Social Science Inquiry II

Week 5: Uncertainty and inference, part II

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Loading packages for this class

- > library(ggplot2)
- > set.seed(60637)

Continuing inference

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- ▶ Last class, we assumed we had a finite population that we observe all of, and the source of randomness in what we observed was due to random assignment of treatment. This is called *randomization inference*.
- Now, we'll assume that our data is produced from a random generative process, where we're sampling from some (potentially infinite) population distribution that is not fully observed. This is the type of inference we use for survey sampling.
- ▶ It's important to consider what the source of randomness is and what population we're making inferences about.

▶ Returning to our example where we flip a coin twice, let *X* be the number of heads we observe. Our coin is *not* fair, and the probability of getting a heads is 0.8.

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- ► The random variable's probability distribution is then:

$$f(x) = \begin{cases} 1/16 & x = 0\\ 3/8 & x = 1\\ 9/16 & x = 2\\ 0 & \text{otherwise} \end{cases}$$

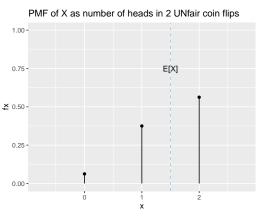
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- ► For example, with respect to the Pager (2003) data, we can use a process like this to model the probability that an employer will hire a white applicant without a criminal record.

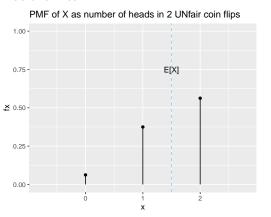
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- ► For example, with respect to the Pager (2003) data, we can use a process like this to model the probability that an employer will hire a white applicant without a criminal record.
- ▶ We might say that there are different random processes, with different probabilities of success, for whites with and without criminal records, and blacks with and without criminal records.

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- ▶ We might say that there are different random processes, with different probabilities of success, for whites with and without criminal records, and blacks with and without criminal records.
- ► Here, where we have mutliple coin flips, we can compare that to the probability distribution of hires for two people with the same profile.

Let's take a look at the mean.



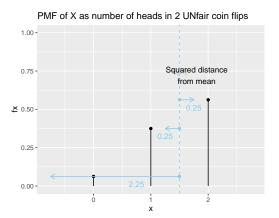
Let's take a look at the mean.



$$E[X] = \sum_{x} xfx$$

$$= 0 \times \frac{1}{16} + 1 \times \frac{3}{8} + 2 \times \frac{9}{16} = \frac{24}{16}$$

And the spread.



Variance = average squared distance from the mean

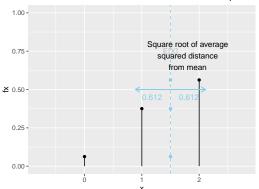
$$Var[X] = E[(X - E[X])^{2}]$$

$$= 2.25 \times \frac{1}{16} + 0.25 \times \frac{3}{8} + 0.25 \times \frac{9}{16}$$

$$= 0.375$$

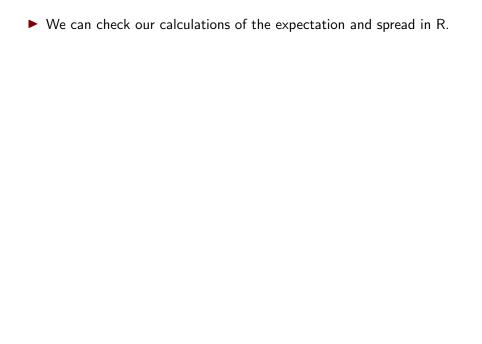
And the spread.

PMF of X as number of heads in 2 UNfair coin flips



SD = square root of variance

$$=\sqrt{0.375}=0.612$$



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```
> n <- 1000
> X <- c(0, 1, 2)
> probs <- c(1/16, 3/8, 9/16)
> x_observed <- sample(X, prob = probs,
+ replace = TRUE,
+ size = n)
> head(x_observed)
[1] 1 0 1 1 2 2
> mean(x_observed)
```

- [1] 1.514
- > var(x_observed)
- [1] 0.3661702
- > sd(x_observed)
- [1] 0.60512

► The process that we just did – sampling and estimation based on observed data – is a very common process in empirical research.

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- ▶ But we may notice that the mean, variance, and standard deviation are not exactly what we calculated analytically.

Let's try it again.

```
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> x_observed <- sample(X,</pre>
                         prob = probs,
                         replace = TRUE,
                          size = n)
> mean(x_observed)
[1] 1.496
> var(x_observed)
[1] 0.3823664
> sd(x_observed)
```

[1] 0.6183578

► The values that we get are close, but not identical.

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- ► This is because what we are observing in practice is a *sample* from the data.

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- ► This is what we use statistics for, and why we talk about probability AND statistics.
- ▶ Probability gives us a model of the world.
- ▶ Statistics give us a way to relate the data that we see to the model.

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- Formally, that educated guess is called *estimation*.

Let's repeat our random sampling from the double coin flip, but we'll consider a smaller sample, of size n = 100.

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> mean(x_observed)

[1] 1.44

▶ We differentiate the *sample mean* from the *population mean* because the sample mean will vary with every new sample we draw.

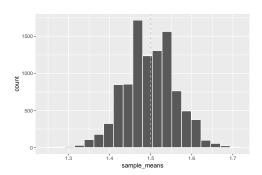
- ▶ We differentiate the sample mean from the population mean because the sample mean will vary with every new sample we draw.
- We'll use a simulation with replicate() to see what would happen if we took a sample of size n=100 from the population distribution many times.

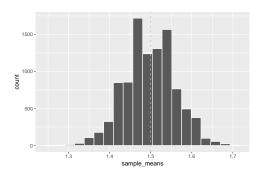
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```
> n iter <- 10000
> x_mat <- replicate(n_iter, sample(X,
                                    prob = probs,
+
+
                                    replace = TRUE,
                                    size = n)
> dim(x mat)
[1]
      100 10000
> head(x_mat[.1])
[1] 0 2 2 2 2 2
> head(x_mat[.2])
[1] 2 2 1 2 1 0
```

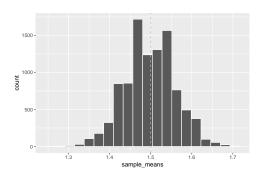
```
> sample_means <- apply(x_mat, 2, mean)
> length(sample_means)
[1] 10000
> head(sample_means)
```

[1] 1.59 1.51 1.54 1.47 1.43 1.46





We see the sample means are roughly distributed around the mean of the underlying population, Ex.



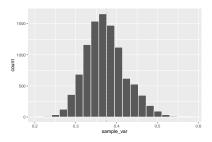
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The expected value of the sample mean is the population mean.

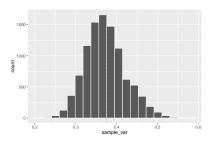
We can estimate the mean of the population using the sample mean. What about the sample variance?

We'll do the same process with our simulations.

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We see the sample variances are roughly distributed around the variance of the underlying population, sdx^2 .

The formula for the unbiased sample variance is:

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Why do we divide by n-1, instead of n?

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- ▶ However, because it is made up of the $1, ..., n X_i$ that we actually observe, the expected difference between $(X_i \bar{X}_n)$ is a little bit smaller than the expected difference between $(X_i E[X])$.

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- ▶ However, because it is made up of the $1, ..., n X_i$ that we actually observe, the expected difference between $(X_i \bar{X}_n)$ is a little bit smaller than the expected difference between $(X_i E[X])$.
- ▶ To account for this, we divide by n-1, instead of n.

> head(x_observed)

[1] 1 1 1 2 2 1

- > head(x_observed)
- [1] 1 1 1 2 2 1
- > var(x_observed)
- [1] 0.329697

- > head(x_observed)
- [1] 1 1 1 2 2 1
- > var(x_observed)
- [1] 0.329697
- > sum((x_observed mean(x_observed))^2)/(n-1)
- [1] 0.329697
- >

```
> head(x_observed)
```

>

R uses the formula for the unbiased sample variance.

► The sample mean is itself a random variable, and so it has its own mean and variance. The mean of the sample mean is the population mean. The variance of the sample mean is:

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▶ And the standard deviation of the sample mean is:

$$\sqrt{\mathrm{Var}[\bar{X}_n]} = \sqrt{\frac{\mathrm{Var}[X]}{n}}$$

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 - ► The *standard error* describes the sampling variation of an **estimator**; i.e., how much our estimates will vary based on the random sample that we draw.
 - standard deviation describes the underlying variation in the population distribution.

Let's check this in our simulation. We saw that mathematically, Var[X] was 0.375. So

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- ▶ In fact, from our 10000 separate samples, we calculate 10000 separate sample means. From the variation in these sample means, we again *estimate* the variance of the sample mean.
- ▶ But *this estimate is itself a random variable*, with, again, its own sampling distribution. We will get slightly different estimates of the sampling variance of the sample mean each time we take our 10000 separate samples.
- ► In practice, we will estimate the standard error of the sample mean by plugging our unbiased sample variance formula into the standard error formula:

$$\hat{\rm se} = \sqrt{S_n^2/n}$$

References I

Pager, D. (2003). The mark of a criminal record. <u>American journal of sociology</u>, 108(5):937–975.