

FUNDAMENTAL CONCEPTS IN MEDICAL STATISTICS

... with applications to randomized controlled trials

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Makerere

7-13 June 2023



COURSE HOME PAGES

<https://gjessing.super.site> + <https://muele.mak.ac.ug>



Fundamental concepts in medical statistics with applications to randomized controlled trials, Kampala 7-13 June 2023

COURSE RESOURCE PAGE

Organizers



Makerere University
College of Health Sciences
SCHOOL OF PUBLIC HEALTH



Victoria Nankabirwa
Lecturer



Halvor Sommerfelt
Professor

KAMPALA, 1993



UNIVERSITY MAIN BUILDING, 1993



FERTILITY AND HEALTH

UNIVERSITY GUEST HOUSE, 1993



AT THE MUSEUM, 1993



ABOUT THIS COURSE...

- More on principles & ideas rather than recipes
- The exercises will support with hands on practical examples
- Selection of topics mostly based on personal interests (and what I had available)
- but it should all be very relevant

SCHEDULE: WEDNESDAY, 7 JUNE

▼ Wednesday 7

FOCUS OF THE DAY: Overview, and fundamental concepts

▼ 09:00 - 10:00 Welcome and startup

- ▶ Welcome/introduction round
- ▶ Quick overview over course + practical info
- ▶ Quick overview over some basic stuff (but note that participants are expected to have a basic understanding of/experience with basic statistical concepts)

▼ 10:00 - 12:00 Some fundamentals

- ▶ Central limit theorem (most importantly, why it's useful)
- ▶ General background on Confidence Intervals
- ▶ Testing, p-values, one-sided vs. two-sided
- ▶ Confidence intervals vs. testing
- ▶ Transformation of data, the log-transform. General discussion of measurement scales

▶ 12:00 - 13:00 LUNCH

▶ 13:00 - 14:00 Exercises

▼ 14:00 - 15:00 Sample size and statistical power / statistical precision

- ▶ What it means
- ▶ How to restrict the probability of Type I/Type II errors, i.e. achieve sufficient statistical power at a given test significance level
- ▶ ... or sometimes better, compute sample size needed for sufficient precision of your effect estimate
- ▶ How to compute sample size for study planning, simple situations

▶ 15:00 - 16:00 Exercises

SCHEDULE: THURSDAY, 8 JUNE

▼ Thursday 8

FOCUS OF THE DAY: Generalized linear models (GLMs)

▼ 09:00 - 11:00 Linear regression

- ▶ Standard linear regression
- ▶ Multiple linear regression, confounding, interpretation
- ▶ Basic regression diagnostics

▶ 11:00 - 12:00 Exercises

▶ 12:00 - 13:00 LUNCH

▼ 13:00 - 15:00 Binomial regression

- ▼ Binomial regression (including logistic regression)
 - ▶ Different link functions: logit, log, and identity
 - ▶ Corresponding interpretations: logistic, log, and additive
 - ▶ Odds Ratio (OR), Relative Risk (RR), and Risk Difference (RD)
 - ▶ Basic regression diagnostics
- ▶ Poisson regression (Introduced here as a GLM, but will for the most part be covered on Monday, in the context of event history analyses.)

▶ 15:00 - 16:00 Exercises

SCHEDULE: FRIDAY, 9 JUNE

▼ Friday 9

FOCUS OF THE DAY: RCTs, other experimental designs

▼ 09:00 - 10:00 Fundamentals

- ▶ The causal reasoning behind randomization
- ▶ Adjustment for baseline variables? Other adaptations.
- ▶ Regression to the mean, need for trial comparator arm, etc.
- ▶ Dropouts, intention to treat

▼ 10:00 - 11:00 Multiple testing, multiple estimation

- ▶ Problems with multiple testing
- ▶ "Winner's curse" in multiple estimation
- ▶ p-hacking, multiple subgroups, etc.
- ▶ Some extreme examples from genetics

▶ 11:00 - 12:00 Exercises

▶ 12:00 - 13:00 LUNCH

▼ 13:00 - 13:30 Weighted analyses

- ▼ Various types of weighting
 - ▶ Sampling
 - ▶ Frequency
 - ▶ Importance (although not really important here)

▼ 13:30 - 15:00 Interaction / effect measure modification

- ▶ Basic concepts
- ▶ Scale dependence
- ▶ Examples
- ▶ How to estimate, variable coding, such as Stata's approach, and dummy variables
- ▶ Testing / estimating differences in effects across subgroups

▶ 15:00 - 16:00 Exercises

SCHEDULE: SATURDAY, 10 JUNE

▼ Saturday 10

FOCUS OF THE DAY: Self study/group work

SCHEDULE: MONDAY, 12 JUNE

▼ Monday 12

FOCUS OF THE DAY: Count data, standardization

▼ 09:00 - 11:00 Poisson regression

- ▶ How and when to use
- ▶ How to handle varying exposure time, use of offset
- ▶ Examples from Covid-19 pandemic analyses

▶ 11:00 - 12:00 exercises

▶ 12:00 - 13:00 LUNCH

▼ 13:00 - 15:00 Standardization methods

- ▶ Standardized averages for continuous data
- ▶ Standardized risks and rates
- ▶ Standardized Mortality Ratios (SMRs)
- ▶ Model-based standardization, examples of how to present results

▶ 15:00 - 16:00 Exercises

SCHEDULE: TUESDAY, 13 JUNE

▼ Tuesday 13

FOCUS OF THE DAY: Survival and event history analysis. Recap of central topics from the course.

▼ 09:00 - 11:00 Survival Analysis / Event History Analysis

- ▶ Introduction to survival/event history analysis
- ▶ Kaplan-Meier
- ▶ Cox regression

▶ 11:00 - 12:00 Exercises

▶ 12:00 - 13:00 LUNCH

▼ 13:00 - 14:00 Additional topics in survival analysis

- ▶ Alternative models, such as additive hazards model
- ▶ Competing risk

▶ 14:00 - 15:00 Summary, discussion, evaluation

Why Medical Statistics?

New England Journal of Medicine

Editorial, Jan. 6, 2000, p. 42-49

The eleven most important developments in medicine over the past millennium

- Elucidation of human anatomy and physiology
- Discovery of cells and their substructures
- Elucidation of the chemistry of life
- *Application of statistics to medicine*
- Development of anesthesia
- Discovery of the relation of microbes to disease
- Elucidation of inheritance and genetics
- Knowledge of the immune system
- Development of body imaging
- Discovery of antimicrobial agents
- Development of molecular pharmacotherapy

Application of statistics to medicine

“Sir David Cox’s 1972 paper on proportional-hazards regression ignited the fields of survival analysis and semiparametric inference (using partial specification of the probability distribution of the outcomes under investigation). Rapid improvements in computer support were essential to the growing role of empirical investigation and statistical inference.”

TOTALLY INDECENT SELF-PROMOTION

Statistics for Biology and Health

Odd O. Aalen, Ørnulf Borgan and Håkon K. Gjessing
Survival and Event History Analysis

Time-to-event data are ubiquitous in fields such as medicine, biology, demography, sociology, economics and reliability theory. Recently, a need to analyse more complex event histories has emerged. Examples are individuals that move among several states, frailty that makes some units fail before others, internal time-dependent covariates, and the estimation of causal effects from observational data.

The aim of this book is to bridge the gap between standard textbook models and a range of models where the dynamic structure of the data comes to its full right. The common denominator of such models is stochastic processes. The authors show how counting processes, martingales, and stochastic integrals fit very nicely with censored data. Beginning with standard analyses such as Kaplan-Meier plots and Cox regression, the presentation progresses to the additive hazard model and recurrent event data. Stochastic processes are also used as natural models for individual frailty; they allow sensible interpretations of a number of surprising findings seen in population data.

The stochastic process framework is naturally connected to causality. The authors show how dynamic path analysis can incorporate many modern causality ideas in a framework that takes the time aspect seriously.

To make the material accessible to the reader, a large number of practical examples, mainly from medicine, are developed in detail. Stochastic processes are introduced in an intuitive and less technical manner. The book is aimed at investigators who use event history methods and want a better understanding of the statistical concepts. It is suitable as a textbook for graduate courses in statistics and biostatistics.

Odd O. Aalen is professor of medical statistics at the University of Oslo, Norway. He PhD from the University of California, Berkeley in 1975 introduced counting processes and martingales in event history analysis. He has also contributed to numerous other areas of event history analysis, such as additive hazards regression, frailty, and causality through dynamic modelling.

Ørnulf Borgan is professor of statistics at the University of Oslo, Norway. Since his PhD in 1986 he has contributed extensively to event history analysis. He is an author of the monograph *Statistical Models Based on Counting Processes*, and is editor of *Scandinavian Journal of Statistics*.

Håkon K. Gjessing is professor of medical statistics at the Norwegian Institute of Public Health and the University of Bergen, Norway. Since his PhD in epidemiology in 1995, he has worked on a broad range of theoretical and applied problems in biostatistics.

STATISTICS | LIFE SCIENCES,
MEDICINE, HEALTH SCIENCES



springer.com

Aalen · Borgan · Gjessing



Survival and Event History Analysis

Odd O. Aalen
Ørnulf Borgan
Håkon K. Gjessing

Survival and Event History Analysis

A Point Process View

Springer

A digression into Statistical Software

- R
 - Many users find RStudio helpful. Free, open-source version
<https://posit.co/products/open-source/rstudio/>
 - Some simple GUIs are also available
<https://r4stats.com/2022/06/20/updated-comparison-of-r-guis/>
- Stata
 - Stata commands, manuals, and documentation are in general of
— AMAZING quality! —
- SPSS (Statistical Pseudo Science System)
- SAS
- Python
- Julia



wizard



=



muggle

WHAT TYPE OF “RESTAURANT” IS YOUR FAVORITE SOFTWARE?

- STATA
 - Michelin restaurant
 - Excellent, detailed menu and top quality
 - Can meet the cooks or attend cooking classes

WHAT TYPE OF “RESTAURANT” IS YOUR FAVORITE SOFTWARE?

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- R

- YOU ARE THE COOK...
- All the tools & ingredients are there for you to use
- The sky is the limit...
- (...or it all ends in a total mess)

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- McDonald's

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- McDonald's
- Motto: “More than just Burgers & Fries!”

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 - All the tools & ingredients are there for you to use
 - The sky is the limit...
 - (...or it all ends in a total mess)
- SPSS
 - McDonald's
 - Motto: “More than just Burgers & Fries!”
- How about SAS, then?

YES, HOW ABOUT SAS, THEN?



NTRE FOR
FERTILITY AND HEALTH

Excerpt from an R help page for the frequently used *apply* function

apply

If each call to FUN returns a vector of length n, then apply returns an array of dimension $c(n, \text{dim}(X)[\text{MARGIN}])$ if $n > 1$. If n equals 1, apply returns a vector if MARGIN has length 1 and an array of dimension $\text{dim}(X)[\text{MARGIN}]$ otherwise. If n is 0, the result has length 0 but not necessarily the “correct” dimension.

If the calls to FUN return vectors of different lengths, apply returns a list of length $\text{prod}(\text{dim}(X)[\text{MARGIN}])$ with dim set to MARGIN if this has length greater than one.

In all cases the result is coerced by `as.vector` to one of the basic vector types before the dimensions are set, so that (for example) factor results will be coerced to a character array.

R: WONDERFULLY CLEAR & PEDAGOGICAL HELP PAGES

Excerpt from an R help page for the frequently used *apply* function

apply —

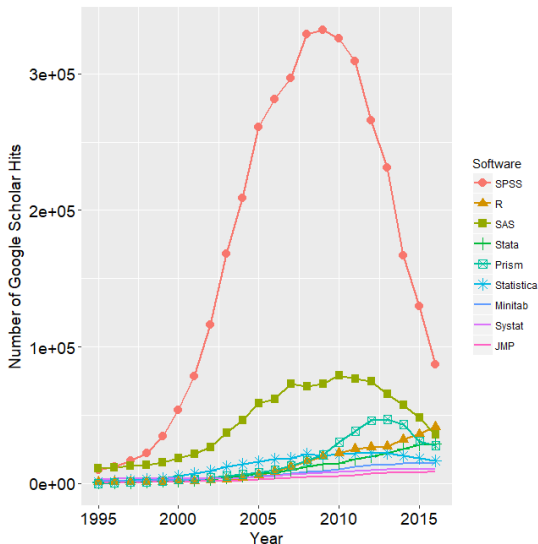
If each call to FUN returns a vector returns an array of dimension $c(n, d_1, \dots, d_m)$. If n equals 1, apply returns a vector and an array of dimension $\text{dim}(X)$ [MARGIN]. If the result has length 0 but not necessarily dimension.

If the calls to FUN return vectors of length n , apply returns a list of length $\text{prod}(\text{dim}(X))$ [MARGIN] if this has length greater than 1.

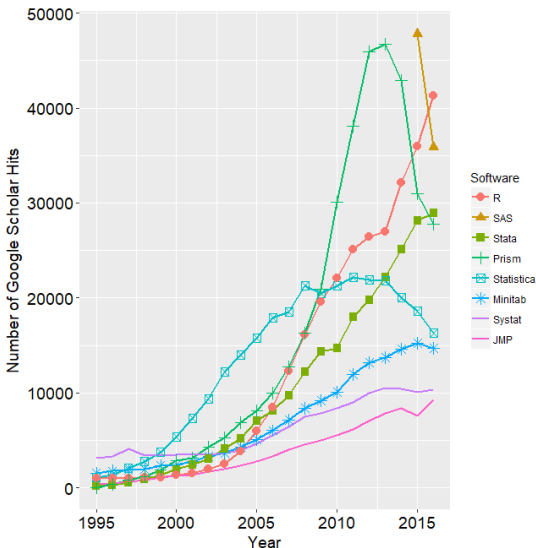
In all cases the result is coerced by FUN to the basic vector types before the dimensionality is determined. For example, factor results will be coerced to character array.



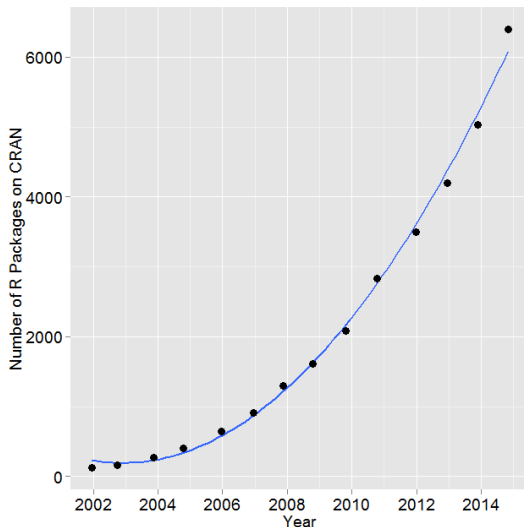
GOOGLE SCHOLAR HITS, BY SOFTWARE ([HTTP://R4STATS.COM/POPULARITY](http://r4stats.com/popularity))



GOOGLE SCHOLAR HITS, BY SOFTWARE (MINUS SPSS, SAS) (HTTP://R4STATS.COM/POPULARITY)



NUMBER OF R PACKAGES ON CRAN ([HTTP://R4STATS.COM/POPULARITY](http://r4stats.com/popularity))

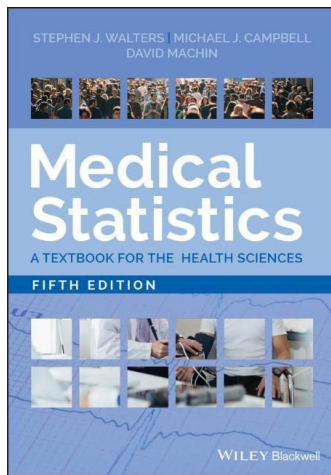


- Python

- (Closer to) a compiled language
- Inherently faster, strong on very large data files
- Suited for larger software development projects
- Two weeks' course in Python → “Data Scientist”(??)

- Julia

- A “modernized” R
- Just-In-Time (JIT) compiled
- Faster and better structured than R
- Can use R, Python and other libraries
- Will it catch on?



- Pretty decent
- Wide (but shallow) coverage of important topics
- Good as a reference text

DATA TYPES

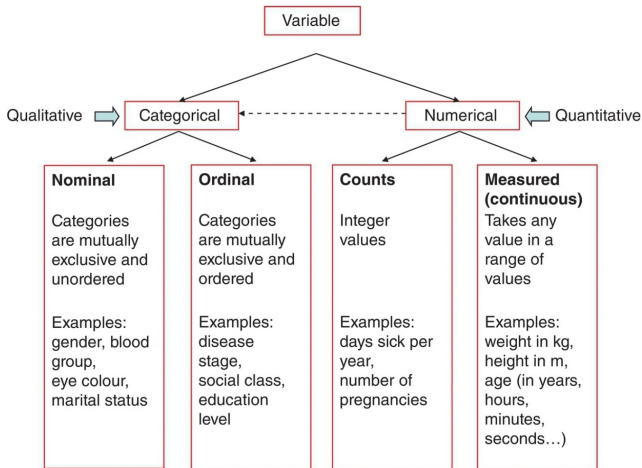
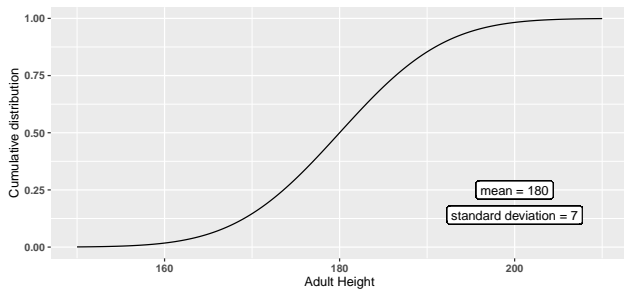
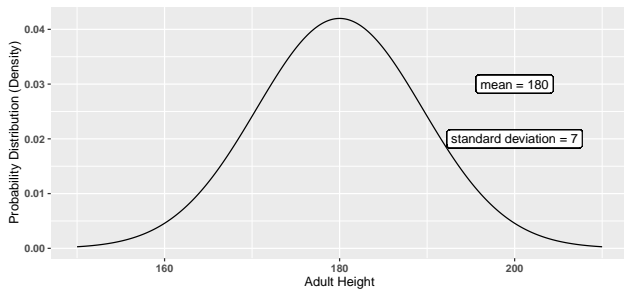


Figure 2.1 Broad classification of the different types of data with examples.

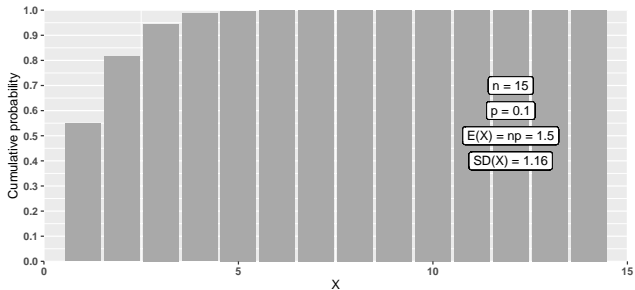
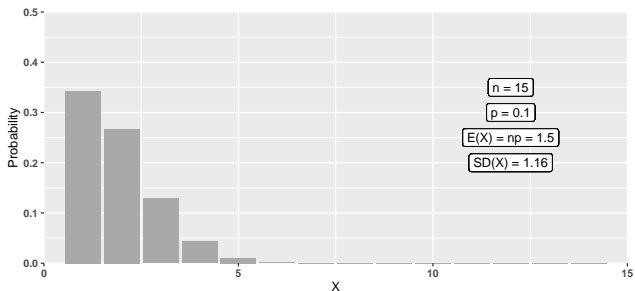
A Random variable X

- X is *discrete* or *continuous* (or both)
- Typical examples are Binomial and Poisson (discrete) and Normal (continuous)
- $P(X = 15)$ is the probability that X is equal to 15, say.
- This is OK for e.g. a Poisson distribution, but not useful for a continuous distribution ($P(X = 15) = 0$ in all continuous distributions!)
- For continuous distributions, intervals make more sense: $P(X > 15)$, $P(X \leq 15)$ etc.
- E.g. birth weight: $P(BW < 2500)$ is the risk of Low Birth Weight
- Conditional probability:
E.g. $P(BW < 2500 | GA = 258)$, i.e. the probability of being Low Birth Weight conditional on Gestational Age at birth being 258 days.

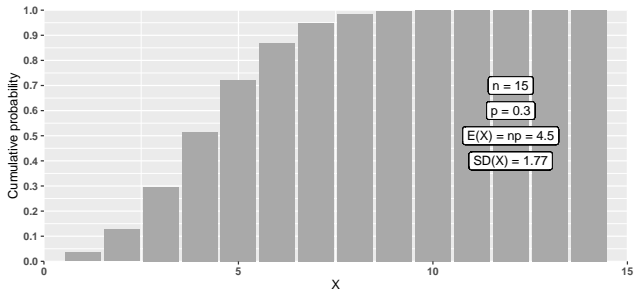
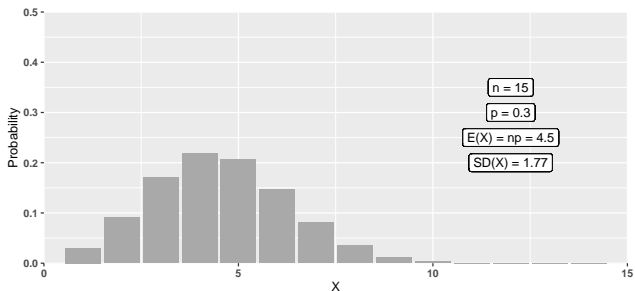
NORMAL DISTRIBUTION



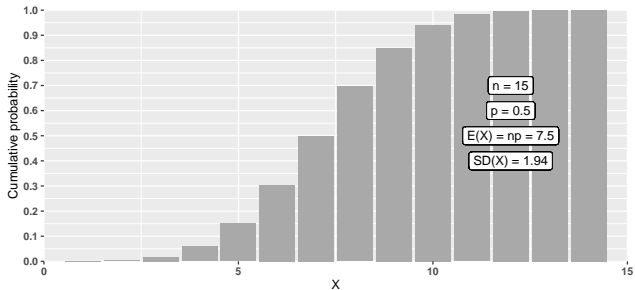
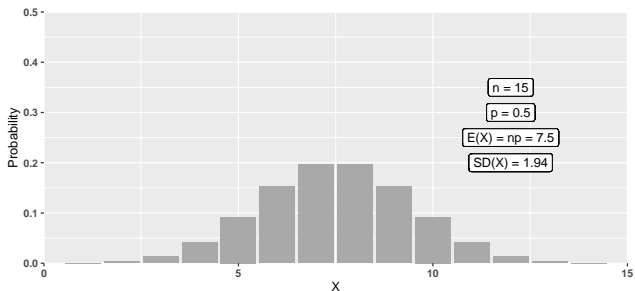
BINOMIAL DISTRIBUTION



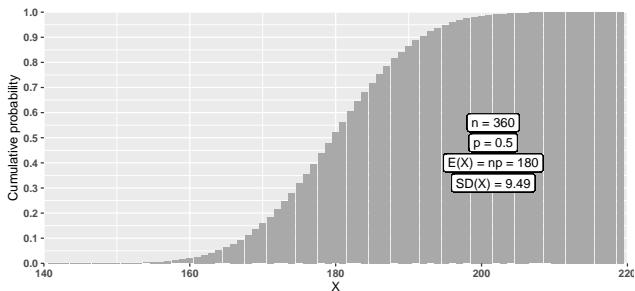
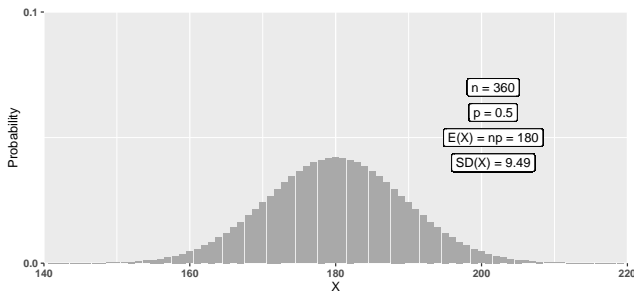
BINOMIAL DISTRIBUTION



BINOMIAL DISTRIBUTION



BINOMIAL DISTRIBUTION



EXPECTED VALUE, STANDARD DEVIATION

- Expectation EX is the *mean* of X

You often see

- μ (mju) in a population
- \bar{X} in a sample
- $\hat{\mu} = \bar{X}$ since $\hat{\mu}$ means *an estimate* of μ

$$\bar{X} = \frac{1}{n} \sum_{i=1}^n X_i = \frac{1}{n} (X_1 + X_2 + \dots + X_n)$$

- $SD(X)$ is the *Standard deviation* of X

You often see

- σ (sigma) in a population
- s or $SD(X)$ in a sample
- $\hat{\sigma} = s = SD(X)$ since $\hat{\sigma}$ means *an estimate* of σ

$$s^2 = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X})^2 = \frac{1}{n-1} \{ (X_1 - \bar{X})^2 + (X_2 - \bar{X})^2 + \dots + (X_n - \bar{X})^2 \}$$

STUDENT'S T-TESTS

Traditionally,

- One-sample t-test
 - Two independent samples t-test
 - Two paired samples t-test (take difference and use one-sample)

 - With small sample sizes, say, less than 30, a t-distribution (with degrees of freedom) is used instead of a Normal distribution
-

Today often replaced by regression models

- Larger sample sizes are the norm
- Software chooses distribution automatically
- More flexible in terms of adjustments etc.
- Paired samples t-test replaced by mixed-effects models

PARAMETRIC VERSUS NON-PARAMETRIC

THE GOOD, THE BAD, AND THE UGLY

The good(?), old days:

- Small sample size
- Test for Normal distribution
- If not very Normal, choose non-parametric (for instance the Wilcoxon-Mann-Whitney test)
- Else use t-test

Dilemma, model choice:

- Test for Normal distribution insensitive when n small
- When n is big, a normal distribution will often be rejected regardless
- ... BUT when n is big, the Central Limit Theorem helps you, so you usually don't need to assume a completely normal distribution anyway, and there is less need for the non-parametric

Dilemma, power:

- When n is small, parametric tests have little power...
- ... but non-parametric tests are even worse...
- And non-parametric tests usually only provide p-values

PARAMETRIC VERSUS NON-PARAMETRIC

THE GOOD, THE BAD, AND THE UGLY

Conclusion:

PARAMETRIC VERSUS NON-PARAMETRIC

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Conclusion:

Get over it. Get more data.

PARAMETRIC VERSUS NON-PARAMETRIC

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Conclusion:

Get over it. Get more data.

(Having said that, used in the right places, non-parametric and semi-parametric methods can actually be great! **Particularly when n is big!**)