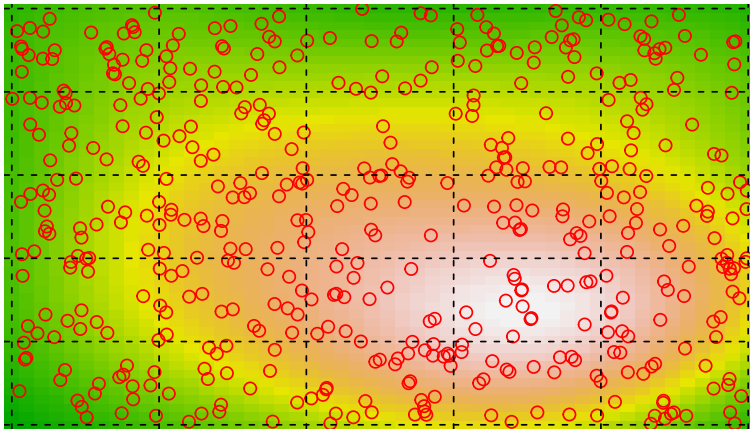
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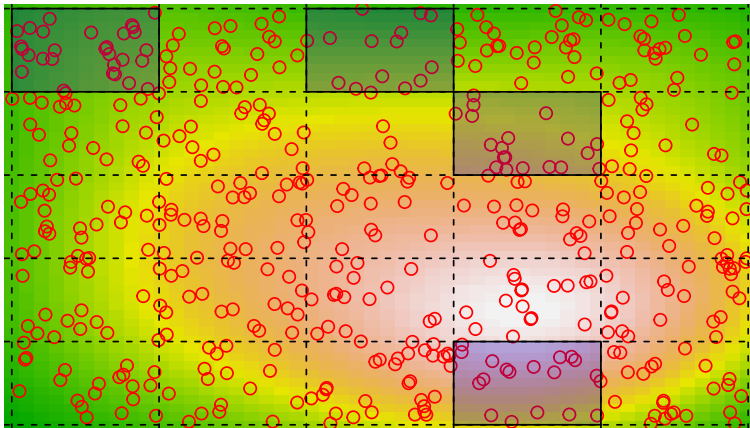
$$\mu=2.0494009 \quad , \quad \bar{y}=2.3547727$$

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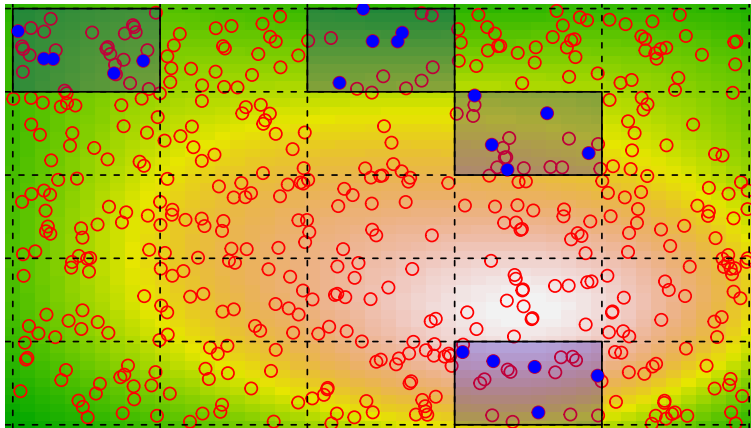
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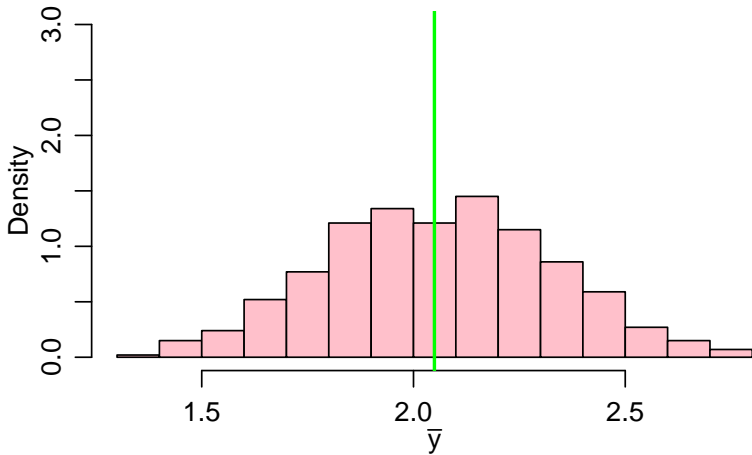
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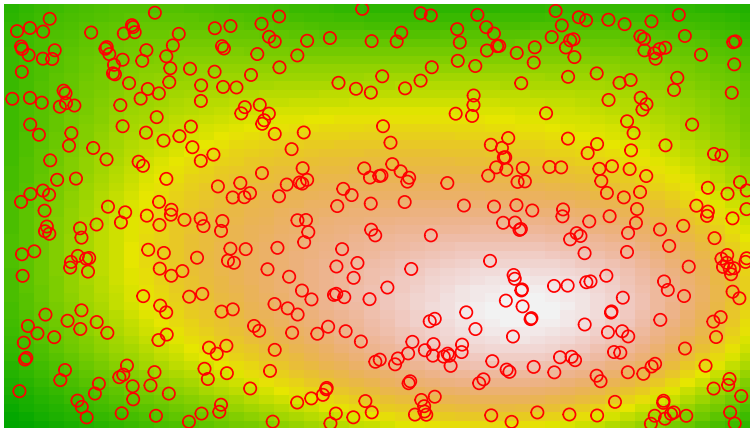


$$\mu=2.0494009 \quad , \quad \bar{y}=1.896463$$

Variability of sample mean

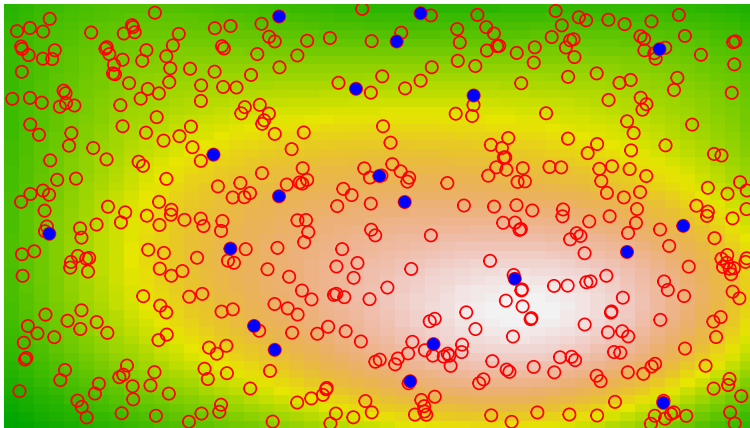


Comparison to SRS



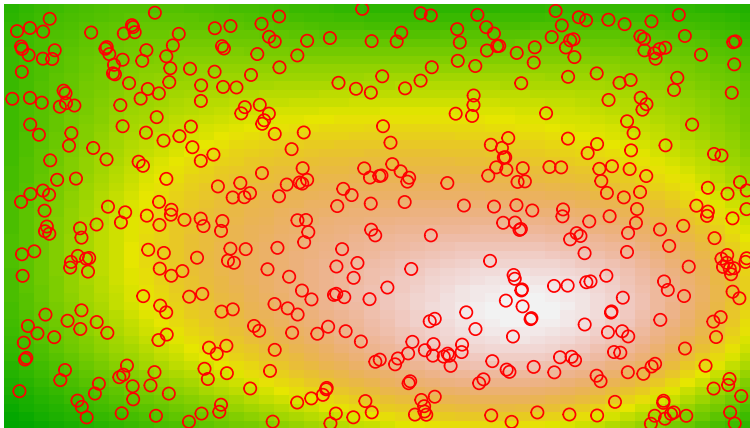
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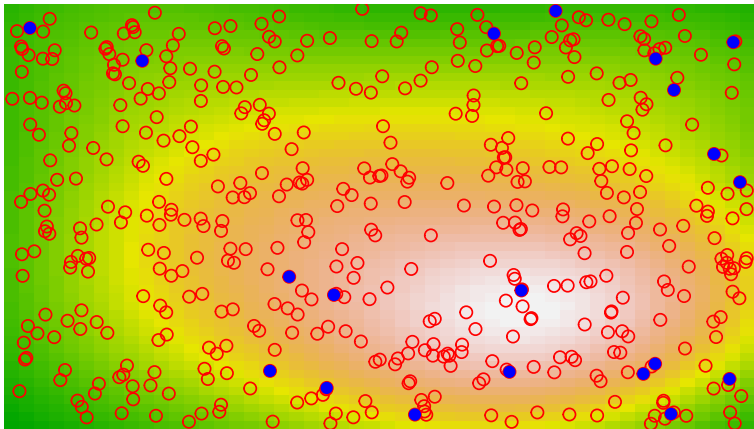
$$\mu=2.0494009 \quad , \quad \bar{y}=2.1696295$$

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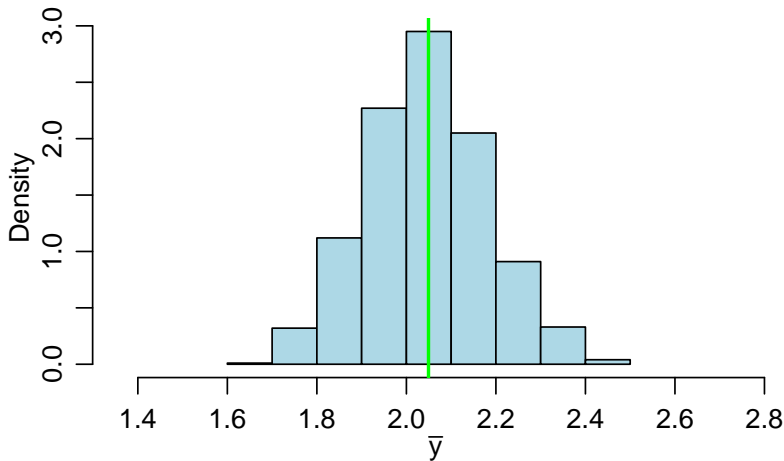
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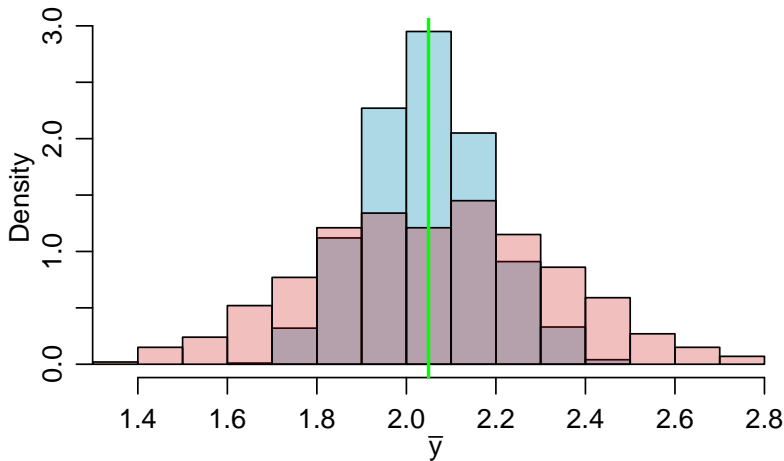


$\mu=2.0494009$, $\bar{y}=1.9804926$

Variability of sample mean



Comparison of sampling variability



Heterogeneity, homogeneity and dependence

As we will show mathematically,

across-group heterogeneity \Leftrightarrow within-group homogeneity
 \Leftrightarrow within-group correlation or dependence

Across-group heterogeneity increases the variance of the sample mean, and so

$$\text{Var}[\bar{y}_{tss}] \geq \text{Var}[\bar{y}_{srs}]$$

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Confidence interval from a SRS

Task: Construct a 95% CI for the population mean.

t-interval for SRS:

If y_1, \dots, y_n is an iid sample with $E[y_i] = \mu$ and $\text{Var}[y_i] = \sigma^2$,

$$E[\bar{y}] = \mu, \text{Var}[\bar{y}] = \sigma^2/n.$$

By the central limit theorem,

$$\bar{y} \sim N(\mu, \sigma^2/n), \quad \frac{\bar{y} - \mu}{\sigma/\sqrt{n}} \sim N(0, 1).$$

As σ^2 is generally unknown, we use

$$\frac{\bar{y} - \mu}{s/\sqrt{n}} \sim t_{n-1}, \quad \text{where } s^2 = \frac{1}{n-1} \sum (y_i - \bar{y})^2.$$

From this, we have

$$\bar{y} \pm t_{n-1, .975} \times s/\sqrt{n} \text{ is a 95\% CI for } \mu.$$

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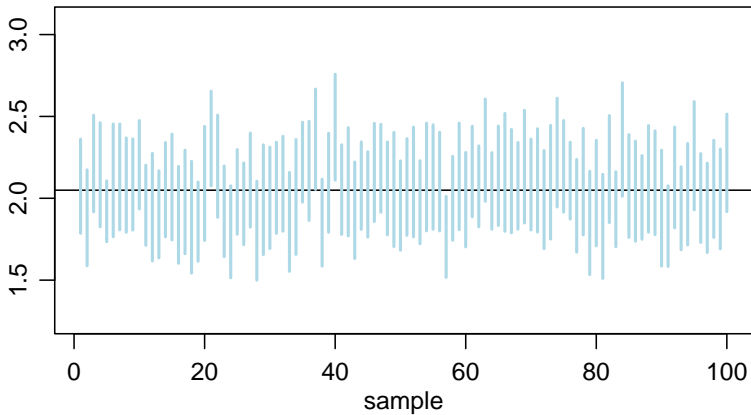
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Frequentist coverage



Ignoring across-group heterogeneity

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What if we apply the formula to data from a cluster sample?

If y_1, \dots, y_n are from a SRS, then

$$\text{Var}[\bar{y}] = \sigma^2/n = E[s^2/n].$$

s/\sqrt{n} provides a good estimate of the sd of \bar{y} .

If y_1, \dots, y_n are from a cluster sample, then generally

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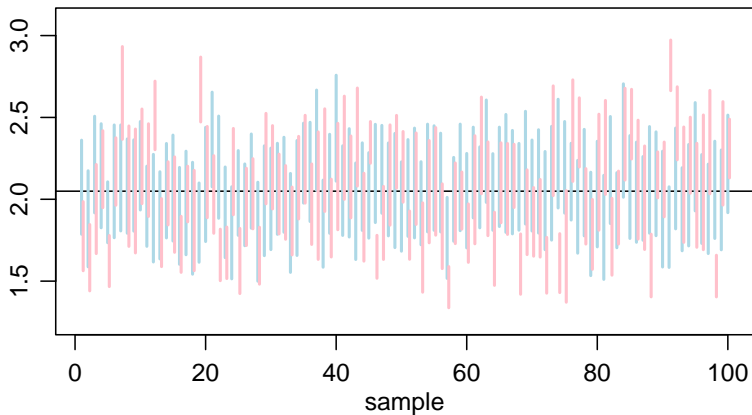
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Ignoring across-group heterogeneity

Summary:

- **Across-group heterogeneity = within-group similarity.**
- Within-group similarity leads to positively correlated cluster sample data.
- The variance of the sample mean from (positively) correlated data is higher than that of the mean of uncorrelated data.
- Statistical inference ignoring such correlation will be inaccurate.

Remedy: We will develop techniques to

- evaluate within- and across-group heterogeneity;
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Estimation of a treatment effect

Suppose

- $x \in \{0, 1\}$
- $\mu_1 = E[y|x = 1]$
- $\mu_0 = E[y|x = 0]$

Task: Estimate the difference $\delta = \mu_1 - \mu_0$ based on cluster sample data.

Data: For each group j , we have $(y_{1,j}, x_{1,j}), \dots, (y_{n,j}, x_{n,j})$.

Question: What could go wrong by ignoring the multilevel nature of the data?

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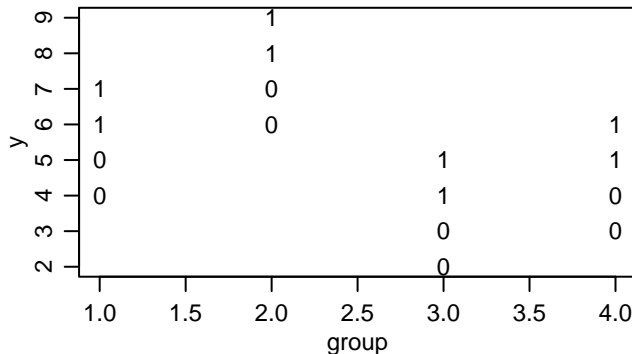
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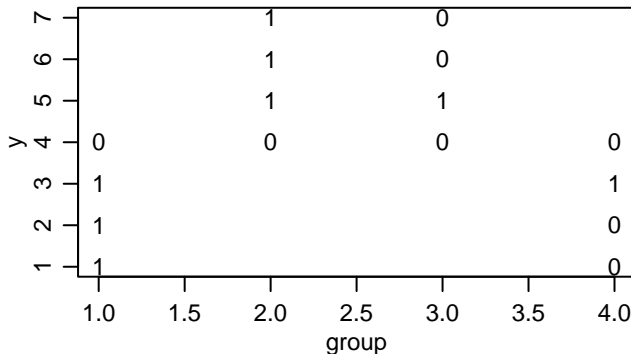
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Overconservative analysis



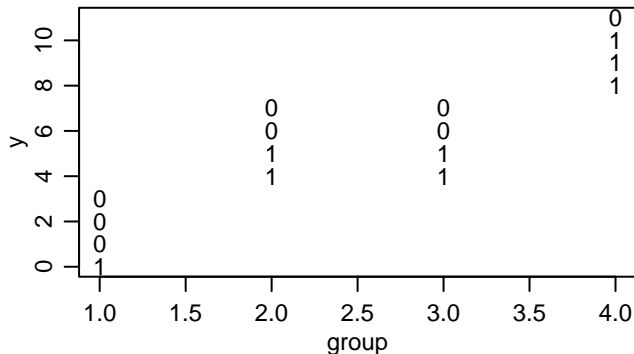
- Overlap across groups, no overlap within groups.
- Across-group variation is *large* compared to the treatment effect.
- Ignoring group differences can lead to *overconservative analysis*.

Underconservative analysis



- The population mean difference is zero.
- The sample mean difference based on pairs of two groups is not zero.
- Ignoring group differences can lead to *underconservative analysis*.

Effect reversal



- $\mu_1 - \mu_0 > 0$ in population, $\mu_{1,j} - \mu_{0,j} < 0$ in every group.
- Within-group effects may be different from population effects.
- This is sometimes called *Simpson's paradox*.

Consequences of across-group heterogeneity

Summary:

- Across-group heterogeneity can lead to *over or under* conservative analysis.
- Population-level effects may be different from group-level effects.
- Data analysis ignoring groups can be inaccurate in *unpredictable* ways.

Remedy: We will develop techniques to

- differentiate between macro and micro level effects;
- appropriately control for within and between-group heterogeneity.

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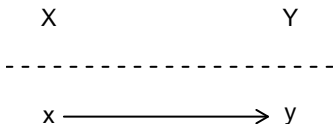
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Macro and micro effects

X , x are macro and micro level explanatory variables

Y , y are macro and micro level outcome variables



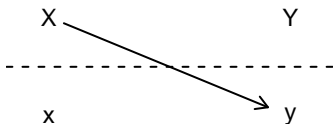
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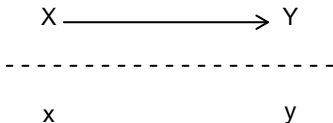
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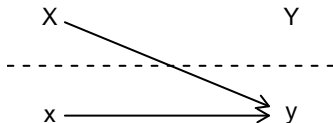


What are the effects of State GDP (X) on statewide political opinion (Y)?
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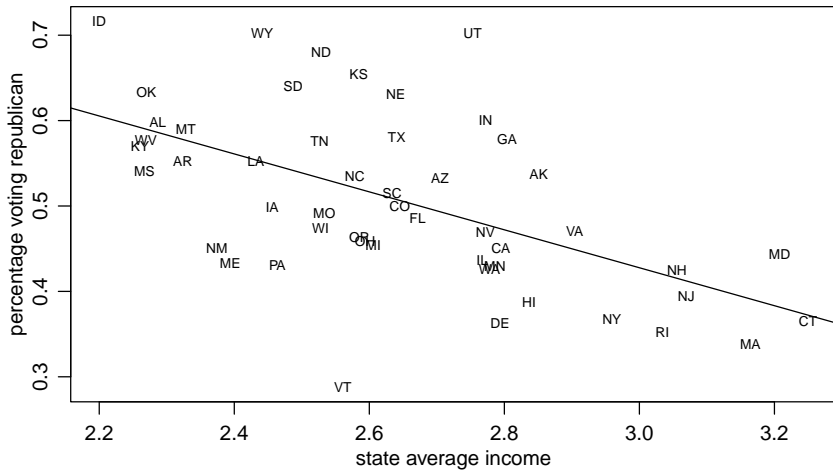
What are the effects of State GDP (X) and SES (x) on political opinion (y)?
(*multilevel effects*)

Example: Income and voting patterns

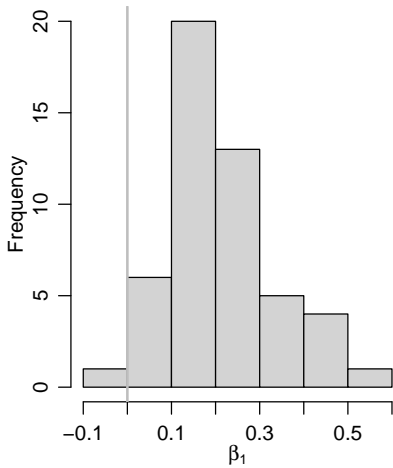
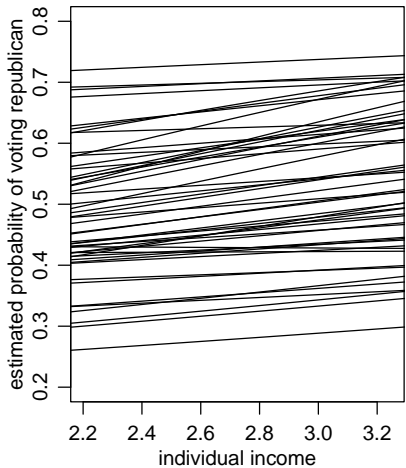
Exit poll data from 2004 presidential election

- $j \in \{1, \dots, 50\}$ indexes the states,
- $y_{i,j}$ is the voting variable for person i in state j ,
- $x_{i,j}$ is a measure of income for person (i, j) .

Macro effects



Micro effects



Joint estimation of effects

In general we may be interested in understanding all of the following:

- **macro level effects,**
- micro level effects,
- macro effects on micro variables,
- heterogeneity of micro effects across groups.

Inference for these items can be made with LME and GLME models:

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