

Interplay Between Science and Statistics

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Common Goals of Statistics

- Scientific Investigation
- Role of Statistics
- Quantifying Distributions
- Comparing Distributions
- Cluster Analysis
- Factor Analysis
- Prediction

Statistical Tasks

- Study Design
- Common Study Designs
- Statistical Analysis

Scientific Investigation

First Stage of Scientific Investigation

- ▶ Hypothesis generation
 - ▶ Observation
 - ▶ Measurement of existing populations or systems
 - ▶ Disadvantages:
 - ▶ Confounding
 - ▶ Limited ability to establish cause and effect

Further Stages of Scientific Investigation

- ▶ Refinement and confirmation of hypotheses
 - ▶ Experiment
 - ▶ Intervention
- ▶ Elements of experiment
 - ▶ Overall goal and specific aims (hypotheses)
 - ▶ Materials and methods
 - ▶ Collection of data
 - ▶ Analysis
 - ▶ Interpretation; Refinement of hypotheses

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Role of Statistics

- ▶ Answering scientific questions in presence of variable response
- ▶ Scientific questions often reduce to comparing the magnitude of some measurement across groups
- ▶ Outcome measures are rarely constant
 - ▶ Inherent randomness
 - ▶ Hidden (unmeasured) variables
- ▶ Use of probability models for describing variability in the real world
 - ▶ Distribution of measurements
 - ▶ Summary measure (functional) for scientific tendency
 - ▶ Quantification of uncertainty in (contrast of) functional(s) (Signal and noise)

“Statistics means never having to say you’re certain.”

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Major goals of statistics typically include

1. Quantifying distributions (or functionals of distributions)
2. Comparison of distributions between groups
3. Identification of clusters
4. Factor analysis
5. Prediction

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1. Quantifying distributions

- ▶ Scientific questions about tendencies for specific measurements within a population
 - ▶ Point estimates of summary measures
 - ▶ Interval estimates of summary measures
 - ▶ Quantifying uncertainty
 - ▶ Decisions about hypothesized values

Example 1: Median life expectancy in stage II breast cancer

- ▶ General goal: Want to know prognosis
- ▶ Follow a cohort of newly diagnosed patients and measure survival time (may be censored)
 - ▶ What defines the cohort?
 - ▶ All patients from a particular healthcare plan?
 - ▶ All patients diagnosed at a particular hospital?
 - ▶ All patients in a certain location?
- ▶ Best estimate of the median survival (??)
- ▶ Quantify uncertainty in that estimate
- ▶ Compare to some clinically important time range (e.g., 10 years)

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1. Quantifying distributions

Example 2: Women suffering from primary biliary cirrhosis

- ▶ Primary biliary cirrhosis: Serious liver disease often leading to liver failure
- ▶ Measure proportion of PBC patients that are women
- ▶ Best estimate of the proportion (What does this mean?)
- ▶ Quantify uncertainty in that estimate
- ▶ Compare to the known proportion of women in the general population (approximately 50%)
- ▶ FYI: About 90% of patients with PBC are women

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2. Comparing distributions across populations

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Can be classified into two main subgroups

2a. *Quantifying differences in the distribution of some measurement across predefined groups (effects or associations)*

2b. *Quantifying differences in effects across subgroups (interactions or effect modification)*

2a. Quantifying differences in the distribution of some measurement across predefined groups (effects or associations)

- ▶ Typically wish to quantify
 - ▶ Existence of differences
 - ▶ Direction of tendency of effect
 - ▶ First, second order relationships in a summary measure
 - ▶ Characterization of dose-response in a summary measure

Example: Effect of serum albumin levels on risk of mortality in end-stage renal disease patients

- ▶ Possible approaches to the analysis:
 1. Compare incidence of mortality across groups of subjects defined by serum albumin
 2. Compare serum albumin level across groups of subjects defined by mortality
- ▶ In either case, comparison can be at many levels of detail regarding nature of differences

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2b. Quantifying differences in effects across subgroups (interactions or effect modification)

- ▶ Typically wish to determine
 - ▶ The existence of interaction
 - ▶ The direction of interaction (synergy, antagonism)
 - ▶ Quantification of exact relationship of interaction

Example: Differential effect of cholesterol on heart attacks by sex?

- ▶ Wish to compare the association between serum albumin level and incidence of mortality between racio-ethnic groups
- ▶ Statistical analysis
 - ▶ Quantify association in Hispanic White
 - ▶ Quantify association in Black
 - ▶ Quantify association in Asian
 - ▶ Quantify association in Non-Hispanic White
 - ▶ Compare measures of association

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3. Cluster analysis

- ▶ Focus is on identifying similar groups of observations
- ▶ Divide a population into subgroups based on patterns of similar measurements
 - ▶ Univariate or multivariate
 - ▶ Known or unknown number of clusters
- ▶ All variables treated symmetrically with no delineation between outcomes and groups

Example: Gene expression of cancerous tumors

- ▶ Dependent upon genetic association, gene expression will be differential between various cancer types
- ▶ Goal: Identify gene expression patterns that separate subpopulations of patients (hopefully by cancer type)
 - ▶ Array of genes (thousands) with corresponding expression levels
 - ▶ Cluster observations into one of K groups to minimize total variability (K-means clustering)
 - ▶ Tabulate cancer type by cluster assignment

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3. Cluster analysis

Example: Gene expression of cancerous tumors

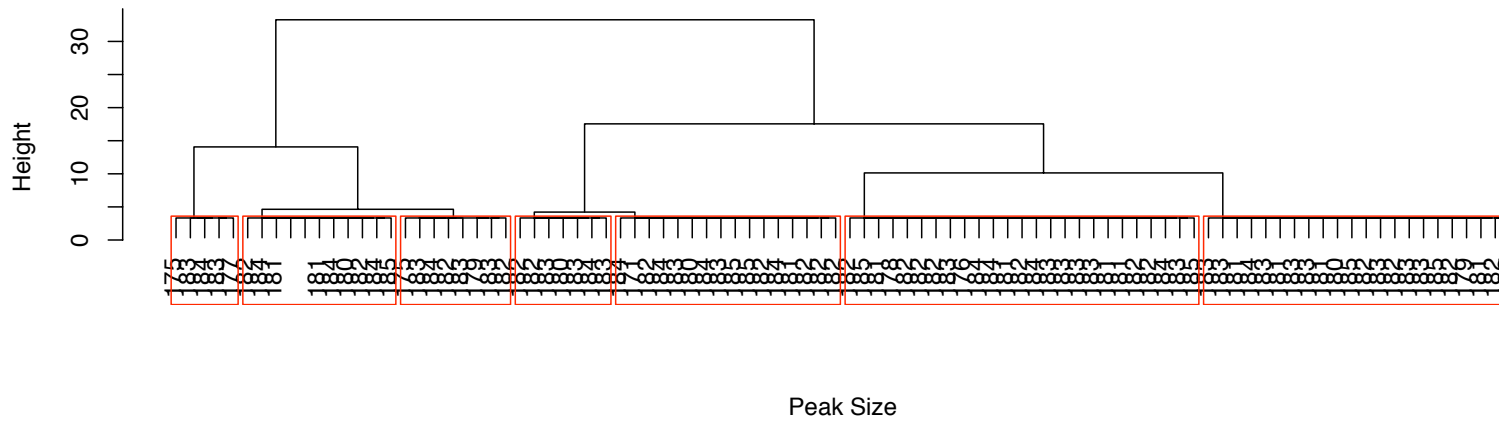
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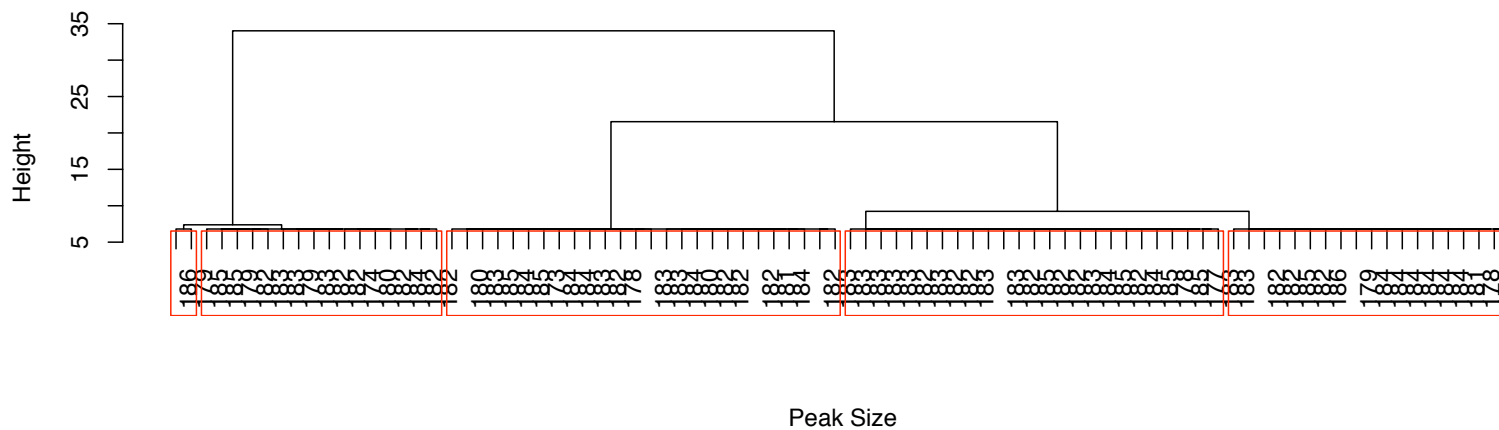
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80.81 : Normal



80.81 : Tumor



4. Factor analysis

- ▶ Identification of hidden variables indicating groups that tend to have similar measurements of some outcome
- ▶ Interest in some particular outcome measurement
- ▶ Predictors that imprecisely measure some abstract quality
- ▶ Desire to find patterns in predictors that more precisely reflect the abstract quality

Example: Barriers to patient compliance in clinical trials

- ▶ In the Health Behavior Questionnaire, multiple variables might be used to measure
 - ▶ Self-perceived health
 - ▶ Social support
 - ▶ Depression
- ▶ Goals:
 1. Find subset of questions that would suffice
 2. Identify hidden variables that affect compliance

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5. Prediction

- ▶ Focus is on future measurements

Point prediction

- ▶ Best single estimate for the measurement that would be obtained on a future observation
 - ▶ Continuous measurements
 - ▶ Binary measurements (discrimination)

Interval prediction

- ▶ Range of measurements that might reasonably be observed for a future observation

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5. Prediction

Example 1 (continuous): Measure of renal function

- ▶ Creatinine is produced by breakdown of creatine, which is used by muscles for energy transfer
- ▶ Removed by kidneys via filtration with little secretion/reabsorption
- ▶ Amount of creatinine cleared by the kidneys in 24 hours used as a measure of blood processed by the kidneys
- ▶ Problem: Need to collect urine (and blood creatinine) for 24 hours
- ▶ Goal: Find blood/urine measures that can be obtained instantly, yet still provide an accurate estimate of a patient's creatinine clearance

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5. Prediction

Example 2 (categorical): Diagnosis of prostate cancer

- ▶ Goal: Use other measurements to predict whether a particular patient might have prostate cancer
 - ▶ Demographic: Age, race, (sex)
 - ▶ Clinical: Symptoms
 - ▶ Biological: Prostate specific antigen (PSA)

Example 3 (interval): Determining normal range for PSA

- ▶ Goal: Identify the range of PSA values that would be expected in 95% of males in the “healthy” population
 - ▶ Typically consider age and race specific values

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Distinctions Without a Difference? No!

- ▶ The largest distinctions in these five goals arise between (1-2) and (3-5)

(1) and (2) are often the focus of hypothesis driven science

- ▶ Testing a well-defined scientific hypothesis
- ▶ Avoidance of data-driven models to guard against inflation of the false positive rate

(3), (4) and (5) necessitate model building and data driven results

- ▶ Allows for the use of much more flexible models
- ▶ Requires the need for stronger assumptions to make probability statements and/or
- ▶ Requires the need for additional *independent* data samples to validate models (this is also the case in hypothesis driven analyses as well)

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Methods

- ▶ Observational study
 - ▶ Data collecting from existing state without intentional interference
- ▶ Interventional study (experiment)
 - ▶ Covariate levels dictated by study researcher

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Time frame

- ▶ Cross-sectional sampling
 - ▶ Real (calendar) time
 - ▶ Event time (e.g. at diagnosis or birth)
- ▶ Longitudinal sampling
 - ▶ Prospective
 - ▶ Follow subjects for occurrence of a specified event
 - ▶ Retrospective
 - ▶ Event has already occurred but exposure measured earlier in time

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Subjects

- ▶ Independent cohorts
 - ▶ Randomly sample groups to be compared
- ▶ Matched groups
 - ▶ Match on potentially important factors that you are not interested in comparing across groups

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Aside: Why match?

- ▶ Usually to control for potential *confounding* factors
 - ▶ Suppose we are interested in the relationship between X (a predictor of interest) and Y (the outcome).
 - ▶ A *confounder* is a third variable Z that is causally associated with X and Y .
- ▶ Lack of adjustment for confounding can lead to a different estimated relationship (unintended) between X and Y .
- ▶ Classic example of confounding
 - ▶ Consider the relationship between alcohol use and the incidence of lung cancer.
 - ▶ Many studies have shown a positive relationship between the two, but why?

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Strength of statistical evidence

- ▶ Sample size determination
- ▶ Statistical power is defined as

$$\Pr[\text{Reject } H_0 | H_0 \text{ false}]$$

where H_0 is some particular hypothesis (eg. “no treatment effect”)

- ▶ Because of variability, power is a function of sample size
- ▶ It is possible for a study to be either *over-* or *under-powered*
 - ▶ A *over-powered* study is designed to detect differences in outcome that are too small to be clinically relevant
 - ▶ “Our sample size ensures that we will have 99% power to detect a difference of 1 day in median survival among newly diagnosed pediatric acute lymphoma cases.”

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Strength of statistical evidence

- ▶ A *under-powered* study has limited ability to reject the null hypothesis for differences in outcome that are clinically relevant
 - ▶ “Our sample size ensures that we will have 50% power to detect a difference of 10-years in median survival among newly diagnosed pediatric acute lymphoma cases.”
- ▶ At best, an over-powered or under-powered study is a waste of resources (money, time, etc.). At worst it can be unethical (consider quality of life, delay in progress of treatment, etc.)

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1. Survey studies

- ▶ Cross-sectional
- ▶ Real (calendar) or event time
- ▶ Efficiency for examining:
 1. Common diseases and risk factors
 2. Associations (not cause and effect)
- ▶ Often limited ability to control for confounding

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1. Survey studies

- ▶ Example Is there an association between ethnicity and health care?
 - ▶ Exposure (ethnicity) and outcome (health care) are easy to measure
 - ▶ How does one choose survey participants?
 - ▶ What about those that refuse to participate? Different from others?
 - ▶ If association is found, do we attempt to explain it?

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2. Cohort studies

- ▶ Groups defined by risk factor
- ▶ Identified prospectively or retrospectively
- ▶ Followed for some outcome event(s)
- ▶ Efficiency for examining
 1. Common diseases
 2. Many different outcomes for same exposure
 3. Associations (not cause and effect)

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2. Cohort studies

- ▶ Example: Is renal impairment among middle aged individuals associated with an increased risk of cardiovascular disease (CVD)?
 - ▶ Measure renal function on subjects between age 35 and 45 with no history of (CVD) and follow until first (CVD) event
 - ▶ Could also consider the relationship between renal function and death
 - ▶ If an association is found, how could it not be causal?

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3. Case-control studies

- ▶ Groups defined by some outcome event
- ▶ Characterize prior exposures
- ▶ Efficiency for examining
 1. Rare diseases
 2. Many different risk factors for same disease
 3. Associations (not cause and effect)

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3. Case-control studies

- ▶ Example: Risk factors for childhood leukemia
 - ▶ Randomly sample children diagnosed with leukemia at time of diagnosis (cases)
 - ▶ Randomly sample children without leukemia (controls)
 - ▶ Compare multitude of risk factors from environment to genetic profile
 - ▶ Careful though must go into the choice of controls in order to make a fair comparison
 - ▶ Possibly match to reduce confounding

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4. Interventional (clinical trials)

- ▶ Ideally controlled and randomized
- ▶ Efficiency for examining
 - ▶ Common outcomes
 - ▶ Cause and effect

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4. Interventional (clinical trials)

- ▶ Example Does recombinant human thrombin (rhThrombin) reduce incidental bleeding during spinal surgery?
 - ▶ Randomize patients to receive rhThrombin (applied via sponge) or a *placebo* consisting of a saline solution
 - ▶ *Double-blind* study participants so that the patient nor the surgeon knows what is being applied
 - ▶ Measure time from application of sponge until bleeding stops
 - ▶ Note: A *single-blind* study would only blind the patient and not the surgeon
 - ▶ How could this lead to a *placebo effect*?

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What are we able to estimate under each of these study designs?

1. Cross-sectional design

- ▶ Sample from population
- ▶ Measure exposure (E), disease (D)
- ▶ Can estimate:
 1. $\Pr[E, D]$: Joint distn of Exposure, Disease
 2. $\Pr[D|E]$: Conditional distn of D within levels of E
 3. $\Pr[E|D]$: Conditional distn of E within levels of D
 4. $\Pr[D]$: Marginal distn (*prevalence*) of Disease
 5. $\Pr[E]$: Marginal distn of exposure

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What are we able to estimate under each of these study designs?

2. Cohort design

- ▶ Fix sample sizes for each level of exposure (E)
- ▶ Measure disease (D)
- ▶ Can estimate:
 1. $\Pr[D|E]$: Conditional distn of D within levels of E

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What are we able to estimate under each of these study designs?

3. Case-control design

- ▶ Fix sample sizes for each level of disease (D)
- ▶ Measure exposure (E)
- ▶ Can estimate:
 1. $\Pr[E|D]$: Conditional distn of E within levels of D

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What are we able to estimate under each of these study designs?

4. Interventional design (clinical trial)

- ▶ Fix sample sizes for each level of disease (E)
- ▶ Measure disease (D)
- ▶ (so a cohort design)
- ▶ Can estimate:
 1. $\Pr[D|E]$: Conditional distn of D within levels of E

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Key point

- ▶ Each can be used for testing an association between E and D :
 - ▶ $H_0 : \Pr[D, E] = \Pr[D] \times \Pr[E]$ (cross sec)
 - ▶ $H_0 : \Pr[D|E] = \Pr[D|\bar{E}]$ (cohort)
 - ▶ $H_0 : \Pr[E|D] = \Pr[E|\bar{D}]$ (case-control)
- ▶ What about parameter interpretation?

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Description of a sample

- ▶ Identification of measurement or data entry errors
- ▶ Characterization of materials and methods
- ▶ Validity of analysis methods
- ▶ Hypothesis generation (for inference and estimation)

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Inference

- ▶ Goal: Generalize from sample to population
- ▶ Two main components:
 1. Estimation
 - 1a. Point estimation
 - 1b. Interval estimation
 2. Testing

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Inference

1a. Point estimation

- ▶ Identification of clusters
- ▶ Point estimates
- ▶ Individual observations (predictions)
 - ▶ Continuous measurements
 - ▶ Categorical measurements (discrimination, classification)
- ▶ Summary measures of distributions
 - ▶ Within a population
 - ▶ Across populations

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Inference

1b. Interval estimates (quantifying uncertainty)

- ▶ Individual observations
 - ▶ Prediction intervals (continuous measurements)
 - ▶ Accuracy (discrimination, classification)
- ▶ Summary measures
 - ▶ Confidence or credible intervals

2. Decisions (hypothesis testing)

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