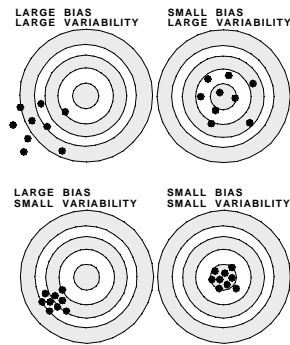


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Being approximately correct and being precisely wrong

PREFACE

Few general statistics textbooks¹ deal with the topic of measurement error. However, it is an important topic. Measurement error is present in all scientific work; while it can sometimes be lessened with careful planning and extra effort, it cannot be completely avoided. In some contexts it just adds noise and makes signals harder to detect. *But in others, its effects are both more subtle and more serious*: it systematically shifts the parameter estimates produced by models.

Its effects become more complicated and unpredictable in larger statistical models with two or more measured-with-error variables. We will restrict attention here to simple comparative situations involving one X and one Y. We will focus on the ‘*classical error model*’ where the errors in the measurements are uncorrelated with the true values being measured. We briefly mention the ‘*Berkson error model*,’ where the errors in the measurement are uncorrelated with the error-containing measurement – and thus correlated with the true value being measured.

More complicated cases (for example, human under-/over-reporting of values) require some knowledge of the subject matter area to know when such are likely to be present.

In the following home-grown material, JH will rely in part on notes he pre-

¹Examples of specialized textbooks focusing just on measurement error are *Measurement error in nonlinear models* by Carroll, Ruppert, and Stefanski 1998 and the 2006 *Measurement error in nonlinear models: a modern perspective*. by Carroll, the 1987 *Measurement error models* by Fuller, the 2004 *Measurement error and misclassification in statistics and epidemiology: impacts and Bayesian adjustments* by Gustafson. See also the excellent **Chapter 1 ‘Reliability of Measurement’** in the (older) book *The design and analysis of clinical experiments* by Fleiss. These, and others, are available from the McGill Library. A Google search will bring up more, as well as a large number of journal articles and other literature on this now-more-widely-studied topic.

pared when teaching measurement concepts/principles to graduate students in the physical and occupational therapy sciences. There, since the qualities/quantities being ‘measured’ were often **psychophysical rather than physical**, he drew on the very long history of ‘**psychometrics**’ in the psychology and education literature. This was a learning experience for him, since, up until then, he had mainly been concerned with the physical quantities measured in medical research.² He was greatly helped by reading parts of the textbook *Psychometric Theory* by Nunnally, and found that Nunnally’s examples of constructing (and measuring something by) exams were concrete and understandable illustrations of the concepts and principles. For those interested, his 1975 ‘lookback’ is a very nice introduction to the subject. There he tells us that

Since Spearman wrote on the topic in 1904, the theory of measurement reliability, or the converse of that, the theory of measurement error, has been a special interest of psychologists. I have always thought that this was partly out of necessity and partly because reliability theory is so neatly mathematized. In his 1904 article, Spearman developed most of the basic statistics pertaining to reliability that are still with us, including corrections for attenuation, the standard error of measurement, the correction of the split-half reliability coefficient for test length, and other statistics that are identified with test reliability. This deductive model, and the attendant mathematical developments, stood as a comprehensive theory of reliability for over 30 years.

QUANTIFYING THE QUALITY OF A MEASURING INSTRUMENT

Not surprisingly, much of our terminology comes from psychology, where measurements are seldom in centimetres, or kilograms. For the latter we have established standards established/regulated by various governmental agencies, such as those in the US, Canada, the EU and the UK

There are two aspects, **Reliability** (‘reproducibility’ / ‘precision’, the degree to which measurements of the same (unchanged) object would stay the same, and thus the relative positions of different objects would remain unchanged, if the measurement were repeated) and **Validity** (the extent to which the measurements measure what they are intended to measure).

But which to consider first?

When asked which they would study first, most students jump immediately to validity. Thus, in the following case, described in a Dr Fowler’s 1982 letter to the Editor of JAMA they want to know how the predictions turned out.

²JH was aware of measurement errors when as part of his PhD work, he made low-tech exhaled-breath measurements of the carbon monoxide in the blood of cigarette smokers, and used the amounts of various substances found in their cigarette butts to estimate their mouth-level exposure to nicotine. But he did nothing about it. Later, when he began working in clinical trials in oncology, he got a rude awakening. Fortunately, this time, after seeing just how much measurement error there was, and heeding the warning in the report, the Eastern Cooperative Oncology Group changed their cutoff for a partial response from a 25% to a 50% reduction in the cross-section area of the measured tumour.

The “Drano Test”: [Drano is a gel, sold by Johnson and Johnson, “formulated thick enough to easily pour through water straight to the clogged drain], dissolving it fast.”]

During the past several years, we have been asked frequently to do the “Drano test” to determine the sex of an unborn baby. It has been published in the lay press that this is a reliable means of sex determination. A Medline search failed to reveal anything in the medical literature concerning the Drano test. As a result, we performed the test in 100 consecutive pregnant women, checking monthly during the last trimester. The test was done by adding a small amount of crystal Drano to approximately 2 mL of urine, agitating, and interpreting results in one minute’s time. Reportedly, the color green indicates a male baby, and yellow to amber indicates a female.

But Dr Fowler knew that *consistency*³ i.e. **repeatability**, is a **prerequisite for accuracy (validity)**

Of the 100 patients, 21 failed to have the *same color* change *consistently*.

Of the babies born to these 21, eleven were girls and ten were boys. Of the remaining 79, we were right in sex determination of 37; of these, there were 20 girls and 17 boys. We were wrong in 42 predications; of these, there were 22 girls and 20 boys.

	P r e d i c t i o n			TOTAL
	Inconsistent	Girl	Boy	
GIRL	11	20	22	53
BOY	<u>10</u> 21	<u>20</u> 40	<u>17</u> 39	<u>47</u> 100

From this brief study, it would appear that the Drano test for antenatal sex determination is roughly equivalent to flipping a coin.

JH has also watched technicians test a photocopying machine. They first test was to see if the machine gave the same copy of a single object. The next was how faithful it was to the original.

⇒ *So we will start with reliability.*

³Not to be confused with its meaning in mathematical statistics, when at issue is the behaviour of statistical estimators.

Roadmap for the remainder of these Notes

The notes on pages 3-19 are from lectures JH gave in a course on measurement for the rehabilitation sciences.

- Pages 3-7: since few of these students had had experience with ANOVA’s (widely used to estimate reliability coefficients) the lectures began with a general introduction to these, showing **how the ANOVA table** – usually used for F *tests* – **can also be used to estimate variance components**. This use of ANOVA (and the Method of Moments) may be new to you too.
- Page 8 is a **nice introduction to reliability and validity**: the ideas are general, not confined to examinations, but to any measurement method/instrument.

Note the **two reliability measures**:

- a standard deviation (SD) measured in the **same scale** (scores, cm, Kg, ...) **that the instrument uses**, [smaller is better]
- a (more abstract) **fraction of overall variance**. Psychometricians prefer the latter because – unlike in physical measurement – the scales/ranges are arbitrary and the object being measured is not a phenomenon, but more a noumenon (abstract). [bigger is better]

These are the ‘converses’ that Spearman mentioned

- Pages 9-17 use ANOVA tables to estimate the variance components used as inputs to the reliability coefficients (**Intra-Class Correlations, p.10**).
- Pages 18-19 are ways to assess **validity**. [NB the different ways terms like accuracy, reliability, validity are used in statistics, and outside statistics].
- Pages 20-22: the **Effects of Measurement Error & role of the ICC**.
- Pages 22-23: **Berkson Error** [a subtle but important difference]

A passing comment on terminology: If JH had his way, we would be using the term reproducibility rather than reliability. One possible problem with the word ‘reliable’ is that it has many meanings in everyday English. What does one mean by a ‘reliable’ friend, or car, or clock, or watch? (and if you look up the history of the word, it was not without controversy grammar-wise). To JH at least, the word reproducible is less ambiguous. There is also the joke about a stopped clock: it is correct 2 times a day! Does that make its result reproducible? reliable? ? ?

RELIABILITY (Reproducibility, Precision): The extent to which one obtains the same (or very similar) answers/values/scores if an object/subject is measured repeatedly under similar situations. **Ways to quantify Reliability** include

- For **one** subject (s) or object:
 - average variation of individual measurements around their mean... either the square root of the average of squared deviations, i.e. standard deviation (SD); or the average absolute deviation, which will tend to be a bit smaller than the the SD.⁴ We could also use the range or other measures such as inter-quartile range (IQR).
 - SD of the measurements as a percentage of the mean of the measurements ... the [within-subject] Coefficient of Variation (CV).
- For **several** subjects (ss.) or objects:
 - average the CV's calculated for the different subjects; if CV's are highly variable, may want to give some sense of this using the range or other measure of spread of the CV's. In biochemical determinations, CV may differ at different concentrations.
 - The (within-ss) CV gives no sense of how well the measurements of different subjects (ss) segregate from each other. We might compare the SD of the within-ss measurements with the SD of the between-ss measurements, but, as we will see below, it makes more sense to focus on variances.
 - Can we use correlation (Pearson or Spearman) if we have 2 assessments of each ss.? What if we have a triplets of measurements for each subject? (cf. study of otitis media in twins and triplets)
 - Using correlation between scores on *random halves* of a test, can estimate how 'reproducible' the full test is (helpful if cannot repeat the test)
- **If the object is a population** (.e.g., the percentage of smokers, or showing immunity, among Canadian adults) and if it is measured (estimated) using a *statistic*: e.g., the proportion in a random sample of 1000 adults, it is possible from statistical laws concerning averages to quantify the reliability of the statistic without having to actually perform repeated measurements (samples). For simple random sampling, the formula

$$SE[\text{sample mean}] = \frac{SD[\text{individuals}]}{\text{sqrt of number of individuals measured}}$$

allows us to quantify the reliability indirectly. If we didn't know this formula, we could also arrive at an answer by various re-sampling methods applied to the individuals in the sample at hand – again without resorting to observing any additional individuals.

- **When the scale is arbitrary**, it is common to use a function of the (theoretical) Variance of the Within-ss measurements and the Variance of the Between-ss values. Classically, these **Components of Variance** were estimated using **Analysis of Variance (ANoVA)**. We will also use a Bayesian approach.

⁴Previous years' notes mistakenly said 'it will usually be quite close to the SD.'

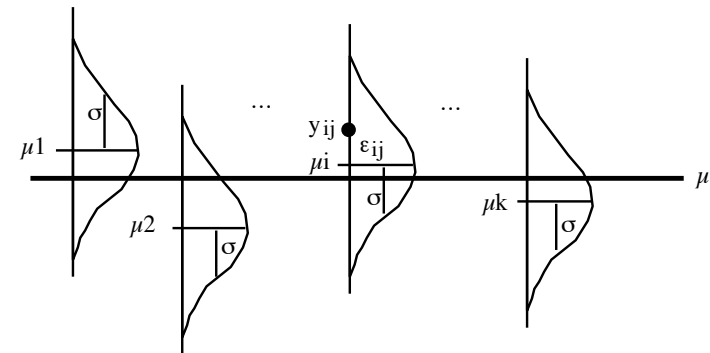
First, a General Orientation to Anova and its primary/classical use: testing differences between μ 's of k (≥ 2) different groups.⁵

E.g. 1-way ANOVA:

DATA:

	Group					
	1	2	...	i	...	k
Subject						
1	y ₁₁
2
...
j	.	.	.	y _{ij}	.	.
...
n	y _{kn}
Mean	\bar{y}_1	\bar{y}_2		\bar{y}_i		\bar{y}_k
Variance	s^2_1	s^2_2				s^2_k

MODEL



σ refers to the variation (SD) of all possible individuals in a group; It is an (unknowable) parameter; it can only be ESTIMATED.

Or, in symbols...

$$y_{ij} = \mu_i + e_{ij} = \mu + (\mu_i - \mu) + e_{ij}$$

⁵ This and the next several pages contain images of pages made with MS Word decades ago. JH hasn't had time to redo them in LaTeX. **2020 additions in red**

DE-COMPOSITION OF OBSERVED (EMPIRICAL) VARIATION

$$\sum\sum(y_{ij} - \bar{y})^2 = \sum\sum(y_i - \bar{y})^2 + \sum\sum(y_{ij} - \bar{y}_i)^2$$

TOTAL Sum of Squares = BETWEEN Groups + Sum of Squares + WITHIN Group Sum of Squares

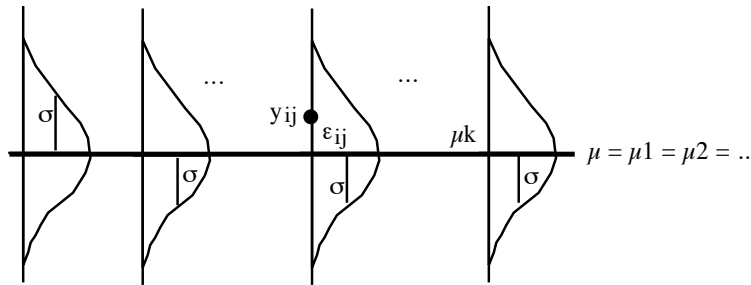
ANOVA TABLE

	Sum of Squares	Degrees of Freedom	Mean Square	F Ratio	P-Value
SOURCE	SS	df	MS (= SS /df)	$\frac{MS_{BETWEEN}}{MS_{WITHIN}}$	Prob(>F)
BETWEEN	xx.x	k-1	xx.x	x.xx	0.xx
WITHIN	xx.x	k(n-1)	xx.x		

LOGIC FOR F-TEST (Ratio of variances) as a test of

$H_0: \mu_1 = \mu_2 = \dots = \mu_i = \dots = \mu_k$

UNDER H0



Means, based on samples of n, should vary around μ with a variance of $\frac{\sigma^2}{n}$

Thus, if H_0 is true, and we calculate the empirical variance of the k different \bar{y}_i 's, it should give us an unbiased estimate of $\frac{\sigma^2}{n}$

i.e. $\frac{\sum[\bar{y}_i - \bar{y}]^2}{k-1}$ is an unbiased estimate of $\frac{\sigma^2}{n}$

i.e. $\frac{n \sum[\bar{y}_i - \bar{y}]^2}{k-1}$ is an unbiased estimate of σ^2

i.e. $\frac{\sum\sum[\bar{y}_i - \bar{y}]^2}{k-1} = MS_{BETWEEN}$ is an unbiased estimate of σ^2

Whether or not H_0 is true, the empirical variance of the n (within-group) values

y_{i1} to y_{in} i.e. $\frac{\sum[y_{ij} - \bar{y}_i]^2}{n-1}$ should give us an unbiased estimate of σ^2

i.e. $s^2_i = \frac{\sum[y_{ij} - \bar{y}_i]^2}{n-1}$ is an unbiased estimate of σ^2

so the average of the k different estimates,

$$\frac{1}{k} \sum s^2_i = \frac{1}{k} \sum \frac{\sum[y_{ij} - \bar{y}_i]^2}{n-1}$$

is also an unbiased estimate of σ^2

i.e. $\frac{\sum\sum[y_{ij} - \bar{y}_i]^2}{k[n-1]} = MS_{WITHIN}$ is an unbiased estimate of σ^2

THUS, under H_0 , both $MS_{BETWEEN}$ and MS_{WITHIN} are unbiased estimates of estimates of σ^2 and so their ratio should, apart from sampling variability, be 1. IF however, H_0 is not true, $MS_{BETWEEN}$ will tend to be larger than MS_{WITHIN} , since it contains an extra contribution that is proportional to how far the μ 's are from each other.

In this "non-null" case, the $MS_{BETWEEN}$ is an unbiased estimate of

$$\sigma^2 + \frac{\sum n[\mu_i - \bar{\mu}]^2}{k-1}$$

and so we expect that, apart from sampling variability, the ratio $\frac{MS_{BETWEEN}}{MS_{WITHIN}}$ should be greater than 1. The tabulated values of the F distribution (tabulated under the assumption that the numerator and denominator of the ratio are both estimates of the same quantity) can thus be used to assess how extreme the observed F ratio is and to assess the evidence against the H_0 that the μ 's are equal.

How ANOVA can be used to estimate Components of Variance used in quantifying Reliability.

The basic ANOVA calculations are the same, but the MODEL underlying them is different. First, in the more common use of ANOVA just described, the groups can be thought of as all the levels of the factor of interest. The number of levels is necessarily finite. The groups might be the two genders, all of the age groups, the 4 blood groups, etc. Moreover, when you publish the results, you explicitly identify the groups.

When we come to study subjects, and ask "How big is the intra-subject variation compared with the inter-subject variation, we will for budget reasons only study a sample of all the possible subjects of interest. We can still number them 1 to k, and we can make n measurements on each subject, so the basic layout of the data doesn't change. All we do is replace the word 'Group' by 'Subject' and speak of BETWEEN-SUBJECT and WITHIN-SUBJECT variation. So the data layout is...

DATA:

Measurement	Subject				
	1	2	i	...	k
1	y ₁₁
2
...
j	.	.	y _{ij}	.	.
...
n	y _{kn}
Mean	\bar{y}_1	\bar{y}_2	\bar{y}_i	...	\bar{y}_k
Variance	s ² ₁	s ² ₂	s ² _k

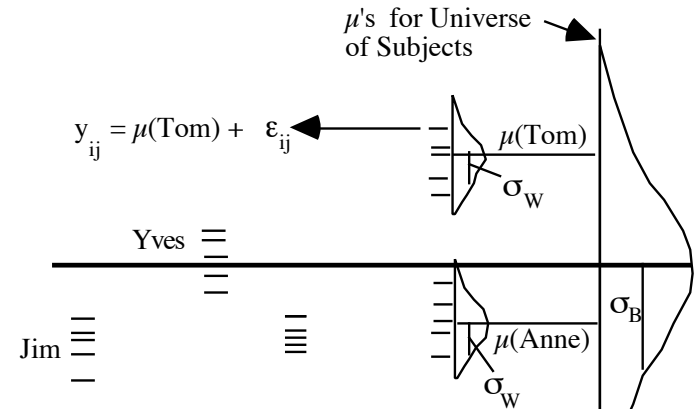
MODEL

The model is different. There is no interest in the specific subjects. Unlike the critical labels "male" and "female", or "smokers", "nonsmokers" and "exsmokers" to identify groups of interest, we certainly are not going to identify subjects as Yves, Claire, Jean, Anne, Tom, Jim, and Harry in the publication, and nobody would be fussed if in the dataset we used arbitrary subject identifiers to keep track of which measurements were made on whom. We wouldn't even care if the research assistant lost the identities of the subjects -- as long as we know that the correct measurements go with the correct subject!

The "Random Effects" Model uses 2 stages:

- (1) random sample of subjects, each with his/her own μ
- (2) For each subject, series of random variations around his/her μ

Notice the diagram has considerable 'segregation' of the measurements on different individuals. There is no point in TESTING for (inter-subject) differences in the μ 's. The task is rather to estimate the relative magnitudes of the two variance components σ^2_B and σ^2_W .



σ_B refers to the SD of the universe of μ 's ; It is an σ_B unknowable parameter and can only be ESTIMATED

σ_W refers to the variation (SD) of all possible measurements on a subject
It is an (unknowable) parameter; it can only be ESTIMATED.

Or, in symbols...

$$y_{ij} = \mu_i + \epsilon_{ij} = \mu + (\mu_i - \mu) + \epsilon_{ij} = \mu + \alpha_i + \epsilon_{ij}$$

$$\alpha_i \sim N(0, \sigma^2_B)$$

$$\epsilon_{ij} \sim N(0, \sigma^2_W)$$

DE-COMPOSITION OF OBSERVED (EMPIRICAL) VARIATION

$$\sum\sum(y_{ij} - \bar{y})^2 = \sum\sum(\bar{y}_i - \bar{y})^2 + \sum\sum(y_{ij} - \bar{y}_i)^2$$

TOTAL Sum of Squares = BETWEEN Subjects + Sum of Squares + WITHIN Subjects + Sum of Squares

ANOVA TABLE (Note absence of F and P-value Columns)

SOURCE	Sum of Squares	Degrees of Freedom	Mean Square	What the Mean Square is an estimate of*
	SS	df	MS (= SS /df)	
BETWEEN Subjects	xx.x	k-1	xx.x	$\sigma^2_W + n \sigma^2_B$
WITHIN Subjects	xx.x	k(n-1)	xx.x	σ^2_W

ACTUAL ESTIMATION OF 2 Variance Components

MS_{BETWEEN} is an unbiased estimate of $\sigma^2_W + n \sigma^2_B$

MS_{WITHIN} is an unbiased estimate of σ^2_W

By subtraction...

MS_{BETWEEN} - MS_{WITHIN} is an unbiased estimate of $n \sigma^2_B$

$\frac{\mathbf{MS_{BETWEEN} - MS_{WITHIN}}}{\mathbf{n}}$ is an unbiased estimate of σ^2_B

This is the **definitional** formula; the **computational** formula may be different.

* Pardon my ending with a preposition, but I find it difficult to say otherwise. These parameter combinations are also called the "Expected Mean Squares". They are the long-run expectations of the MS statistics As Winston Churchill would say, "For the sake of clarity, this one time this wording is something up which you would put".

Example....

DATA:

Measurement	Subject						
	Tom	Anne	Yves	Jean	Claire	Emma	
1	4.8	5.5	5.1	6.4	5.8	4.5	
2	4.7	5.2	4.9	6.2	6.3	4.1	
3	4.9	5.2	5.3	6.6	5.6	4.0	
Mean	4.8	5.3	5.1	6.4	5.9	4.2	Variance = 0.614
Variance	0.01	0.03	0.04	0.04	0.13	0.07	

ANOVA TABLE (Check... I did it by hand!)

SOURCE	Sum of Squares	Degrees of Freedom	Mean Square	What the Mean Square is an estimate of... *
	SS	df	MS (= SS /df)	
BETWEEN Subjects	9.205	5	1.841	$\sigma^2_W + n \sigma^2_B$
WITHIN Subjects	0.640	12	0.053	σ^2_W
TOTAL	9.845	17		

ESTIMATES OF VARIANCE COMPONENTS

MS_{WITHIN} = **0.053** is an unbiased estimate of σ^2_W

$\frac{\mathbf{1.841 - 0.053}}{\mathbf{3}}$ = **0.596** is an unbiased estimate of σ^2_B

1-Way ANOVA Calculations performed by SAS; Components estimated manually

```
PROC GLM in SAS ==> estimating components 'by hand'
DATA a; INPUT Subject Value; LINES;
1 4.8
1 4.7
...
6 4.5
proc glm; class subject; model value=subject / ss3;
random subject ;
```

See worked example using earsize data.

If unequal numbers of measurements per subject, see formula in A&B or Fleiss

This is an example of the METHOD of MOMENTS.

Estimating Components of Variance using "Black Box"

PROC VARCOMP; class subject ; model Value = Subject ;
 See worked example following...

2 measurements (in mm) of earsize of 8 subjects by each of 4 observers

subject	1				2				3				4			
obsr	1	2	3	4	1	2	3	4	1	2	3	4	1	2	3	4
1st	67	65	65	64	74	74	74	72	67	68	66	65	65	65	65	65
2nd	67	66	66	66	74	73	71	73	68	67	68	67	64	65	65	64

subject	5				6				7				6			
obsr	1	2	3	4	1	2	3	4	1	2	3	4	1	2	3	4
1st	65	62	62	61	59	56	55	53	60	62	60	59	66	65	65	63
2nd	61	62	60	61	57	57	57	53	60	65	60	58	66	65	65	65

INTRA-OBSERVER VARIATION (e.g. observer #1)

e.g. observer #1

PROC GLM in SAS ==> estimating components 'by hand'

INPUT subject rater occasion earsize; if observer=1;
 The data set has 16 obsns & 4 variables.

proc glm; class subject; model earsize=subject / ss3;
 random subject ;

General Linear Models Procedure: Class Level Information

Class	Levels	Values
SUBJECT	8	1 2 3 4 5 6 7 8 ; # of obsns. in data set = 16

Dependent Variable: EARSIZE

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	7	341.00	48.71	35.43	0.0001
Error	8	11.00	1.38		
Corrected Total	15	352.00			

R-Square	C.V.	Root MSE	EARSIZE Mean
0.968750	1.80	1.17260	65.0

Source	DF	Type III SS	Mean Square	F Value	Pr > F
SUBJECT	7	341.00	48.71	35.43	0.0001

Source	Type III Expected Mean Square
SUBJECT	Var(Error) + 2 Var(SUBJECT)

$$\begin{aligned} \text{Var(Error)} + 2 \text{Var(SUBJECT)} &= 48.71 \\ \text{Var(Error)} &= 1.38 \\ 2 \text{Var(SUBJECT)} &= 47.33 \\ \text{Var(SUBJECT)} &= 47.33 / 2 = 23.67 \end{aligned}$$

The measurements were made by 4 physical/occupational therapy graduate students (obsr 1-4), on the remaining $n = 8$ subjects – 6 fellow students and 2 instructors, JH and SWD. After a short meeting to standardize themselves, and using clear plastic rulers, each of the 4 measured each of the 8 twice (1st and 2nd). Note: the final subject should be numbered 8, not 6.

Estimating Variance components using PROC VARCOMP in SAS

proc varcomp; class subject ; model earsize = subject ;

Variance Components Estimation Procedure: Class Level Information

Class Levels Values

SUBJECT 8 1 2 3 4 5 6 7 8 ; # obsns in data set = 16

MIVQUE(0) Variance Component Estimation Procedure

Variance Component	Estimate
EARSIZE	
Var(SUBJECT)	23.67
Var(Error)	1.38

• ICC (Fleiss § 1.3)

$$\text{ICC} = \frac{\text{Var(SUBJECT)}}{\text{Var(SUBJECT)} + \text{Var(Error)}} = \frac{23.67}{23.67 + 1.38} = 0.94$$

1-sided 95% Confidence Interval (see Fleiss p 12)

df for F in CI: (8-1)= 7 and 8

so from Tables of F distribution with 7 & 8 df, $F = 3.5$

So lower limit of CI for ICC is

$$\begin{aligned} & 35.43 - 3.5 \\ & = \frac{35.43 - 3.5}{35.43 + (2 - 1) \cdot 3.5} = 0.82 \end{aligned}$$

EXERCISE: Carry out the estimation procedure for one of the other 3 observers.

INTERPRETING YOUR GRE SCORES

(Blurb from Educational Testing Service)

Your test score is an estimate, not a complete and perfect measure, of your knowledge and ability in the area tested. In fact, if you had taken a different edition of the test that contained different questions but covered the same content, it is likely that your score would have been slightly different. The only way to obtain perfect assessment of your knowledge and ability in the area tested would be for you to take all possible test editions that could ever be constructed. Then assuming that your ability and knowledge did not change, the average score on all those editions, referred to as your "true score," would be a perfect measure of your knowledge and ability in the content areas covered by the test. Therefore, scores are estimates and not perfect measures of a person's knowledge and ability. Statistical indices that address the imprecision of scores in terms of standard error of measurement and reliability are discussed in the next two sections.

STANDARD ERROR OF MEASUREMENT

The difference between a person's true and obtained scores is referred to as "error of measurement."* The error of measurement for an individual person cannot be known because a person's true score can never be known. The average size of these errors, however, can be estimated for a group of examinees by the statistic called the "standard error of measurement for individual scores." The standard error of measurement for individual scores is expressed in score points. About 95 percent of examinees will have test scores that fall within two standard errors of measurement of their true scores. For example, the standard error of measurement of the GRE Psychology Test is about 23 points. Therefore, about 95 percent of examinees obtain scores in Psychology that are within 46 points of their true scores. About 5 percent of examinees, however, obtain scores that are more than 46 points higher or lower than their true scores.

Errors of measurement also affect any comparison of the scores of two examinees. Small differences in scores may be due to measurement error and not to true differences in the abilities of the examinees. The statistic "standard error of measurement of score differences" incorporates the error of measurement in each examinee's score being compared. This statistic is about 1.4 times as large as the standard error of measurement for the individual scores themselves. Approximately 95 percent of the differences between the obtained scores of examinees who have the same true score will be less than two times the standard error of measurement of score differences. Fine distinctions should not be made when comparing the scores of two or more examinees.

This excellent blurb is from the days when Graduate Records Examinations (GRE's) were taken on paper and took 3 hours or more. Today, with the 'adaptive testing' that is possible with online tests, the exams can be personalized and speeded up, by starting with questions that are used to get a general sense of one's ability/knowledge, and then using questions of known difficulty to move up or down. So the sequence and number of questions one candidate gets differ from those another candidate gets. This leads into a later topic: the method of *split halves* used to measure reliability.

RELIABILITY

The reliability of a test is an estimate of the degree to which the relative position of examinees' scores would change if the test had been administered under somewhat different conditions (for example, examinees were tested with a different test edition).

Reliability is represented by a statistical coefficient that is affected by errors of measurement. Generally, the smaller the errors of measurement in a test, the higher the reliability. Reliability coefficients may range from 0 to 1, with 1 indicating a perfectly reliable test (i.e., no measurement error) and zero reliability indicating a test that yields completely inconsistent scores. Statistical methods are used to estimate the reliability of the test from the data provided by a single test administration. Average reliabilities of the three scores on the General Test and of the total scores on the Subject Tests range from .88 to .96 on recent editions. Average reliabilities of subscores on recent editions of the Subject Test range from .82 to .90.

Data regarding standard errors of measurement and reliability of individual GRE tests may be found in the leaflet *Interpreting Your GRE General and Subject Test Scores*, which will be sent to you with your GRE Report of Scores.

VALIDITY

The validity of a test—the extent to which it measures what it is intended to measure—can be assessed in several ways. One way of addressing validity is to delineate the relevant skills and areas of knowledge for a test, and then, when building each edition of the test, make sure items are included for each area. This is usually referred to as content validity. A committee of ETS specialists defines the content of the General Test, which measures the content skills needed for graduate study. For Subject Tests, ETS specialists work with professors in that subject to define test content. In the assessment of content validity, content representativeness studies are performed to ensure that relevant content is covered by items in the test edition.

Another way to evaluate the validity of a test is to assess how well test scores forecast some criterion, such as success in grade school. This is referred to as predictive validity. Indicators of success in graduate school may include measures such as graduate school grades, attainment of a graduate degree, faculty ratings, and departmental examinations. The most commonly used measure of success in assessing the predictive validity of the GRE tests is graduate first year grade point average. Reports on content representativeness and predictive validity studies of GRE tests may be obtained through the GRE Program office.

* The term "error of measurement" does not mean that someone has made a mistake in constructing or scoring the test. It means only that a test is an imperfect measure of the ability or knowledge being tested.

This last comment about the meaning of the word 'error' also applies in statistics more generally. When communicating with non-statisticians, we should carefully explain our terminology.

Outline

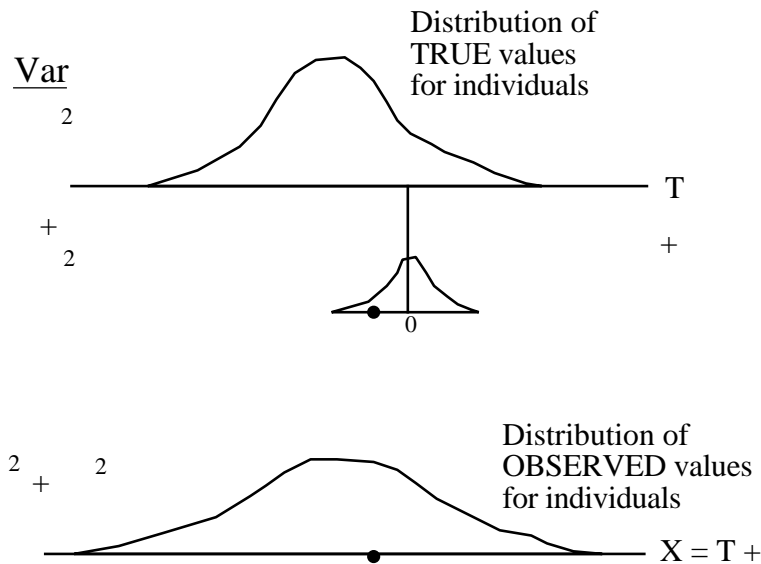
Some ways to quantify Reliability:

- Reliability Coefficient
- Internal Consistency (Cronbach's α)

Implications:

.....

Model for Reliability



"True" scores / values not knowable;

$\epsilon \perp T$
 Variance calculation assumes that the distribution of errors is independent of T

Reliability Coefficient

$$r_{xx} = \frac{\sigma_T^2}{\sigma_T^2 + \sigma_{\epsilon}^2}$$

i.e. the fraction of observed variation that is 'real'

Note that one can 'manipulate' r by choosing a large or small σ_{ϵ}^2

Effect of # of Items on Reliability Coefficient

(if all items have same variance and same intercorrelations)

SCALE 2 N Times more items than SCALE 1

$$r_{SCALE 2} = \frac{N \times r_{SCALE 1}}{1 + [N-1] \times r_{SCALE 1}}$$

e.g.

Scale	# Items	r
1	10	0.4
2	20 (× 2)	0.57
3	30 (× 3)	0.67

The above 'stepped-up reliability' formula is often called the Spearman-Brown prediction formula. See Brown 1910: correlation of mental abilities. See Wainer et al. on why unnecessarily long tests may be 'impeding the progress of Western civilisation.'

Cronbach's α

k items

$$\alpha = \frac{k \times \bar{r}}{1 + [k-1] \times \bar{r}}, \text{ where } \bar{r} = \text{average of inter-item correlations}$$

α is an estimate of the expected correlation of one test with an alternative form with the same number of items.

α is a lower bound for r_{XX} i.e. $r_{XX} \geq \alpha$

$r_{XX} = \alpha$ if items are parallel.

parallel

Average [item 1] = Average [item 2] = Average [item 3] = ...

Variance[item 1] = Variance[item 2] = Variance [item 3] = ...

Correlation[item 1, item 2] = Correlation[item 1, item 3] = ...
=Correlation[item 2, item 3] = ...

INTRAClass CORRELATIONS (ICC's)

- **Various versions**

TEST-RETEST

INTRA-RATER

INTER-RATER...

- **Formed as Ratios of various Variances**

e.g.
$$\frac{\sigma_{\text{TRUE}}^2}{\sigma_{\text{TRUE}}^2 + \sigma_{\text{ERROR}}^2}$$

with estimates of various σ^2 's substituted for the σ^2 's .

Estimates of various components typically derived from ANOVA.

- **Note the distinction between DEFINITIONAL FORM (involving PARAMETERS) and COMPUTATIONAL FORM (involving STATISTICS)**

Fleiss Chapter 1 good here; Norman & Streiner not so good!!)

To prevent cheating, in some multiple choice exams students get versions with the order of questions permuted – or even different questions.

How reproducible are students' scores? Do students rank the same if we use the subtotals for the even- and odd-numbered questions?

We could correlate a random-half versus the reminder. We could average the correlation over many such random splits.

Cronbach's alpha is seen as a way of not having to run a second exam, but to work out the reliability '*internally*' from the one exam

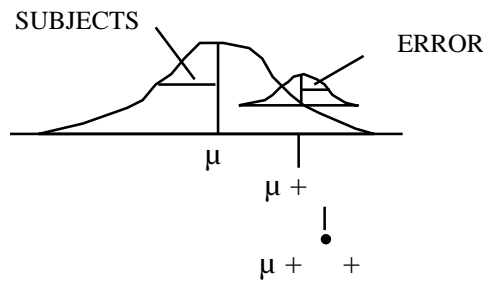
The ETS has a large enough question bank to ensure a high correlation between scores on different versions

See our recent use of 'split halves' in another context.

ICC's (Portnoy and Wilkins)

- (1) multiple (unlabeled) measurements of each subject
- (2) same set of raters measure each subject; raters thought of as a random sample of all possible raters.
- (3) as in (2), but these raters studied are the only raters of interest

.....
 (1) multiple (unlabeled) measurements of each subject



$$ICC = \frac{2 \text{ SUBJECTS}^2}{2 \text{ SUBJECTS}^2 + 2 \text{ ERROR}^2}$$

Model for observed data:

$$y[\text{subject } i, \text{ measurement } j] = \mu + \mu_i + e_{ij}$$

EXAMPLE 1

This example is in the spirit of the way the ICC was first used, as a measure of the greater similarity within families than between families: Study by Bouchard (NEJM) on weight gains of 2 members from each of 12 families: It is thought that there will be more variation between members of different families than between members of the same family: family (genes) is thought to be a large source of variation; the two twins per family are thought of as 'replicates' from the family and closer to each other (than to others) in their responses. Here the "between" factor is family i.e. families are the subjects and the two twins in the family are just replicates and they don't need to be labeled (if we did label them 1 and 2, the labels would be arbitrary, since the two twins are thought to be 'interchangeable'. (weight gain in Kg over a summer)

model: weight gain for person j in family i = $\mu + \mu_i + e_{ij}$

... backstory of example 1 here ; data & R Code* here .

1-way Anova and Expected Mean Square (EMS)

Source	Sum of Sq	d.f	Mean Square	Expected Mean Square
Between (families)	99	11	9.0	$2\sigma_{\text{error}}^2 + k_0 \sigma_{\text{between}}^2$
Error(Within families)	30	12	2.5	$2\sigma_{\text{error}}^2$

Total	129	23		

In our example, we measure k=2 members from each family, so k_0 is simply 2

[if the k's are unequal, k_0 is somewhat less than the average k... $k_0 = \text{average } k - (\text{variance of } k\text{'s}) / (n \text{ times average } k)$...see Fleiss page 10]

Estimation of parameters that go to make up ICC

2.5 is an estimate of $2\sigma_{\text{error}}^2$

9.0 is an estimate of $2\sigma_{\text{error}}^2 + 2 \sigma_{\text{between}}^2$

 6.5 is an estimate of $2 \sigma_{\text{between}}^2$

$\frac{6.5}{2}$ is an estimate of $\sigma_{\text{between}}^2$

$$\frac{\frac{6.5}{2}}{\frac{6.5}{2} + 2.5} = \frac{3.25}{3.25 + 2.5} = 0.57$$

is an estimate of ICC = $\frac{\sigma_{\text{between}}^2}{\sigma_{\text{between}}^2 + \sigma_{\text{error}}^2}$

*R code uses MCMC to obtain a credible interval for ICC

Origin of w'in- ('intra-') vs. b'ween- family ('class') terminology

COMPUTATIONAL Formula for "1-way" ICC

$$\frac{\frac{MS_{\text{between}} - MS_{\text{within}}}{k_0}}{\frac{MS_{\text{between}} - MS_{\text{within}}}{k_0} + MS_{\text{within}}}$$

$$= \frac{MS_{\text{between}} - MS_{\text{within}}}{MS_{\text{between}} + (k_0 - 1)MS_{\text{within}}} \quad [\text{shortcut}]$$

is an estimate of the ICC

Notes:

- Streiner and Norman start on page 109 with the 2-way anova for inter-observer variation. There are mistakes in their depiction of the SSerror on p 110 [it should be $(6-6)^2 + (4-4)^2 + (2-1)^2 + \dots + (8-)^2 = 10$. If one were to do the calculations by hand, one usually calculates the SStotal and then obtains the SSerror by subtraction]
- They then mention the 1-way case, which we have discussed above, as "the observer nested within subject" on page 112
- Fleiss gives methods for calculating CI's for ICC's.

EXAMPLE 2: INTRA-OBSERVER VARIATION FOR 1 OBSERVER**Computations performed on earlier handout...**

$$\text{Var}(\text{SUBJECT}) = 23.67 \quad \text{Var}(\text{ERROR}) = 1.38$$

$$\hat{\text{ICC}} = 23.67 / (23.67 + 1.38) = 0.94$$

An estimated 94% of observed variation in earsize measurements by this observer is 'real' .. i.e. reflects true between-subject variability.

Note that I say 'an estimated 94% ...'. I do this because the 94% is a *statistic* that is subject to sampling variability (94% is just a point estimate or a 0% Confidence Interval). An interval estimate is given by say a 95% confidence interval for the true ICC (lower bound of a 1-sided CI is 82% ... see previous handout)

Computational versions, used to save steps when computing statistics by hand, hide the concept (definition) behind these statistics.

The MCMC method yields a more realistic credible interval than Fleiss' CI. More examples [here](#).

Increasing Reliability by averaging several measurements

In 1-way model: $y_{i,j} = \mu + \alpha_i + e_{ij}$

where $\text{var}[\alpha_i] = \sigma^2_{\text{between subjects}}$; $\text{var}[e_{ij}] = \sigma^2_{\text{"error"}}$

Then if we average k measurements, i.e.,

$$\bar{y}_{i} = \mu + \alpha_i + \bar{e}_{i}$$

then

$$\text{Var}_i[\bar{y}_{i}] = \sigma^2_{\text{between}} + \frac{\sigma^2_{\text{"error"}}}{k}$$

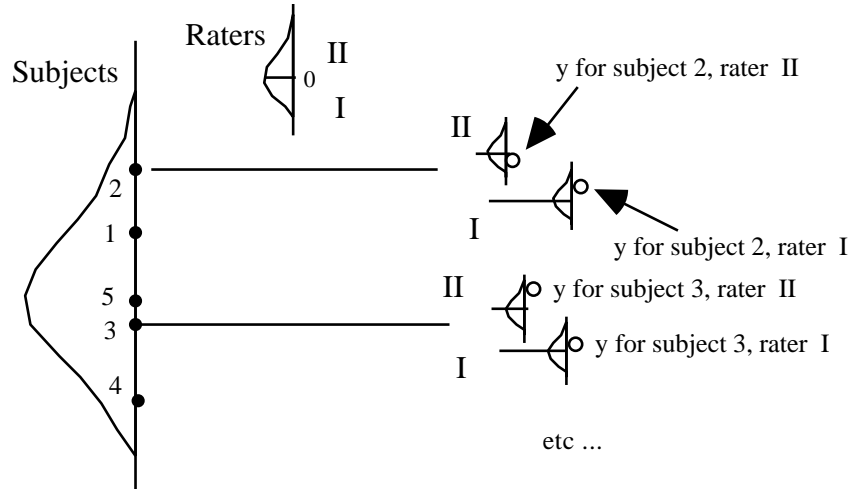
$$\text{So ICC}[k] = \frac{\sigma^2_{\text{between}}}{\sigma^2_{\text{between}} + \frac{\sigma^2_{\text{"error"}}}{k}}$$

This is called "**Stepped-Up**" Reliability.

ICC's (Portnoy and Wilkins).

(2) same set of raters measure each subject; raters thought of as a random sample of all possible raters.

Model



$$\mu + [subject] + [rater] + error$$

2 subjects 2 raters 2 error

From 2- way data layout (subjects x Raters)

estimate 2_{subjects} , 2_{raters} and 2_{error} by 2-way ANOVA

Substitute variance estimates in appropriate ICC form

e.g. 2 measurements (in mm) of earsize of 8 subjects by each of 4 observers

subject	1				2				3				4			
obsr	1	2	3	4	1	2	3	4	1	2	3	4	1	2	3	4
1st	67	65	65	64	74	74	74	72	67	68	66	65	65	65	65	65
2nd	67	66	66	66	74	73	71	73	68	67	68	67	64	65	65	64
subject	5				6				7				8			
obsr	1	2	3	4	1	2	3	4	1	2	3	4	1	2	3	4
1st	65	62	62	61	59	56	55	53	60	62	60	59	66	65	65	63
2nd	61	62	60	61	57	57	57	53	60	65	60	58	66	65	65	65

ESTIMATING INTER-OBSERVER VARIATION from occasion=1;

```
PROC GLM in SAS ==> estimating components 'by hand'
INPUT subject rater occasion earsize; if occasion=1; (32 obsns)
```

```
proc glm; class subject rater; model earsize=subject rater / ss3;
random subject rater;
```

General Linear Models Procedure: Class Level Information

Class	Levels	Values
SUBJECT	8	1 2 3 4 5 6 7 8
RATER	4	1 2 3 4

Number of observations in data set = 32

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	10	764.500	76.45	78.80	0.0001
Error	21	20.375	0.97		
Corrected Total	31	784.875			

R-Square	C.V.	Root MSE	EARSIZE Mean
0.974040	1.534577	0.98501	64.1875

Source	DF	Type III SS	Mean Square	F Value	Pr > F
SUBJECT	7	734.875000	104.98	108.20	0.0001
RATER	3	29.625000	9.87	10.18	0.0002

Source Type III Expected Mean Square

SUBJECT Var(Error) + 4 Var(SUBJECT)

RATER Var(Error) + 8 Var(RATER)

So... solving 'by hand' for the 3 components...

$$\text{Var(Error)} + 4 \text{Var(SUBJECT)} = 104.98$$

$$\text{Var(Error)} = 0.97$$

$$\implies 4 \text{Var(SUBJECT)} = 104.01$$

$$\implies \text{Var(SUBJECT)} = 104.01 / 4 = 26.00$$

$$\text{Var(Error)} + 8 \text{Var(RATER)} = 9.87$$

$$\text{Var(Error)} = 0.97$$

$$\implies 8 \text{Var(RATER)} = 8.90$$

$$\implies \text{Var(RATER)} = 8.90 / 8 = 1.11$$

$$\text{Var(Error)} = 0.97$$

Estimating Variance components using PROC VARCOMP in SAS

```
proc varcomp; class subject rater; model earsize = subject rater;
```

Variance Component	Estimate EARSIZE
Var(SUBJECT)	26.00
Var(RATER)	1.11
Var(Error)	0.97

• **ICC: "Raters Random"** (Fleiss § 1.5.2)

$$\text{ICC} = \frac{\text{Var}(\text{SUBJECT})}{\text{Var}(\text{SUBJECT}) + \text{Var}(\text{RATER}) + \text{Var}(\text{Error})} = \frac{26.00}{26.00 + 1.11 + 0.97} = 0.93$$

1-sided 95% Confidence Interval (see Fleiss p 27)

df for F in CI: (8-1)= 7 and v* , where

$$v^* = \frac{(8-1)(4-1)(4 \cdot 0.93 \cdot 10.18 + 8[1+(4-1) \cdot 0.93] - 4 \cdot 0.93)^2}{(8-1) \cdot 4^2 \cdot 0.93^2 \cdot 10.18^2 + (8[1+(4-1) \cdot 0.93] - 4 \cdot 0.93)^2} = 8.12$$

so from Tables of F distribution with 7 & 8 df, F = 3.5

So lower limit of CI for ICC is

$$= \frac{8(104.98 - 3.5 \cdot 0.97)}{8 \cdot 104.98 + 3.5 \cdot [4 \cdot 9.87 + (8 \cdot 4 - 8 - 4) \cdot 0.97]} = 0.78$$

• **ICC: if use one "fixed" observer** (see Fleiss p 23, strategy 3)

$$\text{ICC} = \frac{\text{Var}(\text{SUBJECT})}{\text{Var}(\text{SUBJECT}) + \text{Var}(\text{Error})} = \frac{26.00}{26.00 + 0.97} = 0.96$$

lower limit of 95% 1-sided CI (eqn 1.49: F = 2.5 ; 7 & 7x3=21 df)

$$\text{ICC} = \frac{104.98 - 2.5}{104.98 + (4-1) \cdot 2.5} = 0.91$$

USING ALL THE DATA SIMULTANEOUSLY

(can now estimate subject x Rater interaction .. i.e extent to which raters 'reverse themselves' with different subjects)

Components of variance when use **both** measurements (all 64 obsns)

```
proc varcomp;                                proc varcomp;
  class subject rater;                        class subject rater;
  model earsize = subject rater;              model earsize = subject rater
                                              subject*rater;
```

Variance Component	Estimate EARSIZE	Variance Component	EARSIZE
Var(SUBJECT)	25.52	Var(SUBJECT)	25.47
Var(RATER)	0.70	Var(RATER)	0.67
Var(Error)	1.37	Var(SUBJECT*RATER)	0.31
		Var(Error)	1.13

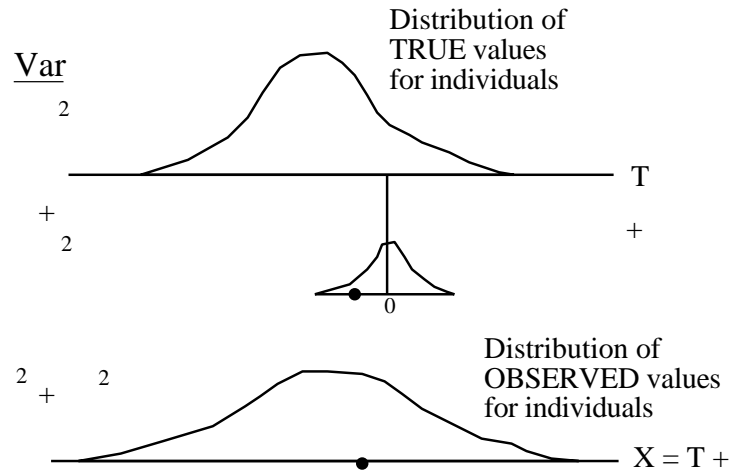
LINK between STANDARD ERROR OF MEASUREMENT and RELIABILITY COEFFICIENT

Example: GRE Tests (cf blurb from Educational Testing Service)

Standard Error of Measurement = 23 points

Reliability Coefficient: R = 0.93

recall...



$$\sigma_e^2 = 23 \implies \sigma_e = 529;$$

$$R = \frac{\sigma_T^2}{\sigma_T^2 + \sigma_e^2} = 0.93 \implies \sigma_T^2 = \frac{R \times \sigma_e^2}{1 - R} = \frac{0.93 \times 529}{1 - 0.93} = 7028$$

$$\sigma_T^2 + \sigma_e^2 = 7028 + 529 = 7557 \implies \sqrt{\sigma_T^2 + \sigma_e^2} = \sqrt{7557} = 87$$

So if 3 SD's on either side of the mean of 500 covers most of the observed scores, this would give a range of observed scores of $500 - 261 = 239$ to $500 + 261 = 761$.

Another way to say it (see Streiner and Norman, bottom of page 119) :-

$$\sigma_e = \sqrt{\sigma_T^2 + \sigma_e^2} \times \sqrt{1 - R} = \text{SD}[\text{observed scores}] \times \sqrt{1 - R}$$

See section 12.7.2 (Measurement error, [here](#) for a colour-based depiction of the mixing of 'true' and 'error' distributions.

Confidence Intervals / Sample Sizes for ICC's

see Fleiss...

CI's based on F distribution tables;

CI's not symmetric;

More interested in 1-sided CI's i.e. (lower bound, 1) i.e. ICC 0.xx;

See also Donner and Eliasziw.

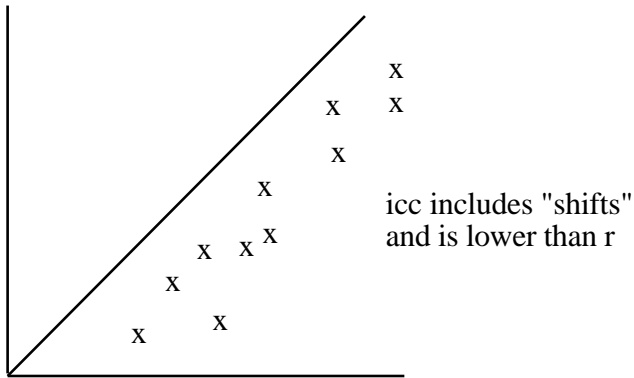
NOTE: If interested in ICC that incorporates random raters, then sample size must involve both # of raters and # of raters

CI will be very wide if use only 2 or 3 raters

Approach sample size as "n's or raters and subjects needed for a sufficiently narrow CI.

Why Pearson's r is not always a good [or practical] measure of reproducibility

1. It does not pick up "shifts"

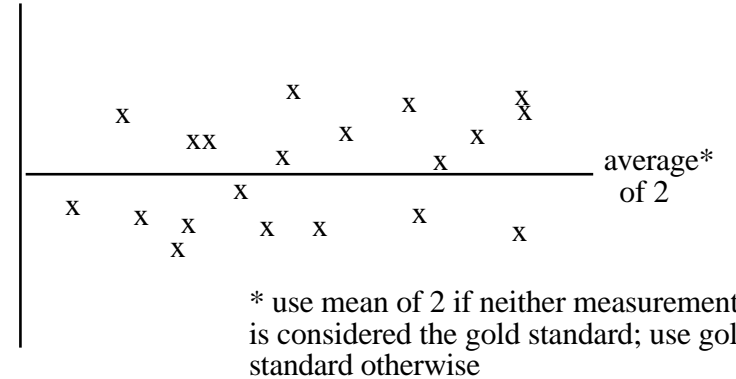


2. not practical if > 2 measurements or variable # of measurements per subject

ICC 'made for' such situations

Method of Bland & Altman [Lancet]

Difference of 2 measurements



+++ see biases quickly

can explain to your in-laws (can you explain ICC to them?)

emphasises errors in measurements scale itself (like ±23 in GRE score)

--- if don't know real range, magnitudes of standard error of measurement not helpful (see Norman & Streiner)

cannot use with > 2 measurements

doesn't generalize to raters

Assessing reproducibility of measurements made on a CATEGORICAL scale

Categorizations of n subjects by RATER 2

		C1	C2	C3
Categorizations of subjects by RATER 1	C1			
	C2			
	C3			
				n

See chapter 13 in Fleiss's book on Rates and Proportions
or pp 516-523 of Chapter 26 of Portnoy and Wilkins

- Simple Measure

$$\% \text{ agreement} = \frac{\# \text{ in diagonal cells}}{n} \times 100$$

- Chance-Corrected Measure

$$= \frac{\% \text{ agreement} - \% \text{ agreement expected by chance}^*}{100\% \text{ agreement} - \% \text{ agreement expected by chance}}$$

* expected proportion = $p[\text{row}] * p[\text{col}]$ --- over the diagonals

(see Aickin's arguments against 'logic' of chance-correction:
Biometrics 199)

can give weights for 'partial' agreement

if > 2 raters, use range or average of pairwise kappas

with quadratic weights, weighted kappa = icc

Validity Statistics

<u>INSTRUMENT</u>	<u>CRITERION</u>	<u>STATISTIC(S)</u>
numerical	numerical - same scale	<ul style="list-style-type: none"> • calculate discrepancies (Altman & Bland) • describe distribution of discrepancies
numerical	numerical - different scale	<ul style="list-style-type: none"> • correlation [Pearson or Spearman]
numerical	ordered categories (e.g. known groups)	<ul style="list-style-type: none"> • correlation [ranks] • ANOVA [groups]
ordered categories	ordered categories	<ul style="list-style-type: none"> • Kendall's "tau"
numerical or ordered categories	binary	<ul style="list-style-type: none"> • difference in means [parametric/ nonparametric test] • ROC curve
binary	binary	<ul style="list-style-type: none"> • sensitivity (Probability + on instr. if Criterion +) & specificity (Probability – on instr. if Criterion –) • predictive values • (phi) statistic (see Streiner & Norman)

For many reasons, this treatment of Validity is short: if the quality is physical, we usually have a gold standard, so assessment is direct (example: Tracking Physical Activity Data). If it is psycho-physical or conceptual the challenge is less in the statistical methods, and more in finding indirect ways to assess it: see the paragraph on VALIDITY in the blurb about the GRE, or the q. on readability in Exercises 1 and 2 below.

CONSEQUENCES

of

MEASUREMENT ERROR

Effect of Measurement Errors in X and Y on measured correlation and slope

typeset with L^AT_EX

Mathematical statisticians will probably instantly see the flaws in the following ‘intuitive’ (and shortcut) derivations by JH, which only apply exactly when the distribution of X is Gaussian – see Notes added 2021.09.13 in blue a few pages further down. But, since the results are so easily remembered, they are worth remembering – even if sometimes they may not be exactly correct. Remember George Box and his comment about all models being wrong, but ..

Denote the “true” (but unobservable) values by X and Y and the observed (error-containing) measurements by X' and Y' . We quantify the degree to which the errors in X' and Y' distort the correlation $\rho_{X,Y}$ and the slope $\beta_{Y/X}$

From the general formulae

$$\rho_{X,Y} = \frac{E[XY] - E[X] \times E[Y]}{SD[X] \times SD[Y]} = \frac{\text{Covar}[X,Y]}{\sqrt{\text{Var}[X] \times \text{Var}[Y]}} \quad (1)$$

and

$$\beta_{Y/X} = \frac{E[XY] - E[X]E[Y]}{\text{VAR}[X]} = \rho_{X,Y} \frac{SD[Y]}{SD[X]}, \quad (2)$$

we can derive the consequences of the errors in X and in Y .

Let $X' = X + \epsilon_X$ where ϵ_X has mean 0 and variance $\text{Var}[\epsilon_X]$, **and is independent of X** ⁶, so that

$$E[X'] = E[X + \epsilon_X] = E[X] + E[\epsilon_X] = E[X] + 0 = E[X] \quad (3)$$

$$\text{Var}[X'] = \text{Var}[X + \epsilon_X] = \text{Var}[X] + \text{Var}[\epsilon_X] \quad (4)$$

Let $Y' = Y + \epsilon_Y$, where ϵ_Y has mean 0 and variance $\text{Var}[\epsilon_Y]$, and is independent of Y , so that

$$E[Y'] = E[Y + \epsilon_Y] = E[Y] + E[\epsilon_Y] = E[Y] + 0 = E[Y] \quad (5)$$

$$\text{Var}[Y'] = \text{Var}[Y + \epsilon_Y] = \text{Var}[Y] + \text{Var}[\epsilon_Y] \quad (6)$$

$$E[X'Y'] = E[\{X + \epsilon_X\}\{Y + \epsilon_Y\}] = E[XY + \epsilon_X Y + \epsilon_Y X + \epsilon_X \epsilon_Y] = E[XY]. \quad (7)$$

By definition

$$ICC[X] = \frac{\text{Var}[X]}{\text{Var}[X] + \text{Var}[\epsilon_X]} \quad \& \quad ICC[Y] = \frac{\text{Var}[Y]}{\text{Var}[Y] + \text{Var}[\epsilon_Y]}, \quad (8)$$

with

$$0 \leq ICC[X] \leq 1 \quad \& \quad 0 \leq ICC[Y] \leq 1. \quad (9)$$

⁶ This is known as the ‘*Classical*’ error model.

1 $\rho_{X',Y'}$: expected correlation of two error-containing variables

$$\begin{aligned}\rho_{X',Y'} &= \frac{E[X'Y'] - E[X'] \times E[Y']}{SD[X'] \times SD[Y']} \\ &= \frac{E[XY] - E[X] \times E[Y]}{\sqrt{Var[X'] \times Var[Y']}} \\ &= \frac{Covar[X, Y]}{\sqrt{\{Var[X] + Var[\epsilon_X]\} \times \{Var[Y] + Var[\epsilon_Y]\}}}\end{aligned}$$

dividing above and below by $\sqrt{Var[X] \times Var[Y]}$

$$\begin{aligned}&= \frac{\frac{Cov[X, Y]}{\sqrt{Var[X] \times Var[Y]}}}{\sqrt{\frac{Var[X] + Var[\epsilon_X]}{Var[X]} \times \frac{Var[Y] + Var[\epsilon_Y]}{Var[Y]}}} \\ &= \frac{\rho_{X, Y}}{\sqrt{\frac{1}{ICC[X]} \times \frac{1}{ICC[Y]}}}\end{aligned}$$

so that...

$$\rho_{X',Y'} = \sqrt{ICC[X]} \times \sqrt{ICC[Y]} \times \rho_{X, Y} \leq \rho_{X, Y}.$$

Thus, the correlation is **attenuated*** (dampened/weakened) by the imperfections (random errors) in the X and Y measurements.

One can reverse the equation to get a “**de-attenuated**” correlation:

$$\rho_{X, Y} = \frac{\beta_{X', Y'}}{\sqrt{ICC[X]} \times \sqrt{ICC[Y]}}$$

<http://www.m-w.com/dictionary/attenuate>

*Main Entry: 1atátenáúáate ; Function: adjective

Etymology: Middle English attenuat, from Latin attenuatus, past participle of attenuare to make thin, from ad- + tenuis thin

1 : reduced especially in thickness, density, or force

2 : tapering gradually usually to a long slender point

2 $\beta_{Y'/X'}$: expected slope of error-containing Y on error-containing X

$$\begin{aligned}\beta_{Y'/X'} &= \frac{E[X'Y'] - E[X'] \times E[Y']}{Var[X']} \\ &= \frac{E[XY] - E[X] \times E[Y]}{Var[X] + Var[\epsilon_X]} \\ &= \frac{Covar[X, Y]}{Var[X] + Var[\epsilon_X]} \\ &= \frac{Covar[X, Y]}{Var[X]} \times \frac{Var[X]}{Var[X] + Var[\epsilon_X]} \\ &= \beta_{Y/X} \times ICC[X].\end{aligned}$$

so that...

$$\beta_{Y'/X'} = \beta_{Y/X} \times ICC[X] \leq \beta_{Y/X}$$

i.e., the slope is **attenuated** (dampened / weakened / flattened / moved towards 0) by the imperfections in the X measurements. Random errors in Y add to the residual variation, and thus increase the instability of the estimated slope, but do not (on average) attenuate the slope.

One can reverse the equation to get a “**de-attenuated**” slope:

$$\widehat{\beta}_{Y/X} = \frac{\widehat{\beta}_{Y'/X'}}{ICC[X]}$$

As Spearman told us, this method goes back to 1904. In the last 40 years, there have been major developments in the statistical handling of measurement error (see the textbooks mentioned at the beginning). One of the more creative is **Simulation Extrapolation Estimation** (**SIMEX**) in Parametric Measurement Error Models, Cook and Stefanski, (1994) JASA , 89, 1314-1328. See also the article Simulation-Extrapolation: The Measurement Error Jackknife by the same authors, as well as the simex Package for R.

3 Relationship between test-retest (X' , X'') and ICC[X]

X' and X'' denote 2 independent measurements of the X on a randomly selected individual, e.g., measuring one's cholesterol / height/ IQ twice in a short period of time, where X has not changed, and where ϵ_1 and ϵ_2 are independent.

In psychometrics, the term “*test-retest*” is **reserved** for a **self-administered** test, such as a questionnaire that is completed by the subject rather than an observer or test-administrator. Otherwise (e.g., if one wishes to study intra-observer or inter-observer variation) psychometricians speak of **observer variation**, rather than test-retest, studies.

Exercise: Show that

$$\rho_{X',X''} = ICC[X]$$

4 Relationship between $\rho_{X',X}$ and ICC[X]

This applies when we can think of X as the ‘gold standard’.

Exercise: Show that

$$\rho_{X',X} = \sqrt{ICC[X]}$$

Berkson error model

Results 1-4 are based on the **Classical** error model.

There is another, less common, error model, where the consequences are different. This is the ‘**Berkson**’ error model, named after this Mayo Clinic statistician/epidemiologist, Joseph Berkson.

- Again, we denote the “true” (but unobservable) values by X and the observed (error-containing) measurements by X'
- Let $X' = X + \epsilon_X$ where ϵ_X has mean 0 and variance $Var[\epsilon_X]$, **and is independent of X'**

This type of error is present in some environmental epidemiology studies, as nicely illustrated in the following excerpt from [this teaching piece](#).⁷

⁷Its author, Ben Armstrong, worked in Occupational Health at McGill for some years,

This distinction [between classical and Berkson error models] is not well known and a little tricky to understand, but it has major implications for the effects of the error.

- **Classical:** The average of many replicate measurements of same true exposure would equal the true exposure.
- **Berkson:** The same approximate exposure (proxy) is used for many subjects; the true exposures vary randomly about this proxy, with mean equal to it.

Example:

A study investigates the relation of mean exposure to lead up to age 10 with intelligence quotient (IQ) in 10 year old children living in the vicinity of a lead smelter. The IQ is measured by a test administered at age 10. Consider two study designs for assessing exposure:

- *Design 1:* Each child has one measurement made of blood lead, at a random time during their life. The blood lead measurement will be an approximate measure of mean blood lead over life. However, if we were able to make many replicate measurements (at different random time points), the mean would be a good indicator of lifetime exposure. This measurement error is thus classical.
- *Design 2:* The children's place of residence at age 10 (assumed known exactly) are classified into three groups by proximity to the smelter - close, medium, far. Random blood lead samples, collected as described in design 1, are averaged for each group, and this group mean used as a proxy for lifetime exposure for each child in the group. Here the same approximate exposure (proxy) is used for all subjects in the same group, and true exposures, although unknown, may be assumed to vary randomly about the proxy. This measurement error is thus Berkson type error.

Another situation giving rise to Berkson error is when exposures are estimated from observed determinants of exposure with an exposure prediction model. Often error has both classic and random components, although one usually predominates.

On page 6 of [these notes](#) JH gives an example of Berkson error, which he

and (as you may gather from his piece) was known as an outstanding teacher. When here, Ben worked with Gilles Thériault and other colleagues on the effects of electromagnetic fields on cancer rates in electric utility workers in Québec and France.

thinks he got from Ben Armstrong.⁸

It would apply if you recorded the temperature X' to which a stove or thermostat was 'set', but the true temperature X at the time the response (Y) was observed/measured could be somewhat above or below X' . The red dots in JH's diagram are what one might observe, and thus used to fit the regression model.

One way to think of the effect of Berkson error is that it moves the $Y[X']$ value *vertically*, but keeps it directly above the X' value.

By contrast, the effect of Classical error is that it moves the $Y[X]$ value *horizontally* to X' , but keeps it at the same vertical position.

Panel B (opposite) shows temperature fluctuations about a 'set' value.⁹

You could also 'see' why the Berkson error does not 'flatten the line' if (with X' – and thus X – centered) you examine the structure of $\hat{\beta} = \frac{\sum X' \times Y_{X'}}{\sum (X')^2}$.

NOTES added 2021.09.13

For many years JH has used 'results' (1) and (2) as if they were universally true, even though he has puzzled at the contradictory results obtained from simulations such as those in Q8 and Q9 below.

This year, he managed to get some clarifications from consulting the book *Measurement Error Models* by Wayne Fuller in 1989, and the introductory chapters in the newer version of *Measurement Error in Nonlinear Models* by Raymond Carroll et al. It appears that in order to prove that when X' contains **classical** error (uncorrelated with the true value X), then

$$E[\widehat{\beta}_{Y|X'}] = \beta_{Y|X} \times ICC[X],$$

we have to assume that $X \sim N(\mu_X, \sigma_X)$. Indeed, the specificity of the relationship is clear from a simple $n = 2$ example: take $X = \{-2, 2\}$, and $Y = \{-2, 2\}$, so that $\beta = 1$. Now add symmetric *classical* errors $\epsilon_1 = \{-1, 1\}$, and $\epsilon_2 = \{-1, 1\}$, so that the 4 equi-probable slope estimates are $4/\{6, 4, 4, 2\}$, or $2/3, 1, 1,$ and 2 ; i.e., their expectation is $7/6$! JH's proof ignores the fact that the denominator of $\widehat{\beta}_{Y|X'}$ is itself a random variable, and we know that expectations of ratios of r.v.s are complicated.

In contrast, the denominator of $\widehat{\beta}_{Y|X'}$ in the **Berkson** case of the 4 Fs measured at 4 'set' Cs is *fixed* in advance at $(-3)^2 + (-1)^2 + (1)^2 + (3)^2 = 20$. So the only random fluctuations are in the $\sum (X' - 17)(y - \bar{y})$ numerator: it is

$$(9/5) \times \{20 - 3 \times \epsilon_{C_1} - 1 \times \epsilon_{C_2} + 1 \times \epsilon_{C_3} + 3 \times \epsilon_{C_4}\}.$$

⁸This situation is alluded to in [this article](#).

⁹From the [thesis work](#) of Scott Weichenthal

From this small toy example you can work out the general algebraic form of the estimator.

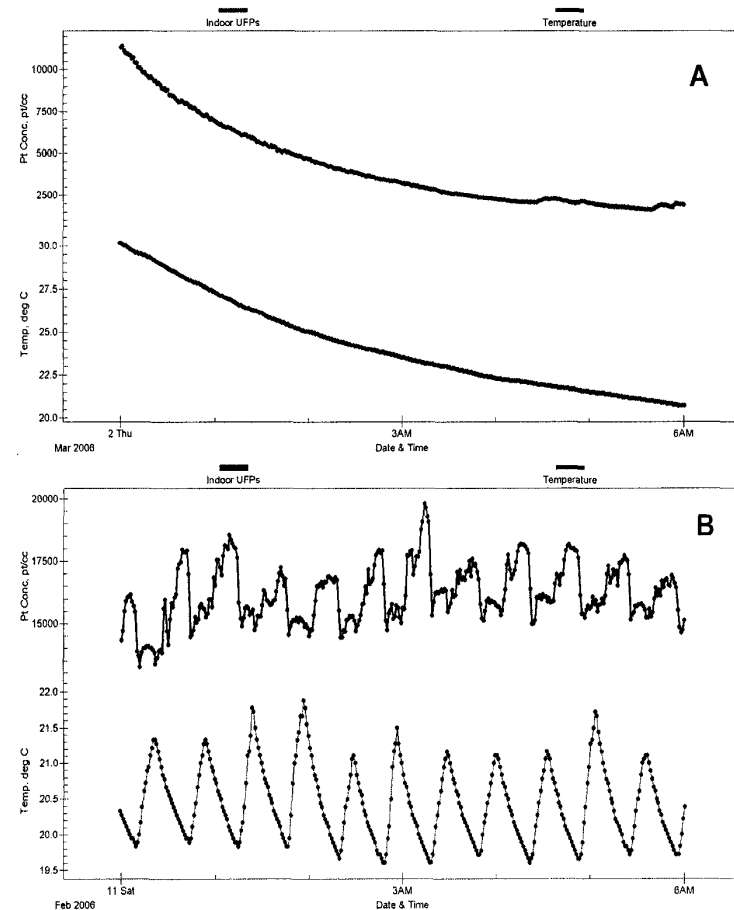


Figure 6a and 6b. Real-time overnight indoor UFP concentrations (cm^{-3} (top line) and temperature ($^{\circ}\text{C}$) (bottom line) as a sign of home heating system activation. (A) Home with a stand-alone wood stove, (B) Home with electric baseboard heaters.

EXERCISES...

1. Refer to the descriptions of the SMOG index, the Fry method, the Flesch Reading Ease, and the Flesch-Kincaid Grade Level, for measuring **readability** (under [Resources for Measurement/Surveys](#)).¹⁰ [Also, see Q29]

For the article or text you have chosen (as per discussion in class), randomly select three separate 100 word passages, and use this *set of three passages* to measure the readability (F_1) using the Fry graph. Rather than do so manually, you can use the SMOG calculator to determine the average number of sentences and syllables per hundred words. Repeat the readability measurement (F_2) with a second *different* set of three passages. Repeat once more (F_3), using a *third set*.

Using these same three sets, calculate the SMOG index, the Flesch Reading Ease, and the Flesch-Kincaid Grade Level.

For each index, use the 3 estimates to calculate the standard error of measurement, and the coefficient of variation. Comment.

2. Propose a method to assess the *validity* of a readability index.
3. [m-s] Derive the link between the standard error of measurement and the (intraclass correlation) reliability coefficient [last line, column 1, p16 in the notes above.] *Hint: it's simply a matter of using the definition of R.*
4. [m-s] Exercise in section 3 (p. 23 of notes) of Relationship between test-retest correlation and ICC(X) [In notes on Effect of Errors in X and Y on measured correlation and slope]
5. [m-s] Exercise section 4: Relationship between correlation(X, X') and ICC(X) [ibid.]
6. Francis Galton (1822-1911) found that the correlation between (*self-reported*) **parental** and (adult) **offspring heights** was strongest for the one between father and son [0.396 ± 0.024], and weakest for the one between mother and daughter [0.284 ± 0.028]. Given the way he obtained

the measurements, can you imagine why this was? ¹¹ [It was 0.302 ± 0.027 for mother & son; 0.360 ± 0.026 for father & daughter.]

7. **Bridging the physical- and the psycho-metric:** The notes on “Increasing Reliability by averaging several measurements” on the right hand column of page 13 of JH’s notes on Quantifying Reliability give the formula for the so-called “Stepped-Up Reliability”. In psychometrics (where the number of items on a test serves as the “several measurements”) this formula serves as the basis for the “Spearman-Brown prediction formula”.¹²

[m-s] Invert the formula on p.13 to derive the one on the right hand column of p.10 for the Spearman-Brown prediction formula relating the reliability of two versions of a test, one with N times more items than the other.

8. You are trying to estimate, from **imperfect observations** of F and C , the values of the two coefficients B_0 and B_1 in the temperature relation $F = B_0 + B_1 \times C$.

For each of the following situations, and using the true values $B_0 = 32$ and $B_1 = 9/5 = 1.8$, simulate¹³ 1000 datasets and investigate the behaviour of the 1000 estimates, b_0 and b_1 , of B_0 and B_1 . In each simulation, use samples of size $n = 4$, with temperatures of $C = 14, 16, 18$ and 20 .

- (a) C measured perfectly, F measured with $\epsilon_F \sim \text{Gaussian}(\mu = 0, \sigma_{\epsilon_F} = 1)$ errors that are independent of F . Check – formally, using a test (or CI) based on the mean of the 1000 estimates – for evidence of bias in b_1 . Also check whether the empirical variance of b_1 agrees with that given by the theoretical formula, namely

$$\text{Var}(b_1) = \sigma_{\epsilon_F}^2 / \sum (x - \bar{x})^2.$$

- (b) F measured perfectly, C measured with $\epsilon_C \sim \text{Gaussian}(\mu = 0, \sigma_{\epsilon_C} = 1)$ errors that are independent of C [*Classical type* error: someone

¹¹After you have thought about it for a while, and looked carefully at Galton’s [Notebook](#), you might wish to compare your answer with Karl Pearson’s explanation: “Why Galton got different parent-offspring correlations in heights” <https://jhanley.biostat.mcgill.ca/bios601/Surveys/pearson1930vol13ach14p17-18.pdf> and also look at “why he (KP) got larger ones” <https://jhanley.biostat.mcgill.ca/bios601/Surveys/PearsonBka1902pp377-378.pdf> using this protocol (p358-) <https://jhanley.biostat.mcgill.ca/bios601/Surveys/PearsonLee1903.pdf> in the ‘Measurement – Lecture Notes, etc’ section of the bios601 resources page for Measurement.

¹²Wikipedia has an entry called ‘Spearman Brown prediction formula’.

¹³If new to simulations, see “Computer code to simulate datasets with measurement error” <https://jhanley.biostat.mcgill.ca/bios601/Surveys/FandC.R.txt> at the bottom of the Resources webpage for measurement/surveys. It gives some ‘starter’ computer code, which you can modify to suit.

¹⁰ToneCheck (<https://techcrunch.com/2010/07/20/tonercheck/>) ‘sounded’ like an interesting tool; it’s not clear if ‘make it’ commercially, or was bought by another company!

else chose situations when C was indeed exactly 14, 16, etc, but didn't tell you what C was, and instead asked you to independently record C using your own imperfect instrument, and to use *your* recordings of C in your estimation of the equation]. Again, formally test for evidence of bias in b_1 .

- (c) F measured perfectly, C recorded from a thermostat that was 'set' to 14, 16, etc, where $\epsilon_C \sim \text{Gaussian}(\mu = 0, \sigma_{\epsilon_C} = 1)$ around the set value¹⁴ [*Berkson Error*]: you use the set values of C in your estimation of the equation]. Again, formally test for evidence of bias in b_1 .

Do your findings line up with the predictions in the Notes? If the patterns are difficult to see, you might change the number of simulations, the sizes of the errors, the range of C or the sample size.¹⁵

9. Attenuation of fitted 'F on C' slopes when progressively greater amounts of error are added to the C measurements

[a] Run the R code provided under the heading 'Animation (in R) of effects of errors in X on slope of Y on X'. <https://jhanley.biostat.mcgill.ca/bios601/Surveys/ErrorsInXAnimation.R.txt> It uses the 'animation' package to add progressively greater amounts of error to the C measurements and show how effects they affect the fitted slopes. Include the plot with your answers. Examine the trace of the fitted slopes, and try to mathematically link the pattern of the 'decay' with the amount of error. *Hint*: as we saw earlier, the attenuation should be a function of (actually, proportional to) the ICC_C ; so use the various amounts of error in C (ranging from $\sigma_{\epsilon_C} = 0$ to $\sigma_{\epsilon_C} = 22$) to calculate the various ICC_C 's and see if the predicted attenuations line up with the trace.

[b] [Added 2022, in light of the Note from 2021.08.13](#): For 15 years in bios601, this has been bothering JH. So now we can use simulations to see if the ICC-predicted relationship hold when the X s are *not spaced uniformly*, but are instead spaced as if they were drawn from a Gaussian distribution with the same mean squared deviation (MSD). So (even though it is hard with just 4 to have a Gaussian pattern*), make up 4 temperatures that have a mean of 17 C, and the same MSD as we have used in Q8, but are more concentrated around 17. and repeat the simulations in question 8(b). [*It might be easier if you gave yourself 40, rather than 4, X locations.]

10. Before we study how well we can digitize survival curves, here is an **exercise on communicating what the curves are meant to convey** and the context in which they were generated.

Refer to the article "Associations between C-reactive protein, coronary artery calcium, and cardiovascular events: implications for the JUPITER population from MESA, a population-based cohort study", available in the Resources link opposite 'Applications' in bios601. We digitized the lowermost (green) curve in Figure 2A of that article.

- (a) Read the Abstract and study the Figures in the article. Then, write, *in your own words*, a short news item of 250 words or so (2-3 minutes or so on radio) for your local newspaper and radio station, where you moonlight as a health reporter. In your piece address (i) the rationale for the study (ii) the principal findings and (iii) the implications of these findings. Also suggest a headline for your story. [You might want to study some health reports to see how they are structured.. the order may not be the (i)-(iii) order listed above. An interesting but slightly more highbrow website devoted to science reporting in general is <http://www.sciencedaily.com/>. The websites <http://www.cnn.com/HEALTH/>, <http://www.nytimes.com/pages/health/index.html>, <http://www.bbc.co.uk/news/health/> and <http://www.cbc.ca/news/health/> are also worth consulting, and indeed monitoring.
- (b) A 65-year old relative of yours reads your story, looks on the internet and finds that a test that measures coronary artery calcium is available in a private clinic in Montreal, and phones you to ask if it would be worth being tested and getting her "score". What would you say to this relative?

11. Errors in digitization

Refer to the duplicate readings you made of the Kaplan-Meier survival curve in the study entitled "Associations between C-reactive protein, coronary artery calcium, and cardiovascular events: implications for the JUPITER population from MESA, a population-based cohort study" available in the Resources link opposite 'Applications' in bios601

For now, ignore the point-wise measures of precision, i.e., the standard errors and confidence intervals, that often accompany such curves. These are (decreasing) functions of the numbers of subjects and the numbers of 'events'; we will cover their calculation later in the term. For now, focus only the loss of precision as a result of your digitization.

¹⁴Figure 6b on page 23 suggests it would not be exactly Gaussian.

¹⁵The article by Hutcheon et al. "Random measurement error and regression dilution bias", <https://jhanley.biostat.mcgill.ca/Reprints/RegressionDilutionBMJ.pdf> in the Resources for Measurement page tries to explain these patterns intuitively.

Focus on your two measurements of each of the reported y -year risks, where $y = 1, 2, 3, 4, 5, 6, 7$:

$$y\text{-year CHD risk} = 100 \times (1 - \text{proportion free of CHD at year } y)\%$$

- (a) From your two measurements at each of the 7 timepoints, obtain a 7d.f. estimate of the ‘standard error of measurement’. Do so using a ‘canned’ statistical routine and also ‘from scratch’ in R

Write out the statistical model that you used to obtain this, and list any assumptions it makes.

- (b) The estimate in (a) is an estimate of the ‘within’ observer variation. In order to estimate the ‘*between*’-observer variation, what is the *minimal* information you would need from each of you co-observers? (since JH has access to all of them, he will supply each of them once you email him with your specific request: he can supply the full raw data that could be then put into a canned statistical routine, but he would prefer that you do the calculations ‘from scratch’ in R).

Again, write out the statistical model that you used to obtain this, and list any assumptions it makes.

- (c) Here the ‘objects’ to be measured were 7 very specific (fixed) timepoints. Assume for the sake of this exercise that the 7 objects were 7 randomly selected human subjects and that we were interested in calculating an intra-class correlation coefficient to serve as a reliability measure. Carry out the ICC calculation. Restrict your attention to years 1-5 and recalculate the new ICC. Comment on why the ICC becomes smaller.

12. **Bernoulli Error?** A not-discovered-for-almost-300-years error in Bernoulli’s book? Or a not-discovered-for-almost-7-years error by A.W.F. Edwards. *Which is it?*

In his ‘Ars conjectandi three hundred years on’ article in Significance Magazine, Cambridge University Professor Edwards tells us that, a few years ago, he was reviewing Sylla’s English translation of (Jacob) Bernoulli’s book. He worked through one of the expectation problems, and came up with a different answer than Bernoulli. In early June of 2013, a week before the Edwards item was published in Significance, Julian Champkin, the magazine Editor, and a journalist by profession, used this ‘300-year-old error’ in the ‘trailer/teaser’ for the upcoming piece, and his question ‘Can you correct it?’ generated a number of responses on the Significance website.

JH has collected together in one .pdf file the item by Champkin, some of the original Bernoulli text in Latin, the full article by Edwards, the Edwards review of the Sylla translation into English, and Sylla’s translation of Bernoulli’s treatment of the problem.

The question arises as to whether it is the probabilities that are incorrect, or the expectation based on them, or whether it is Edwards who is incorrect.

What is your answer? [Remember that Edwards had studied Bernoulli earlier, when writing his book on Pascal’s triangle, and had found an error, that had been reproduced over the centuries in different books, in a table of Bernoulli numbers. So might Bernoulli (or the printers) had been a little bit careless?]

Here is the [R code](#) JH used to count the cases, and here is [Edward’s reply](#).

[Wikipedia entry for Edwards](#).

One wonders what he thinks of the removal of the [Latin Square](#) from the dining hall window at [Gonville and Caius College, Cambridge](#).

13. Imprecision in recording event times

The Introduction to a recent (2013) journal article “Driving under the (Cellular) Influence” by Saurabh Bhargava and Vikram S. Pathania of Carnegie Mellon University begins:

Does talking on a cell phone while driving increase your risk of a crash? The popular belief is that it does – a recent New York Times/CBS News survey found that 80 percent of Americans believe that cell phone use should be banned. This belief is echoed by recent research. Over the last few years, more than 125 published studies have examined the impact of driver cell phone use on vehicular crashes. In an influential paper published in the New England Journal of Medicine, Redelmeier and Tibshirani (1997) – henceforth, RT – concluded that cell phones increase the relative likelihood of a crash by a factor of 4.3. Laboratory and epidemiological studies have further compared the relative crash risk of phone use while driving to that produced by illicit levels of alcohol.

Later, in bios602, you will be introduced to the very clever study design that RT used to arrive at the 4.3.

The 2013 authors then go on to study the topic using a very different but also clever design.

We investigate the causal link between driver cell phone use and crash rates by exploiting a natural experiment induced by the 9pm price discontinuity that characterizes a majority of recent cellular plans. We first document a 7.2 percent jump in driver call likelihood at the 9 pm threshold. Using a prior period as a comparison, we next document no corresponding change in the relative crash rate. Our estimates imply an upper bound in the crash risk odds ratio of 3.0, which rejects the 4.3 asserted by Redelmeier and Tibshirani (1997). Additional panel analyses of cell phone ownership and cellular bans confirm our result.

But while they had very precise data on when cell phones were being used, (see Fig2) the data on crashes were quite messy. To quote the authors:

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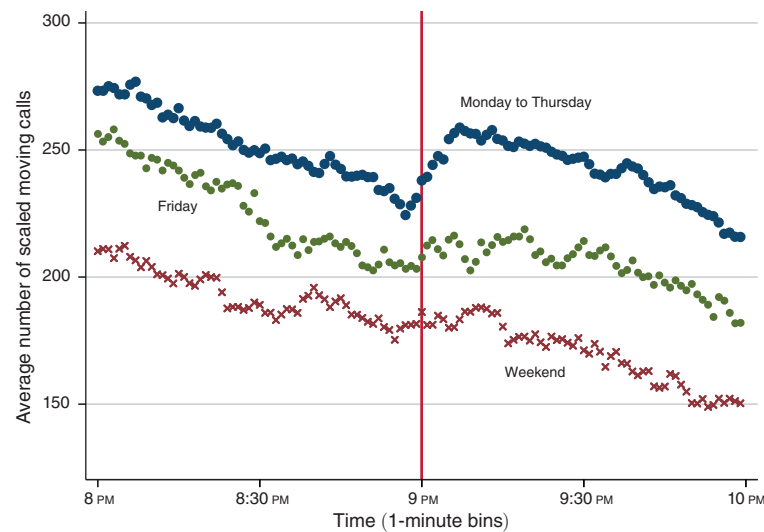


FIGURE 2. CELL PHONE CALL VOLUME FROM MOVING VEHICLES FOR CALIFORNIA FROM 8PM TO 10PM IN 2005

Our analysis principally relies on two sources of crash data. First, the State Data System (SDS) provides data for the universe of reported crashes from 1990 to 2005 for California, Florida, Illinois, Kansas, Maryland, Mississippi, Missouri,

Ohio, and Pennsylvania. A well recognized drawback of using a crash database based on self-reports is the presence of substantive periodic *heaping*.

The trajectory of a crash record helps to illuminate the origins of this bias. Once a vehicular crash is reported, police at the scene document various details of the incident, including the minute of the crash occurrence, and submits the paperwork to one of several possible state agencies. While states vary in the specifics that govern data collection and crash qualification criteria, crash records are ultimately centralized and sent once a year to the NHTSA where they are standardized and maintained.

Figure 4 illustrates the nature of the heaping in reports that characterizes a representative hour in 2005 across the states in our sample. A close examination indicates that nearly 11 percent of crash reports fall exactly on the hour, 31 percent are on the hour, half hour, or quarter hour, and 61 percent reside in a minute ending in either zero or five.

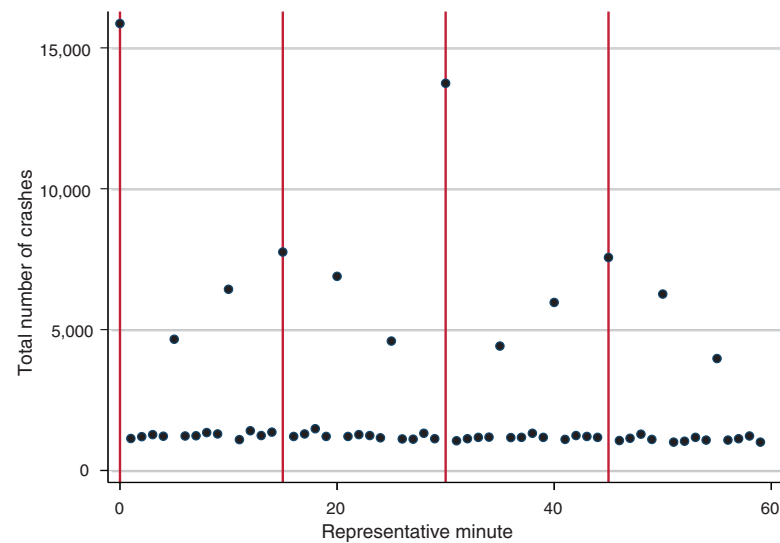


FIGURE 4. PERIODICITY IN SIDS CRASHES ACROSS REPRESENTATIVE HOUR IN 2005 FOR ALL STATES IN SAMPLE

Exercise: In this study, the primary contrast involves crash rates in the 1 hour after and the 1 hour before cellphone calls became “free” at 9 pm. Do you think the heaping errors are an insurmountable problem? If you do, why? If not, suggest ways to deal with them.

14. Galton’s data more than century later

[See also Questions 3-5 above, and see JH’s notes on Quantifying Reliability under the Measurement Lecture Notes heading in the website]

The 1985 article “Galton’s Data a Century Later” re-analyzes the extensive data collected by Francis Galton at his anthropometric laboratory in the South Kensington Museum in London.

JH has contacted one of the authors (Frank Ahern) who replied that “Despite a great deal of searching, neither I or Jerry McClearn have been able to find the original data that were used back in ’85.”

So, we will start again. But this time, instead of having to go to London and photocopy the records, you can take advantage of the scanned copies provided by the Wellcome Library and the Galton archives. To save you having to find the books (each containing about 500 records) in the large amount of material in the Galton archives, JH has down-

loaded them and put them on the bios601 website, in the Resources for Sampling/Masurement folder, under the heading (flagged in red) “Data from Galton’s Anthropometric Laboratory.”

For this exercise, which is designed to familiarize you with how to statistically quantify the psychometric (and psychophysical) properties of different measuring instruments, *we will focus on subjects who have been measured more than once*, so that we can assess the *reliability* of the various measures. For now, we will ignore the fact that there is quite a bit of time between some of the measurements, and that some attributes are age-related (we will try later to see at what age the peak is), and so some of the non-repeatability is for legitimate biological reasons.

So as to get a feel for the (small sample) sampling variability of these measures, and also so that it is not too big a data entry burden, you are asked to enter the complete records for 10 such subjects, i.e., subjects who were measured on more than one date. We can pool these student datasets later to get a more – statistically – reliable estimate of the various reliability measures.

In order to standardize the variable names, and provide a small element of quality control, a .csv file (**Spreadsheet for Data Entry**) with several subjects from the first book is provided on the website, immediately after the data books. Add to it the data for the *first ten* eligible ones you find in the range assigned to you (enter *all* of the records per subject, no matter how close or far apart they are in time). After you have added your entries, delete the ones already there — they were merely provided so as to standardize the naming of variables, and to act as a guide to align the columns correctly, and to make it easier to see any items that are mis-entered.

A few notes at this point (we may discover other oddities that we need to deal with as we go along). JH has noticed that subsequent measurements are some times recorded in metric units rather than Imperial (e.g., cm instead of inches and tenths or inches). We could discuss other ways to enter such mixed units (from JH’s past experience, converting as we enter is not an option!) but JH decided that when he met a metric measurement when he had allocated a pair of fields for say inches and tenths, he simply put the metric measurement in the first field and left the second field blank. It should be relatively easy to use programming to harmonize them later.

In the case of blanks, or illegible recordings, please leave the field blank.

JH has noticed some instances where there were several (4 in subject 0001) rows for the first several items (up to the Snellen test) but fewer

(e.g. 2 in subject 0001) rows for the later items at the bottom of the page, from sitting height to strength of blow with fist. In such instances, use any indications you can to decide which rows at the bottom of the page go with which ones at the top (in the case cited, JH decided that the first and fourth rows were complete, as were both of the bottom ones, so he put these with the first and fourth). In such cases, use the remarks column to flag the case.

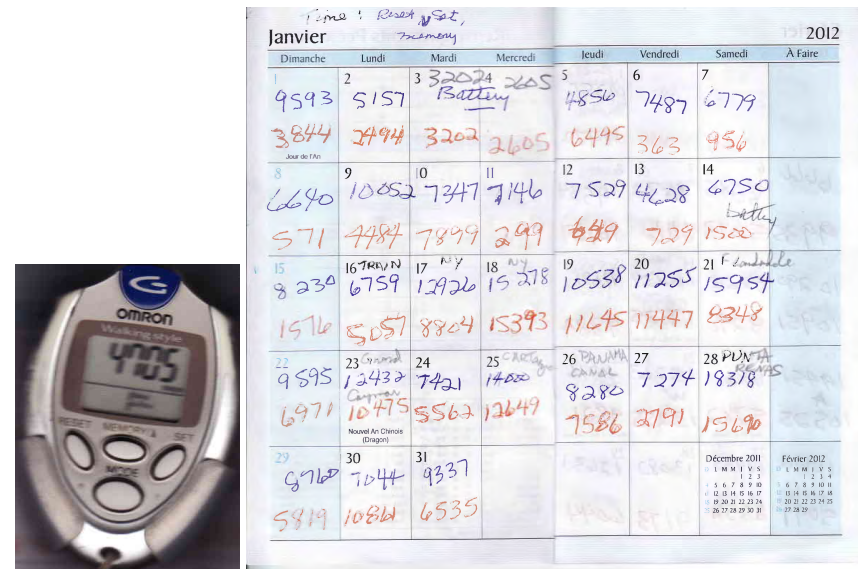
Here are the books assigned to the different students. Contact JH if your ID number is not in the list.

ID	Subjects
JH	0001-0491
26xxxxx21	0511-1028
26xxxxx19	1029-1530
26xxxxx57	1531-2020
26xxxxx99	2021-2520
26xxxxx78	2521-3021
26xxxxx65	3022-3521
26xxxxx58	3522-4000
26xxxxx90	4001-4500
26xxxxx94	4501-5000
	5001-5500
	5501-6000
	6001-6500
	7001-7459

Once you have entered the data, adopt the supplied R code to calculate the ICC for each of the measures shown in Table 1 of the 1985 article. Do not worry about timing or segregation by sex, or age-correction – you will not have enough data to do so; we will do this later when we pool the data. It appears (but JH is not entirely certain) that the 1985 authors used a simple Pearson product moment correlation with paired measurements. The advantage of the ICC is that while it is still connected mathematically with the Pearson correlation (see exercises above), it is more general and it uses whatever number of measurements per person there are. It is less cumbersome than using all possible pairwise correlations, or selecting just two.

Compare the ICCs with the test-retest correlations in Table 1 of the 1985 ‘a century later’ paper, and comment on any substantial differences.

15. Physical Activity: JH 2010-2017



The image shows an Omron step-counter on the left and a handwritten log book on the right. The log book is for the month of January 2012, with columns for each day of the week. The data is handwritten in red and black ink. The log book shows daily step counts and some remarks like 'Battery' or 'F. condole'.

Janvier	Dimanche	Lundi	Mardi	Mercredi	Jeudi	Vendredi	Samedi	A Faire
1	9593	5157	3202 2605 Battery	2605	4856	7487	6779	
2	3844	2494	3202	2605	6445	363	956	
3	6640	10052	7347	7146	7529	4628	6750	
4	571	4487	7899	299	629	729	1500	Battery
5	8230	6759	12926	15218	10538	11255	15954	
6	1576	5057	8804	15393	11645	11447	8348	
7	9595	12432	7421	14000	8280	7274	18378	
8	6971	10475	5562	12649	7586	2791	15690	
9	8960	7044	9337					
10	5819	10866	6535					

Since 2010, JH has used a ‘step-counter’ (pictured above left) to record how many steps he takes each day. His spouse AM has done the same, and has entered the pairs of daily counts onto a log book.

Refer to the six files (2010-2011, 2012-2013, 2014-2015, 2016-2017, 2018, 2019-2020) under the heading “Physical Activity: How many steps a day has JH being doing since 2010?” near the top of the Resources webpage. The 2010-2011 .csv file has the paired recordings for 2010, as well as JH’s ones for 2011. The 2012-2013, 2014-2015, 2016-2017, 2018 and 2019-2020 pdf files have scanned images (see above right) of the pages of paired recordings from the log-book.

The exercise in sampling from these data raised the issue of how many days one needs to sample in order to ensure that the estimate one gets is close to what one would obtain with a census, i.e., a 100% sample of days. Similar issues occur in dietary recall surveys. The least costly method is the food frequency questionnaire (Google for more info); a much more costly one is the x-day 24-Hour dietary recall method. How large x should be for different sub-populations (e.g., children, young adults, the elderly) has been studied. In measuring physical activity, it is common to use quite expensive accelerometers, and so they are usually given to research subjects for just one randomly chosen week.

The Omron model shown costs a lot less, and unlike the accelerometers –

which store minute by minute activity – just records the number of steps for each of the last 7 days. JH’s data help us answer the question of how many weeks are needed to get a good estimate of his yearly activity.

(a) divide the 2010-2111 data into weeks, and derive a (somewhat oversimplified) 1-way analysis of variance table, with week as the factor.

in this greatly oversimplified model, the numbers of steps (y) on any day (j) within week w ($i=1 \dots 104$) can be written as

$$y_{w,j} = \mu + b_w + \epsilon_{w,j}$$

(b) For didactic purposes, treat the model as a random-effects one, i.e., with week as the random factor. Thus, the 104 b_w ’s are assumed to be a random sample drawn from a $N(0, \sigma_w^2)$ distribution.¹⁶ Even though they may have a lot of structure, treat the variations across days within a week as uncorrelated ‘disturbances’ or ‘errors’ ($\epsilon_{yr,w,y,j}$) with variance σ^2 but no structure (i.e. treat all ϵ ’s as exchangeable, so that order of observations within the same week is irrelevant – in the file, you only need to know which week it is, not which day of the week. Clearly, there may be strong intra-week patterns, but for now assume that you are not even told which observation corresponds to which day of the week.

From the Expected Mean Squares (EMS) for this model¹⁷

Source	Sum of Squares	df	Mean Square	EMS
Weeks	SS_w	103	SS_w/df	$\sigma^2 + 7\sigma_w^2$
Error	SS_e	104×6	SS_e/df	σ^2

use the method of moments to estimate the σ_w^2 and σ^2 components.

(c) Using the results from (b), and the same overly simplified model, work out the expected variance of estimators that average recordings from (i) 3 random days in 1 random week (ii) 1 random day in each of 3 random weeks (iii) 3 random days in each of 3 random weeks.

(d) Could you have arrived at the results in (c) using the ‘Stepped-Up’ Reliability formula referred to in page 4 of the Quantifying Reliability notes?

¹⁶Using Roman b’s and Greek β ’s to distinguish random effects from fixed effects is a recent convention: it was not used when JH learned linear models.

¹⁷See also pages 4 and 5 of Notes on Introduction to Measurement Statistics, and pages 3 and 4 of the Notes on Quantifying Reliability (on the Resources website, under the heading ‘Measurement – Lecture Notes, etc’). ‘Weeks’ in the current example correspond to ‘persons’ or ‘subjects’ or ‘families’ in those examples.

See p7-8, [week to week variability in JH’s average steps per day, 2010-11.](#)

16. Repeatability of a Test – and of the statistical analysis itself!

Refer to the report ‘A Novel Test of Endurance Running Performance’ in the Resources website [under the tab ‘Data from various repeatability studies’]. as well as the [data and R code](#).

- Redo the 2-way ANOVA ‘with participant and trial as main effects’ to see if you can reproduce the reported coefficient of variation.
- Use a 1-way ANOVA, with subjects as a random effect, and the 3 trials as replicates (i.e. ignoring the order) and calculate an overall coefficient of variation. [A very similar 1-way ANOVA is shown in the 1st column of page 6 of the ‘Introduction to Measurement Statistics’ Notes on the Resources website. Page 11 of the Notes ‘Quantifying Reliability’ has an example with 2 measurements per family, but the principle is the same.] Which makes more sense to you, the CV based on their 2-way ANOVA, or yours based on a 1-way ANOVA?
- Calculate subject-specific coefficients of variation (just as was reported in Table 1 in the article on breath alcohol – the link to this article can be found just above the one for the endurance test). Summarize the 10 CVs using say the median and the range. Would you report the ‘overall’ CV the authors did, or some summary of the 10 subject-specific ones? Give a reason for your choice.
- Use the results of the 1-way ANOVA¹⁸ to calculate an intra-class correlation (ICC).
- In this setting, which makes more sense, a CV or an ICC? Why?
- Rerun the ICC code several times on random subsets of the subjects. As you reduce the sample size to just 2 or 3, does the ICC stay stable? Use the example to say what the ICC tells us that the CV can not, and what the CV tells us that the ICC can not.
- How could one ‘rig’ (i.e., manipulate) the sample of subjects in the breath alcohol study to (i) maximize (ii) minimize the ICC?

¹⁸The R code supplied makes use of an ICC package, but it is always safer to check with a worked example that a package you don’t know is doing what you want it to do.

17. **How reproducible and accurate are free smartphone apps to track your steps, calories burned, distance and active time?**

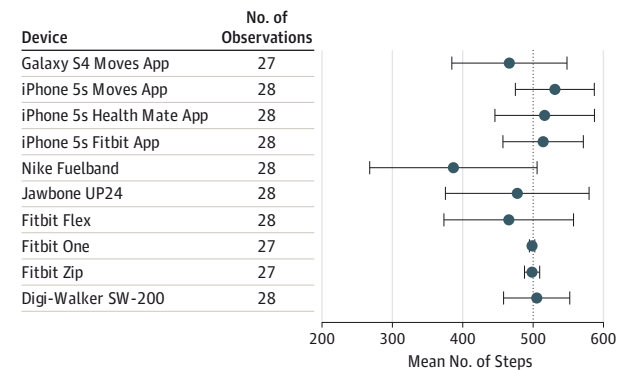
The letter ‘Accuracy of Smartphone Applications and Wearable Devices for Tracking Physical Activity Data’ in JAMA in February 2015 [under the tab ‘Data from various repeatability studies’] reports

This prospective study recruited healthy adults aged 18 years or older through direct verbal outreach at a university. Participants gave verbal informed consent to walk on a treadmill set at 3.0 mph for 500 and 1500 steps, each twice, for no compensation. An observer (M.A.C.) counted steps using a tally counter in August 2014. This study was approved by the University of Pennsylvania institutional review board.

A convenience sample of 10 applications and devices was selected from among the top sellers in the United States. On the waistband, each participant wore the Digi-Walker SW-200 pedometer (Yamax), which has been well validated for research,⁶ and 2 accelerometers: the Zip and One (Fitbit). On the wrist, each wore 3 wearable devices: the Flex (Fitbit), the UP24 (Jawbone), and the Fuelband (Nike). In one pants pocket, each carried an iPhone 5s (Apple) simultaneously running 3 iOS applications: Fitbit (Fitbit), Health Mate (Withings), and Moves (ProtoGeo Oy). In the other pants pocket, each carried the Galaxy S4 (Samsung Electronics) running 1 Android application: Moves (ProtoGeo Oy).

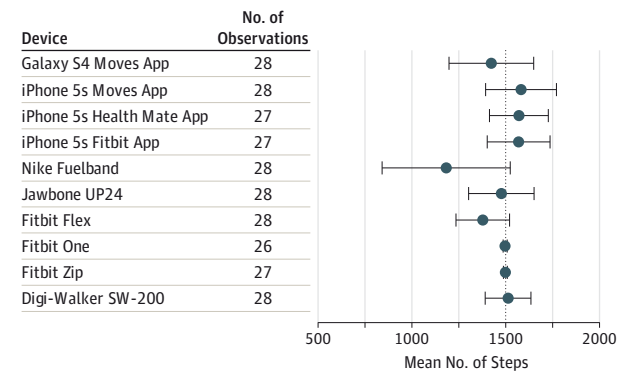
Across all devices, 552 step count observations were recorded from 14 participants in 56 walking trials. Participants were 71.4% female, had a mean (SD) age of 28.1 (6.2) years, and had a mean (SD) self-reported body mass index (calculated as weight in kilograms divided by height in meters squared) of 22.7 (1.5).

Figure 1. Device Outcomes for the 500 Step Trials



The vertical dotted line depicts the observed step count. The error bars indicate ± 1 SD.

Figure 2. Device Outcomes for the 1500 Step Trials



The vertical dotted line depicts the observed step count. The error bars indicate ± 1 SD.

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Figure 1 shows the results for the 500 step trials by device and Figure 2 shows the results for the 1500 step trials. Compared with direct observation, the relative difference in mean step count ranged from -0.3% to 1.0% for the pedometer and accelerometers, -22.7% to -1.5% for the wearable devices, and -6.7% to 6.2% for smartphone applications. Findings were mostly consistent between the 500 and 1500 step trials.

- (a) Rewrite the authors findings using the words ‘under-’ and ‘over-counted.’
- (b) For which instruments is there evidence that this ‘bias’ is non-zero? *You can use your eye to determine the means and SDs, or use the ones in the .pdf file shared by senior author (‘I’m attaching the raw data that we have to share’) and available on the course website.*
- (c) The data summaries were in response to an email from JH to the author, asking if there was ‘any chance you would be able to share the Excel file of raw data, so we should see if the deviations from the target were all over the place, or peculiar to a few people or a few devices. I can imagine the pockets on some people being a bit deep and wide.. and that the machines in them slosh around – I sometimes keep my \$20 dollar step counter in my pocket instead of on my belt.’

Imagine that the author had shared these data as 552 separate lines, each one containing a step count, a participant ID (1-14), the target (500 or 1500), the occasion (1st or 2nd) and the name of the device.¹⁹ Write out a plan for analyzing them, including the model you would use, the meaning of each component (parameter) in the statistical model, how you would estimate each component, a table of results (use made up, but realistic numbers), and a sketch of one or more graphs that would quickly tell the same story.

- (d) In the Fall of 2016, the EPIB601 class carried out its own investigations. The Epidemiology teacher tested an app called Pacer - Pedometer plus Weight Loss and BMI Tracker By Pacer Health, Inc that is available for free for both the iPhone and Android devices. Dr Patel (senior author of the letter) ‘particularly like[d] Withings HealthMate because it has a good user interface and works with both iPhones and Androids. Fitbit is also good but works with a limited set of Androids.’

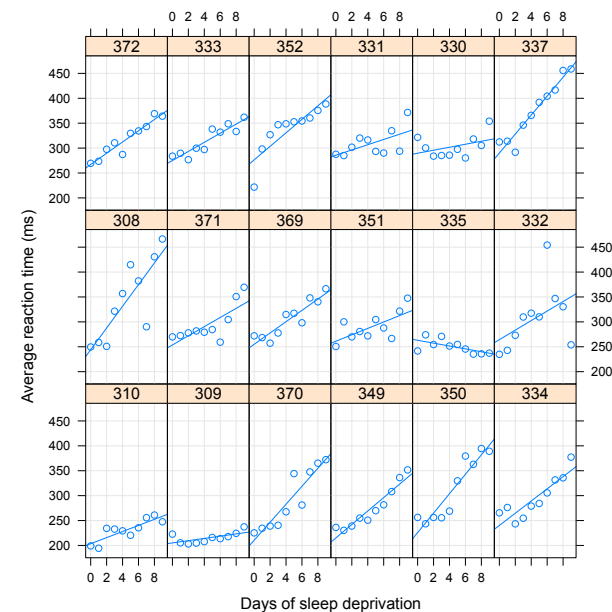
For the BIOS601 of 2016, students were asked to prepare to participate in a planning session, where together they would design (and subsequently carry out) their own investigation into the reproducibility and validity of a few smartphone apps with respect to steps, distance, calories, etc

¹⁹At the end of each trial, step counts from each device were recorded. In rare instances that a device was not properly set to record steps (8 of 560 observations), these data were not included. The mean step count and standard deviation for each device was estimated using Excel (Microsoft). Across all devices, 552 step count observations were recorded from 14 participants in 56 walking trials.

18. Reaction times

The orientational material below is from the **sleepstudy** data re-analyzed in Ch. 3 of the excellent (online) book ‘lme4: Mixed-effects modeling with R, dated June 25 2010, by Douglas M. Bates. The data are included in the **lme4** package – and were used again in the 2017 Epidemiology (teaching) article Sample Size Estimation for Random-effects Models: Balancing Precision and Feasibility in Panel Studies by Weichenthal, Baumgartner and Hanley.

Belenky et al. [2003] report on a study of the effects of sleep deprivation on reaction time for a number of subjects chosen from a population of long- distance truck drivers. These subjects were divided into groups that were allowed only a limited amount of sleep each night. We consider here the group of 18 subjects who were restricted to three hours of sleep per night for the first ten days of the trial. Each subject’s reaction time was measured several times on each day of the trial.



‘Average reaction time versus number of days of sleep deprivation by subject for the **sleepstudy** data. Each subject’s data are shown in a separate panel, along with a simple linear regression line fit to the data in that panel. The

panels are ordered, from left to right along rows starting at the bottom row, by increasing intercept of these per-subject linear regression lines. The subject number is given in the strip above the panel.⁷

The 2003 article [European Sleep Research Society, *J. Sleep Res.*, 12, 1-12] that Bates cites is more specific about the Psychomotor vigilance test (PVT), and the number of trials (JH estimates 100 or so) that went into each datapoint shown in the graph [note that Bates used the average response latency whereas Belenky used its reciprocal.]

The PVT measures simple reaction time to a visual stimulus, presented approximately 10 times/minute (interstimulus interval varied from 2 to 10 s in 2-s increments) for 10 min and implemented in a thumb-operated, hand-held device (Dinges and Powell 1985). Subjects attended to the LED timer display on the device and pressed the response button with the preferred thumb as quickly as possible after the appearance of the visual stimulus. The visual stimulus was the LED timer turning on and incrementing from 0 at 1-ms intervals. In response to the subject's button press, the LED timer display stopped incrementing and displayed the subject's response latency for 0.5 s, providing trial-by-trial performance feedback. At the end of this 0.5-s interval the display turned off for the remainder of the foreperiod preceding the next stimulus. Foreperiods varied randomly from 2 to 10 s. Dependent measures, averaged or summed across the 10-min PVT session, included mean speed (reciprocal of average response latency), number of lapses (lapse = response latency exceeding 500 ms), and mean speed for the fastest 10% of all responses.

In bios601, each of you will make some rough ('amateur') reaction time measurements, so as to learn what your reaction times are like, and to plan a study into whether they are faster when using your dominant rather than your non-dominant hand.

The 2003 measurements relied on a thumb-operated, hand-held device and a microcomputer program described in 1985.

To make your own measurements, you can choose this quite intuitive web tool <https://faculty.washington.edu/chudler/java/redgreen.html> – and use either the keyboard or the mouse/trackpad. It only performs and shows the results of 5 trials at a time. So – since you will need to calculate the mean and SD of 10 individual times – you will need to copy the individual times into R, 5 at a time.

[To get around this, JH wrote a simple R program that may not be as accurate or fancy but that stores the individual times from however many you do into a vector. Links to web-based tools, and to some scholarly and newspaper articles on reaction times) are available under [Online Tools](#) on the webpage for the Resources for measurement.]

The main objective is to gain experience with 'hands on' data, and with sample size planning, so try both tools and choose between them.

[If you have energy to spare, you can try to empirically determine how closely this R-based instrument and the web-based instrument agree.]

Before running the measurements, be sure to practice first.

- (a) Run 10 trials using your dominant hand, and calculate the mean reaction time, the SD, and the SE of the mean (SEM).

Convert the SEM into a coefficient of variation (CV²⁰). How does *this* CV (which measures the 'instability' of the *mean*) relate to the CV for *individual* measurements?

Use the SEM to calculate a 95% confidence interval to accompany your point estimate of the true mean. Why use a larger-than-1.96 multiplier to calculate the margin of error?

- (b) Suppose you wished to perform enough trials that the margin of error would be less than 5% of the mean. Using the SD (or SEM, or CV) you already obtained²¹, calculate how many trials you would need.

Guidance on such sample size *considerations* (JH prefers this term over sample size *requirements*) can be found in section 4 of his bios601 Notes on Mean/quartile of a quantitative variable:- models / inference / planning

- (c) Suppose you wished to (i) test whether, or (ii) measure how much, the mean of reaction times (*r.t.*) obtained with your dominant hand (D) differs from the mean of reaction times obtained with your non-dominant hand (ND).

²⁰When reporting a CV, it is customary to do it so as a *percentage*

²¹Of course, if you were to run that many trials, there is no guarantee that the SD would be the same as the SD you got for the 10 – it could be higher or it could be lower. But use the SD of the 10 as the best guess for planning purposes

You will make n measurements with each hand. Assume that there is no ‘fatigue factor’ or ‘order-of-testing’ effect, so that it doesn’t matter whether you first do the n with one hand and then the n with the other. [If there were a fatigue factor, or order effect, then we would want to think of other designs, possibly involving pairing/blocking].

The 2 n ’s may be large enough that the relevant sampling distribution of the difference of two independent sample means (Student’s t) is close to a Z distribution; otherwise, use trial and error. Also assume that the variability is about the same in both r.t. series.

For (i) you will use a 95% confidence interval for the difference of two unknown means, $\mu_D - \mu_{ND}$.

For (ii) you will use the test statistic $\frac{\overline{r.t.D} - \overline{r.t.ND}}{SE \text{ of this difference}}$, and $\alpha = 0.05$ (2-sided).

For the *estimated difference* determine the n per hand that would yield a margin of error of at most: 10 milliseconds; 5 milliseconds.

For the *statistical test* determine the n per hand that would give you an 80% chance of obtaining a ‘statistically significant’ test result if the true difference in milliseconds were: 5, 10, 25.

For the *statistical test*, also determine the chance of obtaining a ‘statistically significant’ test result (the statistical ‘power’, or $1-\beta$) if each n is fixed at 25, but the true difference in milliseconds was: 1, 5, 10, 25.

What if the SD you used for planning was too large? too large?

- (d) Do a few trials using the tool <https://www.justpark.com/creative/reaction-time-test/> that was featured in the newspaper story ‘[Brain test judges how old you are based on your reaction time.](#)’ Consider their reaction-time vs. age curve, and how it was fitted. The website don’t say (i) how they selected the 2,000 people aged 18 and above that they surveyed, or (ii) how many trials they asked each of them to do.

As for (i), describe one scenario where the curve they obtained would be ‘flatter’ than the one that would be obtained if representative population-based samples were recruited at each age.

Suppose²² that each of the very large number of subjects in each 1-year-wide age-bin was tested a very large number of times. Suppose then that within each age-bin we sorted the persons from slowest to fastest and selected the ‘median’ (middlemost) person. Suppose further²³ that from age 25 to age 64, these medians made an almost perfect straight line with slope 2 ms per year of age, or 0.5 years of age per ms of response latency if we plot age on the vertical (y) axis and response latency on the horizontal (x) axis.

For now, we will retain these 40 people from this ‘ideal’ world. As for (ii), we will ask them to make *just 1 trial each*, and (like the website) use these 40 values to fit the LS line of age(y) upon latency(x).

Assuming within-person variation of the same magnitude as in your own set of measurements, what is your best estimate of what the fitted slope will be? *Hint*: remember some earlier exercises.

The above scenario selected the *median* person in each bin. If you picked one *random* person from each bin, what is your best estimate of what the fitted slope will be? (State your assumptions).

Write a few sentences summarizing why (even if their sample of subjects is representative) the age-latency graph in the website may be inaccurate, and in what respect.

- (e) What if each median-person’s latency was measured perfectly (large n), but ages were in bins (intervals) 5 years wide (so that, e.g., the persons aged 25, 26, 27, 28 and 29 are put at age 27), and we fitted the LS line of latency(y) upon the *midpoint* (x) of each age bin?

²²This ideal universe where subjects are easily recruited, and have lots of patience and can maintain their attention over a very large number of trials, is just for didactic purposes.

²³Now we are really dreaming! While we are at it, we will assume symmetric age-specific distributions.

Note re terminology:

In the situation where x = latency, the errors in measuring the true X values are uncorrelated with these true values of X . This is called the *classical* ‘errors in X ’ situation. It is the nastier case.

$$X = \text{true value}; \quad x = X + \epsilon_X, \text{ with } \epsilon_X \perp X$$

In the situation where x = the mid-age of the bin, the errors in measuring the true X values (ages) are correlated with the true values of X , but uncorrelated with the observed x ’s. This is called the *Berkson* ‘errors in X ’ situation. It is less nasty, but it does increase the (sampling) variability of the estimated slope.

$$X = \text{true value}; \quad x = X + \epsilon_X, \text{ with } \epsilon_X \perp x$$

JH’s favourite example of Berkson error (one he adapted for the earlier exercise on F v.s C temperatures) is one that may have come from Berkson himself: An investigator wished to measure temperatures in an oven at various times.

- An unreliable thermometer, i.e., one that gives readings that fall equally on both sides of the truth, would generate classical errors.
- The temperatures shown on the thermostat are as likely to be above/below the true temperature at any given moment of interest; as you can check, these would be Berkson errors.

For more on these topics, you might consult JH’s Ch. 4 notes, Applied Linear Models, course 679, or the books and presentations by the (measurement-expert) statistician Raymond Carroll <https://www.stat.tamu.edu/~carroll>

19. **Instead of measuring heights with a tape, how about using a smartphone app?**

Q. prompted in 2019 by revisiting Pearson’s protocol, and by this piece, <https://lifehacker.com/which-ar-measuring-app-is-more-accurate-1827242756>, found when searching ‘measuring heights smartphone’. As of 2020, one can find this for the iPhone, iPad, or iPod touch. Think about ways to test its validity and reproducibility.

20. **Who seems to age faster?** The following are the reported ages of the 40 students in JH’s course 513-607 (Inferential Statistics) in 1986.

```
AGE.1986 = c( rep(22,4), 23, 25, rep(26,3), rep(27,4), rep(28,3),
             rep(29,2), rep(30,5), rep(31,4), rep(32,2),
             rep(33,2), rep(34,2), 35, 36, 37, rep(38,2), 39,42 )
```

- Make a new variate AGE.1999 from the AGE.1986 variate.
- use the `lm` function to estimate, from the regression of AGE.1999 on AGE.1986, how much these students aged in the intervening 13 years.
- Notice the use of uppercase AGE to denote the true age. What if these 40 students had reported their 1986 ages as their true 1986 ages ± 5 years (with the - or + determined at random, without regard to the person’s true age)? i.e. as (say) `age.1986 = AGE.1986 + 5*sample(c(-1,1), 40, replace=TRUE)`. Note the use of lowercase `age` to denote the ‘error-containing’ value [In the measurement error literature, and in JH’s notes, it is common to use X and Y for the true values and X^* and Y^* , or X' and Y' , for the error-containing values.]
Now, again, use the `lm` function to estimate, from the regression of AGE.1999 on `age.1986`, how much these students aged in the intervening 13 years, and who aged the most and who the least. Comment on your findings, and give a non-technical explanation that your engineer-sibling would understand.
You might want to simulate several `age.1986` vectors to convince yourself that the effects are reproducible. Or – if keen on algebra – work out how much, on average, the attenuation is.
- Apply the `lm` function again, but this time with AGE.1999 – `age.1986` as the ‘ y ’ variate, and `age.1986` as the ‘ x ’ variate. Comment.

21. **Some ‘big-ticket’ epidemiology examples.** Refer to the 1997 article The INTERSALT study: background, methods, findings, and implications. Am. J. Clin. Nutr. 65, 626S?642S. by J Stamler²⁴ It is available here: <https://jhanley.biostat.mcgill.ca/bios601/Surveys/InterSALT.pdf>

The last 2 columns of Table 2 on p630S are entitled ‘Observed coefficient as percentage of true coefficient’ when there is one measurement and when there are four. The footnote explains how it is calculated from the ratio of the intra- to inter-individual variance and the number of measurements.

²⁴cited on p. 72. of Chapter 4 (Principles of measurement) of Cox and Donnelly’s 2011 book Principles of Applied Statistics, available as an eBook from McGill. <https://www-cambridge-org.proxy3.library.mcgill.ca/core/books/principles-of-applied-statistics/E7225E64F86B2C8193CA3C57621B6338>.

- (a) Verify the calculations for 24-h Urinary Na excretion.
- (b) Suppose you had the ICC (rather than the Intra- to inter- variance) as the column header. Alter the wording of the footnote accordingly.
- (c) What is the relation between these and the ‘stepped-up reliability’ measures addressed in question 7?
- (d) Show how the numbers in row 2 of Table 3 were derived from those in row 1.
- (e) Use JH’s daily steps in 2010 (see Q. 15) to work out an ‘intra-’ variance. We will assume JH’s intra- is typical of the ‘intra-’ variance of other people of JH’s age.
- (f) We don’t have average steps per day for that year for many other people his age, but assume we did have it for a large number of individuals, and that the mean of this large number of person-specific yearly averages is 6,000 steps/day, and the (inter-individual) SD is 2,000 steps/day. Using this SD, and the results from (c), to add a row, entitled say ‘activity, measured as steps/day,’ to Table 2.
- (g) Comment on the slope for the leftmost portions of the relationship in the top left panel (Total physical activity (cpm)) of Figure 2 of the article “Dose-response associations between accelerometry measured physical activity and sedentary time and all cause mortality: systematic review and harmonised meta-analysis.” The file is here: <https://jhanley.biostat.mcgill.ca/bios601/Surveys/PhysActivityAllCauseMortality.pdf> Since the y axis is on a log scale, it shows $\log(\text{Hazard ratio})$ v.s physical activity as being approximately linear for the early part.
Try to de-attenuate the slope, by assuming (a bit unrealistically) that the measurements shown on the X axis are based on 7 random days over a year, and that the intra- to inter- variance ratio you calculated in part (d) applies to the Total physical activity (cpm) measurements in this study.
- (h) Refer to the article “Association of Office and Ambulatory Blood Pressure With Mortality and Cardiovascular Outcomes.” The file is here <https://jhanley.biostat.mcgill.ca/bios601/Surveys/BP-1-time-24-hr.pdf> Which blood pressure index would you expect to have the strongest relationship with mortality rates, and why? Are your expectations borne out?

22. Measuring COVID-19 Antibody Seroprevalence in Santa Clara County, California in early April 2020:

The following is adapted from the [full report of April 11](#).

Methods: In early April, 2020, California investigators tested Santa Clara county residents for antibodies to SARS-CoV-2 using an immunoassay. Participants were recruited using Facebook ads targeting a representative sample of the county by demographic and geographic characteristics. They reported the prevalence of antibodies to SARS-CoV-2 in a sample of 3,330 people [2,718 adults and 612 children], adjusting for zip code, sex, and race/ethnicity. They also adjusted for test performance characteristics using 3 different estimates: (i) the test manufacturer’s data, (ii) a sample of 37 positive and 30 negative controls tested at Stanford University, and (iii) a combination of both.

Results: The unadjusted (crude) prevalence of antibodies to SARS-CoV-2 in Santa Clara County was $[50/3,330 =] 1.50\%$ (exact binomial 95CI 1.11-1.97%) [A], and the population-weighted prevalence was 2.81% (95CI 2.24-3.37%) [B]. Under the three scenarios for test performance characteristics, the population prevalence of COVID-19 in Santa Clara ranged from 2.49% (95CI 1.80-3.17%) to 4.16% (2.58-5.70%) [C]. These prevalence estimates represent a range between 48,000 and 81,000 people infected in Santa Clara County by early April, 50- 85-fold more than the number of confirmed cases. [taken from April 11 report]

The questions below are taken directly from the Part A (bios700) exam of August 4, 2020.

For questions a-f, ignore the misleading wording in A and B, which refer to the proportion of positive tests (there were 50 positive tests), not the prevalence of antibodies. (In the statistical analyses section, they use the term ‘frequencies of positive tests as a proportion of the sample size.’)

- (a) Explain how the exact binomial CI in [A] is calculated. How different is it in this instance from the usual ‘non-exact’ test?
- (b) Give a reason why neither the ‘exact’ nor the ‘inexact’ CI around the 1.5% is relevant. (You might delay answering until you have been through the remaining questions)
- (c) Explain in more detail how you think they arrived at [B], i.e., the prevalence of 2.81% and the 95% CI 2.24-3.37.
- (d) Given that there are 57 zip codes, do you see any issues in calculating a SE? Explain.
- (e) Suggest one other way of arriving at a point and interval estimate of the county percentage [incidentally, besides the non-representativeness of the sample with respect to zip code, sex, and race/ethnicity mismatches, the age distribution of the sample did not match that of the county either.]

- (f) They say in the main text that their 2.24-3.37 CI around the 2.81% was computed ‘without clustering the standard errors for members of the same household’ and that it was 1.45-4.16 when this clustering was taken into account. Explain why the CI that takes account of the household clustering is the more appropriate of the two, why it is wider, and one way you would calculate it.
- (g) In A and B, by using the terms ‘test positivity’ and ‘prevalence’ interchangeably, the authors are implicitly assuming that the test is perfect, i.e., always positive when antibodies are present (100% *sensitive*), and always negative when antibodies are absent (100% *specific*). In C, they “adjusted the prevalence for test sensitivity and specificity. Because the SARS-CoV-2 assays are new, [they] applied three scenarios of test kit sensitivity and specificity.” The first scenario used the manufacturer’s validation data (*sensitivity* 91.8%; *specificity* 99.5%).’
- (Unrealistically) for now, take the given *se* and *sp* values as having negligible sampling error, and ignore the fact that the sample of 3,330 is not representative of the county. Derive the estimating equation whose solution gives the Maximum Likelihood estimate of the population prevalence, θ . (You don’t need to solve the estimating equation.)
 - You can get to the same estimating equation faster by the method of moments, so what is the advantage of going the ML route?
 - In fact, the manufacturer’s 91.8% sensitivity was the proportion 78/85, and the 99.5% specificity was 369/371. The smaller pilot dataset from Stanford gave $se = 25/37 = 67.6\%$ and $sp = 30/30 = 100\%$. How would you incorporate the inherent sampling error in *se* and *sp* into the CIs for θ .
- (h) The concepts of sensitivity and specificity are typically taught in the context of positive and negative predictive values of diagnostic tests. In this SARS-CoV-2 immunoassay context, what would predictive values refer to? And how does this focus differ from the focus of the Santa Clara study?

The first version of the report was the subject of a [very extensive statistical blog](#) (as well as many many politically-based) ones.

These resulted in a [second version](#), that prompted another round of statistical attention, and even a [full paper](#). This topic is an old one, and just-now-retired [McGill biostatistics professor Lawrence Joseph](#) played a

pioneering role – well before prevalence estimation became so important. Interestingly, the references in the full paper don’t go back many years.

For a newer [as of July 28, but this topic is moving fast!] report covering more space and time, see [this JAMA article](#) *Seroprevalence of Antibodies to SARS-CoV-2 in 10 Sites in the United States, March 23-May 12, 2020*.

You will see that a key driver remains the *specificity* of the test, and its critical presence in both the numerator and denominator of the prevalence estimator.

As you will find by Googling say ‘dashboard seroprevalence covid’ there are now several dashboards, of variable quality, some showing international comparisons. [Here is one](#). It is not always clear if all of the reported prevalences make the correction for the operating characteristics of the tests used, or how estimates are combined, or how calendar time is taken into account.

23. A 165-year rewind: John Snow’s data on cholera deaths in customers of two Water Companies, South London, Fall 1854, and his additional 1856 report:

Of the internet ‘shrines’ to John Snow and his work elucidating the mode of transmission of cholera, the foremost, and the earliest is the [UCLA site](#). Others are the one at Michigan State University, maintained by authors of the definitive biography of Snow, and the [John Snow Society](#) at the London School of Hygiene and Tropical Medicine.

On JH’s site that he prepared for his lecture in the [EBOH 2014 lecture series for the public](#), you will find further material and links.

While the full John Snow story is quite extensive, most courses limit their coverage to the Broad Street Pump episode. Interestingly, in his 1855 book, Snow gave this fewer pages (and less weight?) to this than he gave to the South London data, which arose from what he thought of as ‘The Grand Experiment.’ The account of this begins in what the UCLA site used to call ‘part3’ of the (now online) recreation of the book. Snow’s own observations began in 1832 in the coal mines in northeastern England 1849, and it was during the 1949 epidemic that he published his short pamphlet. So you might want to start in 1849, and his section *Influence of the water supply on the epidemic of 1849, in London*, and how he exploited the fact of the ‘New water supply of the Lambeth Company,’ and the ‘**Intimate mixture of the water supply of the Lambeth with that of the Southwark and Vauxhall Company**’.

For this exercise, focus on the ‘Result of the inquiry as regards the first

four weeks of the epidemic in 1854’ – an inquiry he carried out by personally visiting the houses of the first 334 who died. The purpose was to learn which water company (or other source) they received their water from. The results are summarized in his Table VII. He went on to say:

According to a return which was made to Parliament, the Southwark and Vauxhall Company supplied 40,046 houses from January 1st to December 31st, 1853, and the Lambeth Company supplied 26,107 houses during the same period; consequently, as **286 fatal attacks of cholera took place, in the first four weeks of the epidemic, in houses supplied by the former Company, and only 14 in houses supplied by the latter**, the proportion of fatal attacks to each 10,000 houses was as follows. Southwark and Vauxhall 71. Lambeth 5. **The cholera was therefore fourteen times as fatal at this period, amongst persons having the impure water of the Southwark and Vauxhall Company, as amongst those having the purer water from Thames Ditton.**

The additional tables extend the study to the end of the year, but he considers the data from the first 4 weeks as the most trustworthy, and also as the ones that gave the sharpest contrast (the least diluted by information quality, and other factors: indeed the mortality rate ratios declined as larger time windows are considered).

He returns to this data quality issue in his J Public Health, Oct. 1856 article. The main purpose of his article was to use newer and more fine-grained (district level) number-of-persons denominators than the overall number-of-houses denominators available to him when he completed the book at the end of 1854.

The main result was in Table V, which covered the entire 1854 epidemic,

TABLE V.
Showing the results of the Inquiry for the whole Epidemic of 1854.

Registration Districts.	Number of inhabited houses in 1851.	Population in 1851.	* Number of houses, and estimated number of persons, supplied in 1854 with water "as under."		Water supply of the houses in which fatal attacks of cholera took place.			Mortality per 10,000 supplied with water as under.			IN THE NORTH DISTRICTS OF LONDON.			
			By the Southwark and Vauxhall Co.		By the Lambeth Company.		Southwark and Vauxhall Co.	Lambeth Co.	Pump-works and other sources.	Supply not stated.		Deaths from cholera in the epidemic of 1854.		
			No. of houses.	Estimated population.	No. of houses.	Estimated population.								
St. Saviour, Southwark	4,600	35,731	78	3,631	19,617	1,689	14,201	406	72	10	3	491	207	90
St. Olave, Southwark	2,360	19,373	82	2,193	18,698	0	1,785	821	0	8	28	313	148	..
Bermondsey	7,007	48,128	69	8,492	57,884	268	1,785	821	0	25	0	846	142	..
St. George, Southwark	6,992	51,824	74	8,419	25,039	3,183	23,712	388	99	0	56	543	155	41
Newington	10,458	64,810	62	5,224	31,949	5,473	33,551	408	58	2	176	694	143	17
Lambeth	30,447	139,325	68	8,077	64,982	11,763	83,786	525	138	24	340	927	96	16
Wandsworth	8,376	50,764	61	3,928	18,290	618	3,870	298	7	106	40	421	145	18
Camberwell	9,412	54,607	58	4,905	29,472	1,835	10,478	322	33	115	49	549	150	31
Tothornhill	4,792	17,205	64	4,336	14,951	0	0	307	0	46	30	283	198	..
Greenwich & sub-dis. Sydenham	4	4	2	1	11
Houses not identified	66	411	2,712	..	165
Totals	72,844	482,435	67	39,730	207,025	34,854	171,528	3,706	411	338	623	5,078	138	23
Non-ascertained cases distributed in proportion of others	561	62
Population (Registrar-General)	286,516	..	173,748	4,267	473	338	..	5,078	160	27

59

In his discussion, he shows a keen **insight** – It is just as relevant today, in the era of ‘Big Data’ – concerning the **consequences of errors in data**, in this case the **addresses of the persons in the Registrar General’s list of deaths from cholera**.²⁵

[Big Data] **‘can bear no comparison in point of accuracy to a personal inquiry, made on the spot, at the time of the epidemic.’**

It concerns the quality of the **numerators**, ie. how the cholera deaths might have been (mis)classified into the (wrong) water companies, especially as the addresses were collected by people who might have realized how critical they would be in the cholera story.

As we recount in this recent item “The 25-page appendix to his 1855 book provided a detailed record of the 334 deaths from cholera that occurred in South London between 8 July and 5 August 1854. According to Snow, *this information was included ‘as a guarantee that the water supply was inquired into, and to afford any person who wishes it an opportunity of verifying the results* (p. 80)’. 6 The information that John Snow recorded included the address at which each cholera-related death occurred, the date of death, the occupation and the age of the deceased, the duration of symptoms before death and the water source (Figure 1).”

In that item, we recount another little-known fact: when the members of the house could not supply evidence as to which was the water supplier, he used a ‘high-tech’ method to determine it (1st column, p. 1795).

But, even though it is difficult to imagine that this would be a big issue today, his 1856 article is **quite concerned with the numbers of wrong addresses**.

²⁵Samantha.

the spot, at the time of the epidemic. In the first place, throughout the greater part of Lambeth, Newington, and the Borough, the houses are either without numbers, or numbered very irregularly, and the numbers are liable to frequent change, as new houses are built, or older ones re-painted; there are also frequently repetitions of the same number in the same street, and although, in some instances, the companies have returned the names of the occupiers, that can be of no assistance in the case of the poor, who occupy but one or two rooms, and form the greater bulk of the population. In the next place, the poor often furnish, unintentionally, a wrong number to the registrar, even when the houses are regularly numbered. They know their own homes perfectly, but, having no occasion to refer to the number, they partially forget it; and, in the greater number of my personal inquiries, I had to call at two or three houses before I found the one in which the death occurred.

and with the **consequences for the rate difference and the rate ratio:**

For these reasons it follows that, in comparing the lists of the water supply with the lists of deaths, many errors must have occurred; and as the deaths were six times as numerous in the houses supplied by the Southwark and Vauxhall Company as in those supplied by the Lambeth Company, the evident result would be that out of every six mistakes five would transfer a death from the former company to the latter, and only one would transfer a death from the latter company to the former. Another source of error, but operating

Exercise for bios601

- (a) Assume the numbers of deaths (4,267 and 473) in the columns of Table V are *correct*. Let P be the probability that the address of a diseased person is recorded correctly. Let it vary from perfect (1), to ‘almost’ (say 0.99), to 0.9 to 0.4. Assume also that the ‘mixing’ is so intimate that there is no ‘clustering’ of water supplier by street. Snow tells us that it was close to random:

In the sub-districts enumerated in the above table as being supplied by both Companies, the mixing of the supply is of the most intimate kind.

The pipes of each Company go down all the streets, and into nearly all the courts and alleys. A few houses are supplied by one Company and a few by the other, according to the decision of the owner or occupier at that time when the Water Companies were in active competition. *In many cases a single house has a supply different from that on either side.* Each company supplies both rich and poor, both large houses and small; there is no difference either in the condition or occupation of the persons receiving the water of the different Companies.

The experiment, too, was on the grandest scale. No fewer than three hundred thousand people of both sexes, of every age and occupation, and of every rank and station, from gentlefolks down to the very poor, were divided into two groups without their choice, and, in most cases, without their knowledge; one group being supplied with water containing the sewage of London, and, amongst it, whatever might have come from the cholera patients, the other group having water quite free from such impurity.

To turn this grand experiment to account, all that was required was to learn the supply of water to each individual house where a fatal attack of cholera might occur. I regret that, in the short days at the latter part of last year, I could not spare the time to make the inquiry; and, indeed, I was not fully aware, at that time, of the very intimate mixture of the supply of the two Water Companies, and the consequently important nature of the desired inquiry.

Show what the ratio of the expected mortality rates would be if the quality of the address reporting was as low as Snow suggested. Since it is difficult to convert ‘*in the greater number of my personal inquiries, I had to call at two or three houses before I found the one in which the death occurred*’ into an exact probability, compute and plot/tabulate the ratio against the $P = c(1, 0.99, (9:4)/10)$ values suggested above. Some might call this a *sensitivity analysis*.

- (b) Assume the split of the numbers of deaths is *incorrect*, i.e., that they are *already distorted* by the errors in the addresses. Thus, your job is to reverse-engineer what the *correct* split of the 4740 is. Repeat part (a) but with the ‘truth’ starting at more extreme splits of the 4740 deaths than the 4,267:473 observed. If (say) $P=0.9$, what would the true split (and thus the mortality ratio) have to be to produce the ‘observed’ ratio?
- (c) What principles can you draw from this assignment, and how well does the lesson fit with the ‘regression dilution’ examples earlier?
- (d) Are these errors in the addresses ‘*differential*’ or ‘*non-differential*’? Be careful with online definitions: the issues are subtle.

Snow mentions one other source of error – one he himself did not make in his ‘shoe-leather’ epidemiology. He made sure to enquire about the water supply in the house the person lived in *before* getting sick.

pany to the former. Another source of error, but operating to a less extent, is, that a number of persons who were attacked with cholera in houses supplied by the Southwark Company died in the workhouses of St. Saviour's, Lambeth, and Newington, which were supplied by the Lambeth Company. It need excite no surprise, therefore, that the supplemental inquiry, embodied in the recent Report, instead of showing a mortality of 160 and 27 for the population supplied by the two water companies, or a difference of 6 to 1, showed a mortality of 125 and 37 per 10,000, or a difference of only $3\frac{1}{2}$ to 1. It must be obvious, however, independently of the above facts, that a difference of three and a-half to one would not explain the great difference in the mortality of the various districts and subdistricts. The epidemic of 1853 is included with that of 1854 in Mr. Simon's Report; but as there were but few deaths in 1853, and those chiefly amongst the population supplied by the Southwark Company, this circumstance would not much affect his results.

24. A more modern (but ? similar type of) example:

Suppose you were asked to calculate the difference in the mean ages of death of the males and females in this dataset of 1,000 persons who died of Covid-19.

The 1000 ages were easy to extract, and had [this distribution](#). But the extraction of the names proved more difficult, and then (electronically – or even manually) converting them to male and female posed added some uncertainty. [This link](#) led JH to the `gender` package for R. Based just on the first name, it can produce a probability of being male for each person listed (of course, one could modify/override these based on any additional information contained in each ‘blurb’).

How would you suggest these probabilities, p_1 to p_{1000} , (some of which are near/at zero and 1, and some are intermediate) be used to calculate a point and *interval* estimate of the difference in mean age at death?

25. Mixing of the ‘real’ and the ‘noise’

Refer to Fig 12.5 in the ‘online book’, section 12.7.2 (Measurement error), [link here](#) for a colour-based depiction of the mixing of ‘true’ and ‘error’ distributions.

There, E has a ‘2-point distribution, namely -0.5 and +0.5, with equal probabilities.’

Extend the diagram, and the VAR and ICC calculations, for the following E distributions:

- 3-point distribution, namely -0.5, 0, and +0.5, with probabilities $1/4, 1/2, 1/4$.
- 5-point distribution, namely -1, -0.5, 0, +0.5 and +1, with probabilities $1/10, 2/10, 4/10, 2/10$ and $1/10$.

26. ‘Berkson’ error model

In your own words, explain why Berkson error in X does not flatten the regression slope. Also, show it by algebra.

27. Effect of ‘Coarsening’ of Age to ‘not publish personally identifiable information’

The following notes accompanied the dataset that authors of the article “Increased mortality in community-tested cases of SARS-CoV-2 lineage B.1.1.7” (now known as the ‘alpha’ variant, rather than the ‘UK’ variant) published in *Nature* on March 15, 2021, shared with the public.

“This repo includes an anonymised data set (in `./dataset/`) which allows partial replication of the analyses in our preprint. This is a `qs` file, which can be read using the ‘`qs`’ R package.

Note that the **anonymised data set randomises ages within 5-year age bands**, and caps the maximum age at 104 for purposes of not publishing personally identifiable information. For this reason, results obtained using this data set will not be exactly equal to those found in the paper, which analyses the non-public data. (This just means we have the exact age in years for each person tested—we have never had access to things like names and addresses!)

The `FINALID` column in the anonymised data set is also randomised, so please be aware that this will *not* link correctly to other datasets held by Public Health England. This is again in order to minimise any risk of disclosure of personal information.”

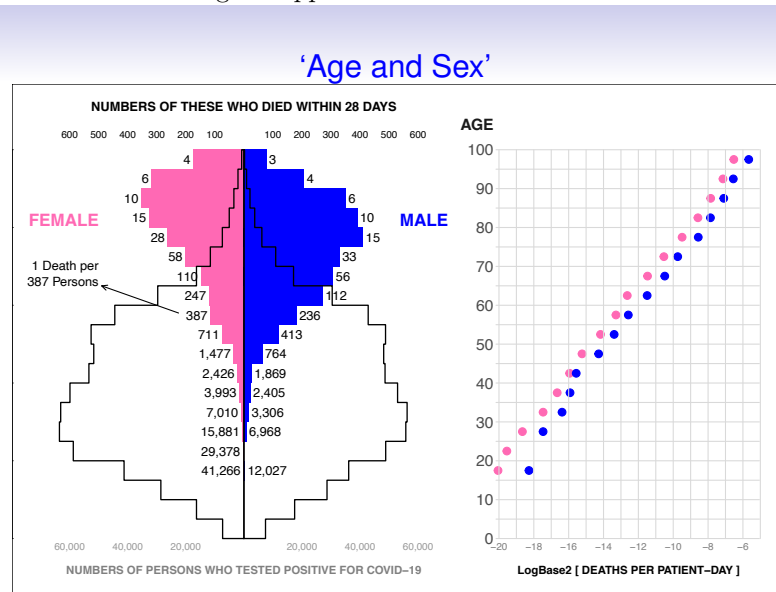
This is the R code they used to coarsen age

```
# Coarsen age
rd = rd[age != 0]; # remove age 0
rd[, age := (pmin(100, age) %/% 5) * 5 +
          sample(0:4, .N, replace = TRUE)];
```

Suppose the distribution of the exact age in years of a million cases has a (??-like) shape like the one produced by this R code

```
age = 1:110
f = 1*age*(age <= 25) +
  (25 - 0.1 *(age-25))*(age > 25 & age <=65) +
  (21.5- 0.7*(age-65))*(age > 65 & age <=80)+
  (11- 0.365 *(age-80))*(age > 80 & age <=110)
f=1000000*f/sum(f) ;
plot(cbind(f/2,age), xlim=c(-max(f/2),max(f/2)),
     pch=19, cex=0.25 )
points(cbind(-f/2,age),pch=19, cex=0.25 )
```

This is a ‘rough’ approximation to the actual distribution,



since it ignores the sex differences.

Suppose also, still ignoring the sex differences, that the age-specific 28-day mortality in the non-public data can be (very roughly!) modelled

as the ‘Gompertz-like’²⁶ function

$$\% \text{ among persons age } a = 100 \times 2^{-22 + 0.2 \times a},$$

so that the log of the proportion is linear in age, and has a slope of 0.2 if we put the in the \log_2 scale. i.e., the proportion or % among those age $a + 5$ is double what it is among those age a .

Exercise:

- (a) Determine, by algebra, or by numerical investigation, what, if any, is the effect of the coarsening on the ‘slope’ of 0.2? (If you prefer a simpler context, just imagine a simple linear regression, where

$$\text{Average blood pressure in persons age } a = 120 + 0.5 \times a,$$

and where there are equal numbers of persons at each age a .)

- (b) What if these UK authors had randomised ages within 10-year rather than 5-year age bands? Do your calculations suggest that the error model is closer to the ‘classical’ or the ‘Berkson’ version?

28. **SGTF is an imperfect test for the presence of B.1.1.7 [formerly known as the ‘UK’ ‘variant of concern’ (VOC)]**

“The presence or absence of B.1.1.7 202012/01] can be identified because mutations in this lineage prevent PCR amplification of the spike (S) gene target (known as S gene target failure (SGTF)).

Before the emergence of B.1.1.7, a number of minor circulating SARS-CoV-2 lineages with mutations in the S gene could also cause SGTF.

Accordingly, a positive test with SGTF is *not necessarily* indicative of infection with B.1.1.7. A negative test of SGTF is assumed to be indicative of an absence of infection with B.1.1.7. Thus, our main analyses are restricted to specimens from 1 November 2020 onwards to avoid diluting the measured effect of B.1.1.7 on mortality due to non-B.1.1.7 lineages that cause SGTF. [Misclassification of an exposure can result in bias in its estimated association with the outcome.]”

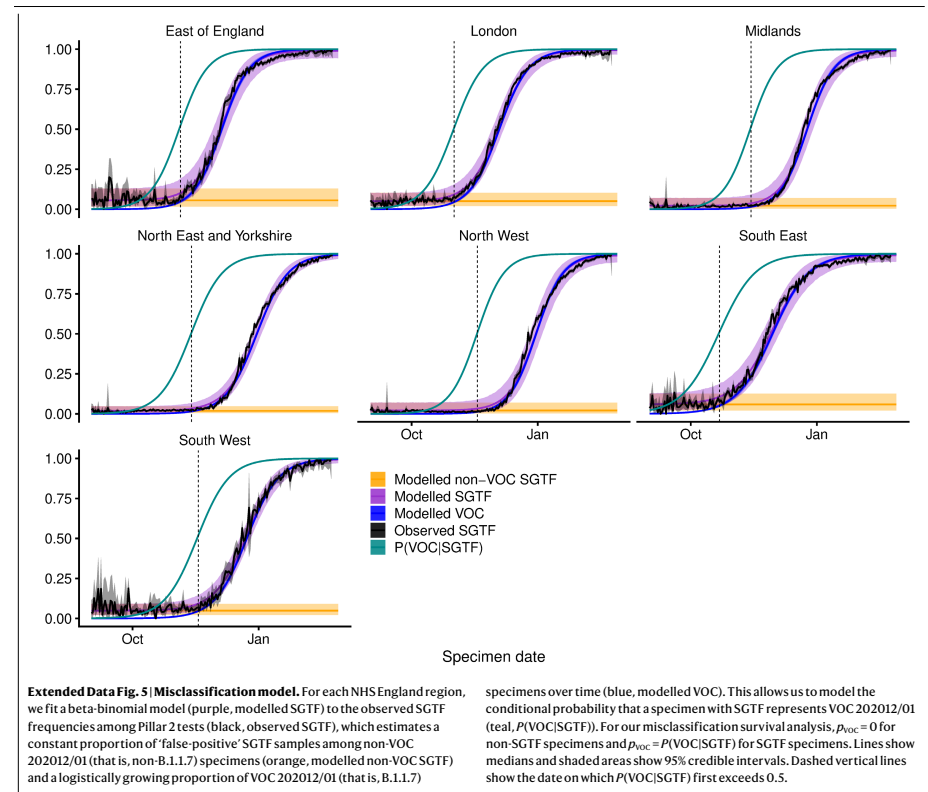
The green curves in Extended Data Fig. 5 show how the (modelled) Prob[VOC+ | SGTF+] evolved over time.

²⁶[Link](#)

To simplify matters, assume that in the period from 1 November onwards, the ‘average’ $\text{Prob}[\text{VOC} \mid \text{SGTF}^+]$ was 0.90, that there were 500,000 SGTF⁺ and 400,000 SGTF⁻ cases, and that there were 4,000 and 2,000 deaths among these, respectively.

Exercise:

- Calculate the naive case fatality rate²⁷ ratio for the VOC⁺ : VOC-contrast, taking an SGTF⁺ test as equivalent to VOC⁺.
- Calculate a corrected case fatality rate ratio for the VOC⁺ : VOC-contrast, i.e., allowing for the false positive tests.
- Do your results fit with those in the article? [See second last sentence of the Abstract, as well as the ‘Misclassification analysis’ on page 4, and the ‘Misclassification analysis’ in the second (un-numbered) page of the Methods section of the article.]
- Is JHs use of the term ‘CFR’ closer to the traditional CFR or to an $\underline{\text{IFR}}$? (See the Nature article, and the authors’ use of the term CFR in their Table 1)



²⁷In early 2020, some politicians downplayed the case fatality rate, and wanted to include the ‘silent’ cases in the denominator in order to make the CFR lower. The CFR is a very old epidemiological term, and typically referred only to the rate in **recognized** cases of the illness/condition. Since 2020, one often sees the term *Infection* fatality rate, based on a denominator of **all infected individuals, including all asymptomatic and undiagnosed persons**. For a well thought out ‘dictionary’ of epidemiology, see the 1-author book *Epidemiological research : terms and concepts* by the late McGill emeritus professor Olli S Miettinen, available as an ebook from the [McGill library](#). Here is the CFR entry in it:

“Case-fatality rate * (synonyms: fatality rate, death rate) – Concerning cases of an illness in general, or recognized cases of it (ones with rule-in diagnosis about the illness), the proportion in which the illness is fatal; that is, such that the outcome of the course of the illness is fatality from it. (Cf. ‘Survival rate.’)

Note: For the concept to be truly meaningful, it commonly is to be specific to particulars of the case (broadly at least) and to the choice of treatment; and it also is to be conditional on absence of intercurrent death from some other, ‘competing’ cause.”

29. Readability (initially addressed in Q1 and Q2, dating from the pre-computer era) revisited

Refer to the article

‘The Past, Problems, and Potential of Readability Analysis’

by N. Lines, in *Chance Magazine* in 2022.

- In 150 words, summarize the article.
- Of the ‘instruments/tools’ mentioned, indicate the ones that are (immediately) accessible to you.
- Try a few sub-samples of text (from the same text) in one/more of them, and tell us what you found.
- Do you like the layout of Figure 1? Can you think of other ways to display the information?

- (e) Look up the ‘SMOG’ index, mentioned in Q1, and see if is different from all of the tools listed in Table 1.
- (f) *FYI only, here is the article that introduced ‘Cloze’ Tests. JH very much liked the old-style way the article was written.*
- (g) Comment on the findings shown in Table 2. In particular, explain the opposite signs, and interpret the low correlations.
- (h) Did you find the article interesting and useful? Should JH include this Q29 in future exercises?

31. What was the point of each of the assignments?

For each of the assigned questions, use one sentence to describe what you think the learning objective was; use another to describe in what situations the concepts and techniques will be of use to you and to those you will work with.

30. A statistical query from a vaccine researcher I know

The following is from an email I got in June 2023.

I have a simple problem and I hope you will help interpret my solution.

We are running a virus neutralization assay, and the scientist performing the task takes each serum sample and runs it in triplicate. This wouldn’t be a problem, except that we have to run the test against many virus strains, and we have a limited amount of serum.

I have tried to convince the scientist that there is no need to run the samples in triplicate because the main source of variability is in the animal subjects, not the assay. Also, more animals can be tested on the same day if only 1 repeat is used.

I would like to prove this. Here is what I propose: I ran a two-way ANOVA and got the following results.

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
group	5	23.24	4.648	9.149	4.89e-07 ***
subject	1	0.80	0.802	1.578	0.212
sample	1	0.00	0.002	0.003	0.955
Residuals	88	44.71	0.508		

group is the vaccine

subject is the animal number

sample is the replicates

(in the data file, the assay result is labelled ‘A’)

How do I interpret the results? Does this show the replicates (sample) have a much smaller contribution to the error than the subjects?

Exercise:

After you have re-analyzed the data, draft a reply to the query.

Endpieces

• Type IV error

The following Table contains the Result of the Experiments.

Exper.	Mot. weight	Mot. arm	Do. corr.	Time vib.	Do. corr.	Density.
1	m. to +	14.32	13.42	" "	-	5.5
	+ to m.	14.1	13.17	14.55	-	5.61
2	m. to +	15.87	14.69	-	-	4.88
	+ to m.	15.45	14.14	14.42	-	5.07
3	+ to m.	15.22	13.56	14.39	-	5.26
	m. to +	14.5	13.28	14.54	-	5.55
4	m. to +	3.1	2.95	-	6.54	5.36
	+ to -	6.18	-	7.1	-	5.29
	- to +	5.92	-	7.3	-	5.58
5	+ to -	5.9	-	7.5	-	5.65
	- to +	5.98	-	7.5	-	5.57
6	m. to -	3.03	2.9	-	-	5.53
	- to +	5.9	5.71	-	-	5.62
7	m. to -	3.15	3.03	7.4	6.57	5.29
	- to +	6.1	5.9	by mean.		5.44
8	m. to -	3.13	3.00	-	-	5.34
	- to +	5.72	5.54	-	-	5.79
9	+ to -	6.32	-	6.58	-	5.1
10	+ to -	6.15	-	6.59	-	5.27
11	+ to -	6.07	-	7.1	-	5.39
12	- to +	6.09	-	7.3	-	5.42
	- to +	6.12	-	7.6	-	5.47
13	+ to -	5.97	-	7.7	-	5.63
	- to +	6.27	-	7.6	-	5.34
14	+ to -	6.13	-	7.6	-	5.46
	- to +	6.34	-	7.7	-	5.3
15	- to +	6.34	-	7.7	-	5.3
16	- to +	6.1	-	7.16	-	5.75
	- to +	5.78	-	7.2	-	5.68
17	+ to -	5.64	-	7.3	-	5.85
	+ to -	5.64	-	7.3	-	5.85

XXI. Experiments to determine the Density of the Earth. By Henry Cavendish, Esq. F.R.S. and A.S.

Read June 21, 1798.

MANY years ago, the late Rev. JOHN MICHELL, of this Society, contrived a method of determining the density of the earth, by rendering sensible the attraction of small quantities of matter; but, as he was engaged in other pursuits, he did not complete the apparatus till a short time before his death, and did not live to make any experiments with it. After his death, the apparatus came to the Rev. FRANCIS JOHN HYDE WOLLASTON, Jacksonian Professor at Cambridge, who, not having conveniences for making experiments with it, in the manner he could wish, was so good as to give it to me.

http://en.wikipedia.org/wiki/Cavendish_experiment: in 1798 Cavendish found that the Earth's density was 5.448 ± 0.033 times that of water (due to a **simple arithmetic error, found in 1821**, the erroneous value 5.48 ± 0.038 appears in his paper).

• Scientific Method, Statistical Method and the Speed of Light

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• Determinations of the parallax of the sun, the mean density of the earth, and the speed of light

----- [Link](#)

• A Historical View of Statistical Concepts in Psychology and Educational Research

----- [Stigler](#) — [Edgeworth \(cited\)](#) — [Peirce \(cited\)](#)