

19: Sampling and the Bootstrap

Jerry Cain

May 10, 2021

Quick slide reference

3	Sampling definitions	19a_intro
11	Unbiased estimators	19b_sample_stats
23	Reporting estimation error	19c_statistical_error
29	Bootstrap: Sample mean	19d_bootstrap_mean
40	Bootstrap: Sample variance	LIVE
*	Bootstrap: Hypothesis testing	LIVE

Sampling definitions

Motivating example

You want to know the true mean and variance of happiness in Bhutan.

- But you can't ask everyone.
- You poll 200 random people.
- Your data looks like this:

Happiness = {72, 85, 79, 91, 68, ..., 71}

- The mean of all these numbers is 83.

Is this the **true mean happiness** of Bhutanese people?



Population



This is a **population**.

Sample



A **sample** is selected from a population.

Sample



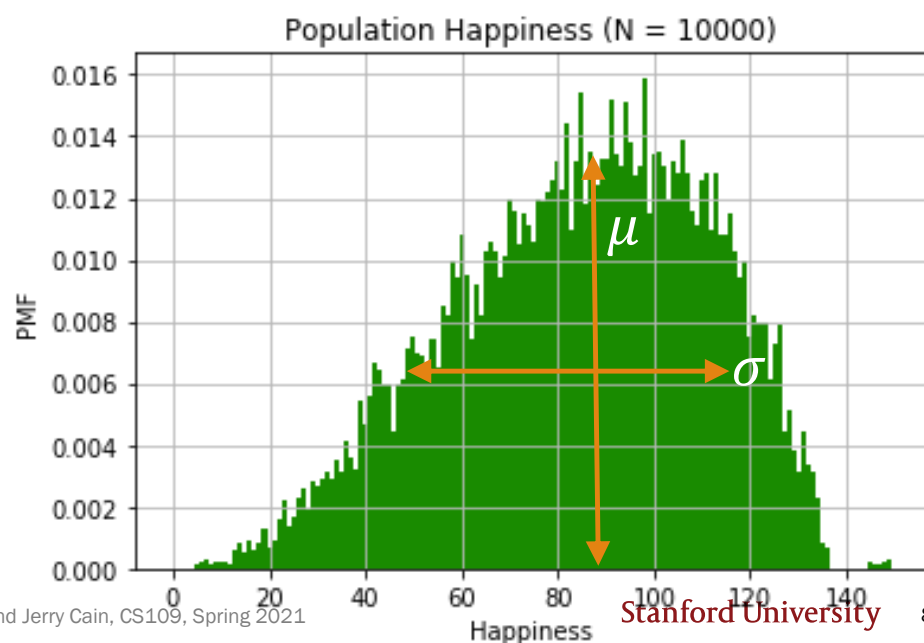
A **sample** is selected from a population.

A sample, mathematically

Consider n random variables X_1, X_2, \dots, X_n .

The sequence X_1, X_2, \dots, X_n is a **sample** from distribution F if:

- X_i are all independent and identically distributed (i.i.d.)
- X_i all have same distribution function F (the **underlying distribution**), where $E[X_i] = \mu$, $\text{Var}(X_i) = \sigma^2$



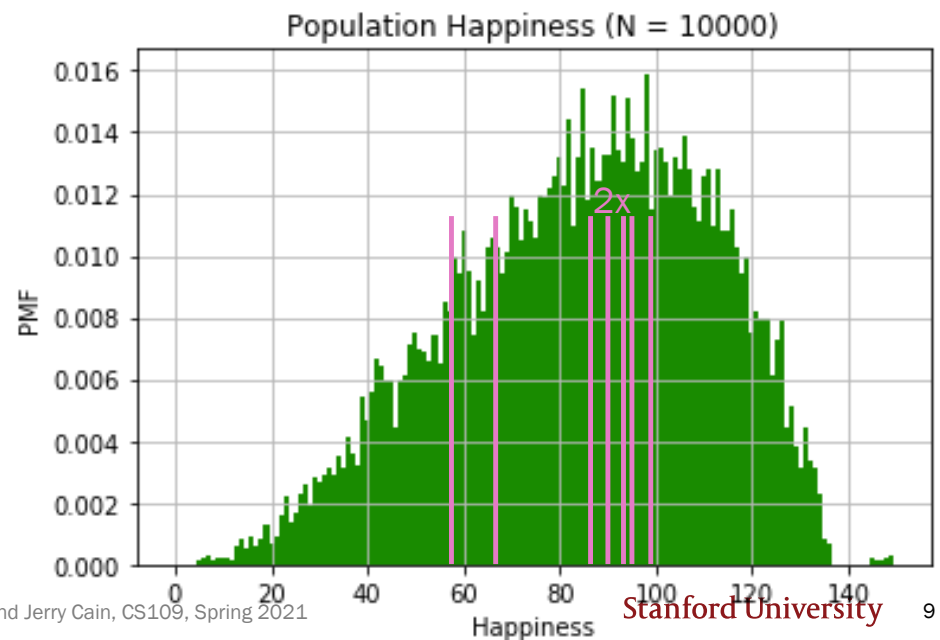
A sample, mathematically

A sample of **sample size** 8:

$(X_1, X_2, X_3, X_4, X_5, X_6, X_7, X_8)$

A **realization** of a sample of size 8:

$(59, 87, 94, 99, 87, 78, 69, 91)$



A single sample



A happy
Bhutanese person

If we had a distribution F of our entire population, we could compute exact statistics about about happiness.

But we only have 200 people (a sample).

Today: If we only have a single sample,

- How do we report *estimated* statistics?
- How do we report estimated error of these estimates?
- How do we perform hypothesis testing?

19b_sample_stats

Unbiased estimators

A single sample



A happy
Bhutanese person

If we had a distribution F of our entire population, we could compute exact statistics about about happiness.

But we only have 200 people (a sample).

So these population statistics are unknown:

- μ , the **population mean**
- σ^2 , the **population variance**

A single sample

If we had a distribution F of our entire population, we could compute exact statistics about about happiness.



A happy
Bhutanese person

But we only have 200 people (a sample).

- From these 200 people, what is our best estimate of **population mean** and **population variance**?
- How do we define best estimate?

Estimating the population mean



1. What is our best estimate of μ , the **mean happiness** of Bhutanese people?

If we only have a sample, (X_1, X_2, \dots, X_n) :

The best estimate of μ is the **sample mean**:

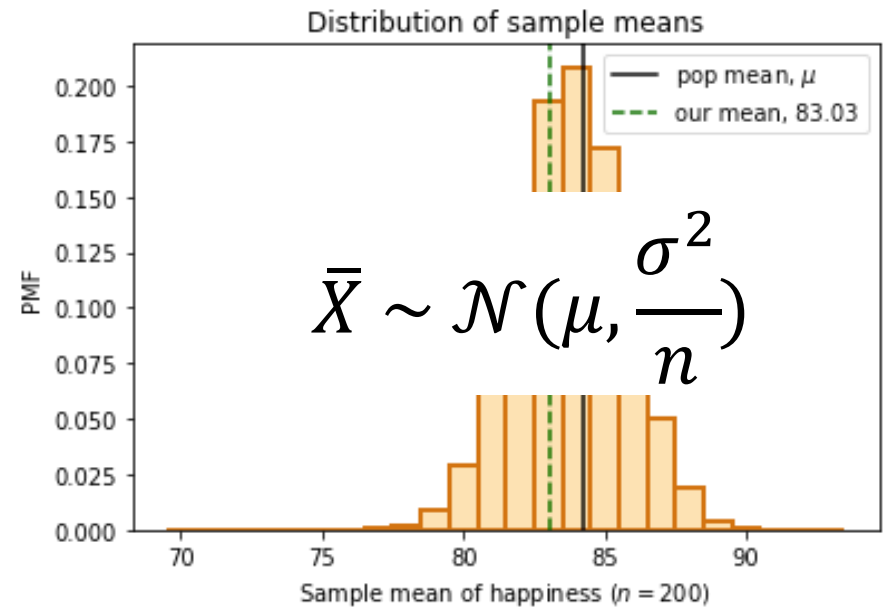
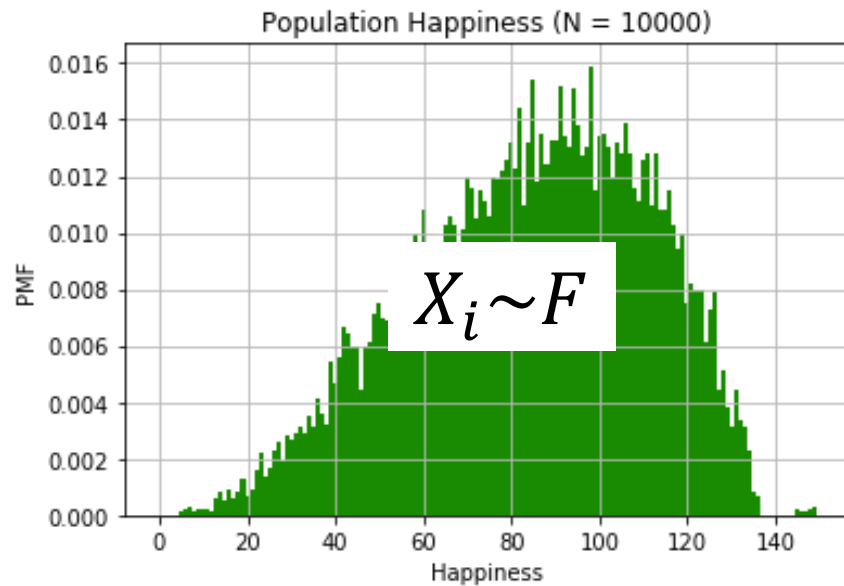
$$\bar{X} = \frac{1}{n} \sum_{i=1}^n X_i$$

\bar{X} is an unbiased estimator of the population mean μ . $E[\bar{X}] = \mu$

Intuition: By the CLT, $\bar{X} \sim \mathcal{N}(\mu, \frac{\sigma^2}{n})$ If we could take *multiple* samples of size n :

1. For each sample, compute sample mean
2. On average, we would get the population mean

Sample mean



Even if we can't report μ , we can report our sample mean 83.03, which is an unbiased estimate of μ .

Estimating the population variance



2. What is σ^2 , the **variance of happiness** of Bhutanese people?

If we knew the entire population (x_1, x_2, \dots, x_N) :

population variance

$$\sigma^2 = E[(X - \mu)^2] = \frac{1}{N} \sum_{i=1}^N (x_i - \mu)^2$$

population mean

If we only have a sample, (X_1, X_2, \dots, X_n) :

sample variance

$$S^2 = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X})^2$$

sample mean

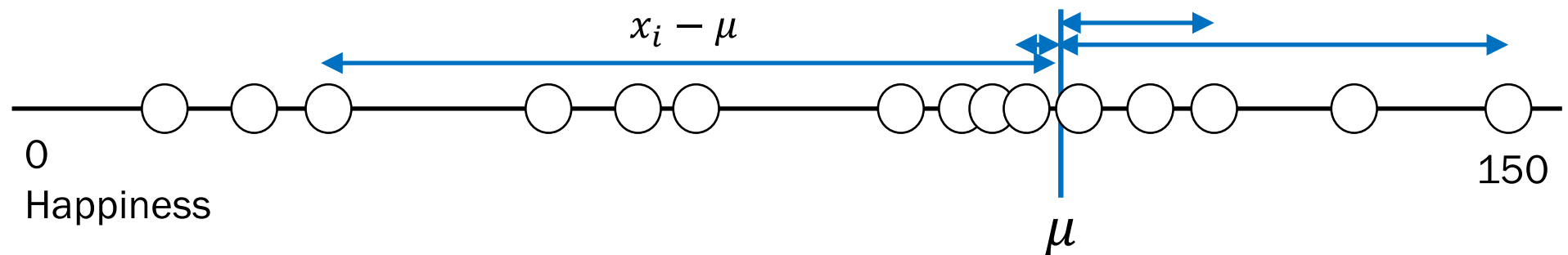
Intuition about the sample variance, S^2

Actual, σ^2

population variance

$$\sigma^2 = \frac{1}{N} \sum_{i=1}^N (x_i - \mu)^2$$

population mean



Population size, N

Calculating population statistics exactly requires us knowing all N datapoints.

Intuition about the sample variance, S^2

Actual, σ^2

population variance

$$\sigma^2 = \frac{1}{N} \sum_{i=1}^N (x_i - \mu)^2$$

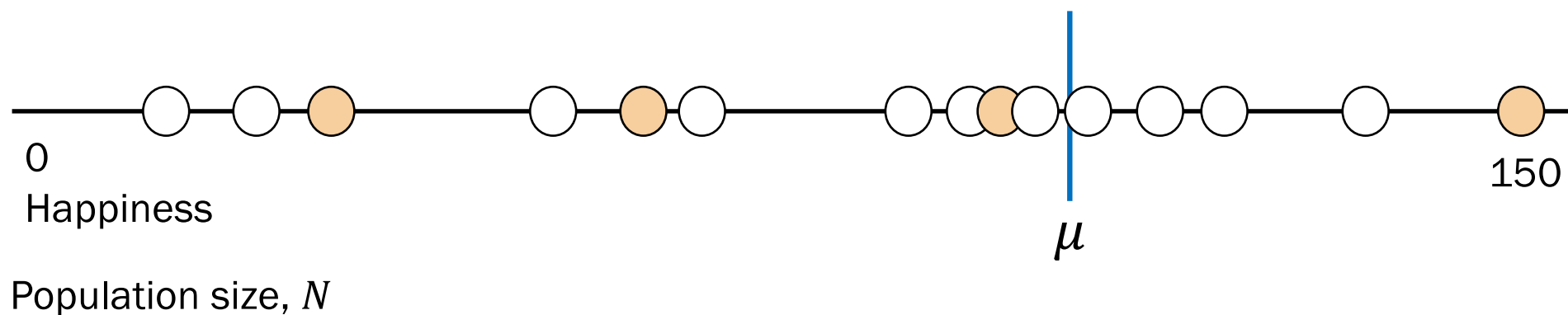
population mean

Estimate, S^2

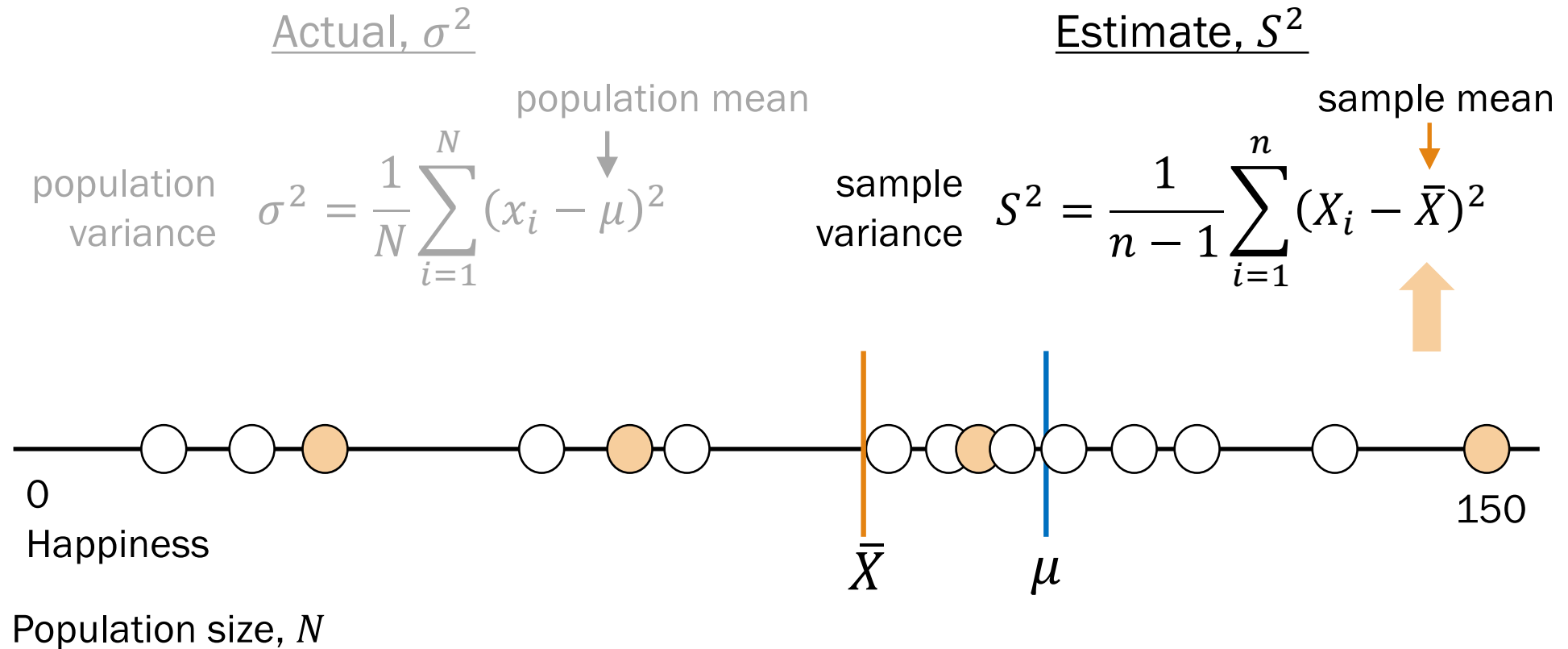
sample variance

$$S^2 = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X})^2$$

sample mean



Intuition about the sample variance, S^2



Intuition about the sample variance, S^2

Actual, σ^2

population variance

population mean

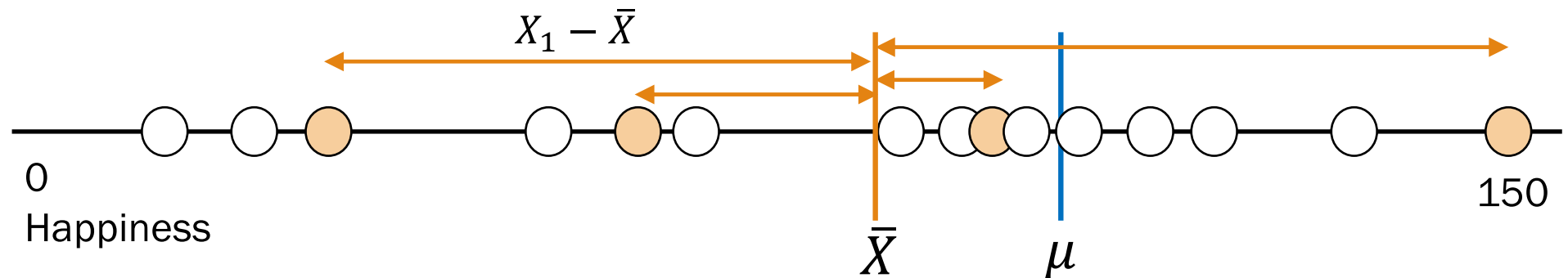
$$\sigma^2 = \frac{1}{N} \sum_{i=1}^N (x_i - \mu)^2$$

Estimate, S^2

sample variance

sample mean

$$S^2 = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X})^2$$



Population size, N

Sample variance is an estimate using an estimate, so it needs additional scaling.

Estimating the population variance



2. What is σ^2 , the **variance of happiness** of Bhutanese people?

If we only have a sample, (X_1, X_2, \dots, X_n) :

The best estimate of σ^2 is the **sample variance**:

$$S^2 = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X})^2$$

S^2 is an **unbiased estimator** of the population variance, σ^2 . $E[S^2] = \sigma^2$

Proof that S^2 is unbiased (just for reference)

$$E[S^2] = \sigma^2$$

$$E[S^2] = E\left[\frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X})^2\right] \Rightarrow (n-1)E[S^2] = E\left[\sum_{i=1}^n (X_i - \bar{X})^2\right]$$

$$(n-1)E[S^2] = E\left[\sum_{i=1}^n ((X_i - \mu) + (\mu - \bar{X}))^2\right]$$

(introduce $\mu - \mu$)

$$= E\left[\sum_{i=1}^n (X_i - \mu)^2 + \sum_{i=1}^n (\mu - \bar{X})^2 + 2 \sum_{i=1}^n (X_i - \mu)(\mu - \bar{X})\right]$$

$$= E\left[\sum_{i=1}^n (X_i - \mu)^2 + n(\mu - \bar{X})^2 - 2n(\mu - \bar{X})^2\right]$$

$$= E\left[\sum_{i=1}^n (X_i - \mu)^2 - n(\mu - \bar{X})^2\right] = \sum_{i=1}^n E[(X_i - \mu)^2] - nE[(\bar{X} - \mu)^2]$$

$$= n\sigma^2 - n\text{Var}(\bar{X}) = n\sigma^2 - n\frac{\sigma^2}{n} = n\sigma^2 - \sigma^2 = (n-1)\sigma^2$$

Therefore $E[S^2] = \sigma^2$

$$\begin{aligned} & 2(\mu - \bar{X}) \sum_{i=1}^n (X_i - \mu) \\ & 2(\mu - \bar{X}) \left(\sum_{i=1}^n X_i - n\mu \right) \\ & 2(\mu - \bar{X})n(\bar{X} - \mu) \\ & -2n(\mu - \bar{X})^2 \end{aligned}$$

19c_standard_error

Standard error

Estimating population statistics

A particular outcome

1. Collect a sample, X_1, X_2, \dots, X_n .

(72, 85, 79, 79, 91, 68, ..., 71)
 $n = 200$

2. Compute **sample mean**, $\bar{X} = \frac{1}{n} \sum_{i=1}^n X_i$.

$\bar{X} = 83$

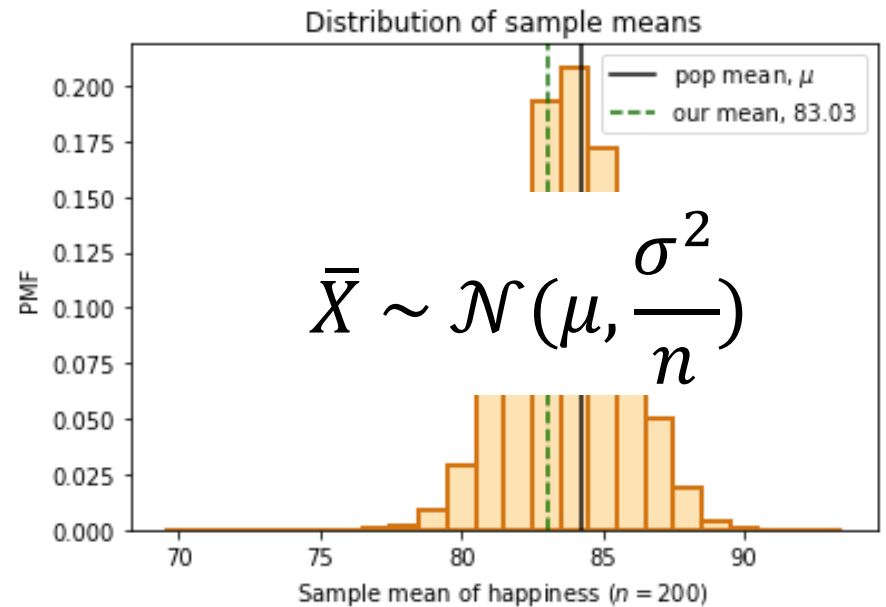
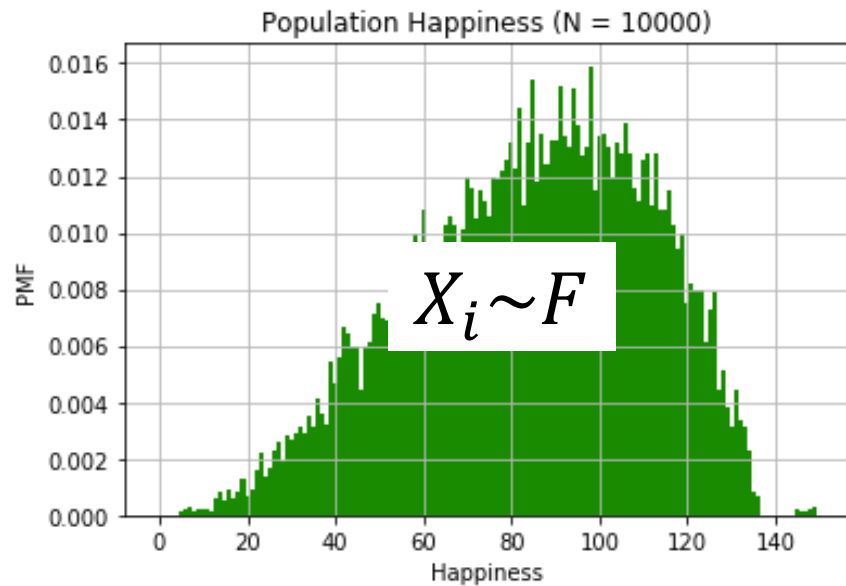
3. Compute sample deviation, $X_i - \bar{X}$. $(-11, 2, -4, -4, 8, -15, \dots, -12)$

4. Compute **sample variance**, $S^2 = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X})^2$.

$S^2 = 793$

How “close” are our estimates \bar{X} and S^2 ?

Sample mean



- $\text{Var}(\bar{X})$ is a measure of how “close” \bar{X} is to μ .
- How do we estimate $\text{Var}(\bar{X})$?

How “close” is our estimate \bar{X} to μ ?

$$E[\bar{X}] = \mu$$

$$\text{Var}(\bar{X}) = \frac{\sigma^2}{n}$$

We want to estimate this

def The **standard error** of the mean is an estimate of the standard deviation of \bar{X} .

$$SE = \sqrt{\frac{S^2}{n}}$$

Intuition:

- S^2 is an unbiased estimate of σ^2
- S^2/n is an unbiased estimate of $\sigma^2/n = \text{Var}(\bar{X})$
- $\sqrt{S^2/n}$ can estimate $\sqrt{\text{Var}(\bar{X})}$

More info on bias of standard error: [wikipedia](https://en.wikipedia.org/wiki/Standard_error)

Standard error of the mean

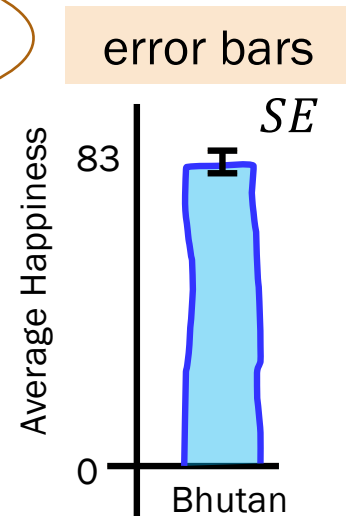
1. Mean happiness:

Claim: The average happiness of Bhutan is 83, with a standard error of 1.99.

Closed form: $SE = \sqrt{\frac{S^2}{n}}$

this is our estimate of how “close” we are

this is our best estimate of μ



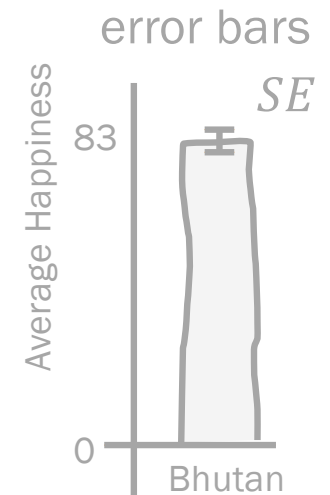
These 2 statistics give a sense of how the sample mean random variable \bar{X} behaves.

Standard error of variance?

1. Mean happiness:

Claim: The average happiness of Bhutan is 83, with a standard error of 1.99.

Closed form: $SE = \sqrt{\frac{S^2}{n}}$



2. Variance of happiness:

Claim: The variance of happiness of Bhutan is 793.

Closed form: Not covered in CS109

But how close are we?



this is our best estimate of σ^2

Up next: Compute Statistics with code!



19d_bootstrap_mean

Bootstrap: Sample mean

Bootstrap

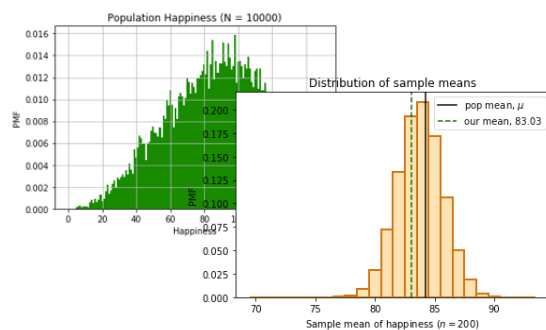
The Bootstrap:

Probability for Computer Scientists

Computing statistic of sample mean

What is the standard deviation of the sample mean \bar{X} ? (sample size $n = 200$)

Population
distribution
(we don't have this)



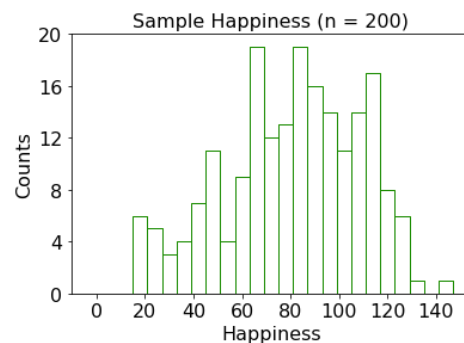
$$\frac{\sigma}{\sqrt{n}} = 1.886$$

Exact statistic
(we don't have this)

1.869

Simulated statistic
(we don't have this)

Sample
distribution
(we do have this)



$$SE = \frac{S}{\sqrt{n}} = 1.992$$

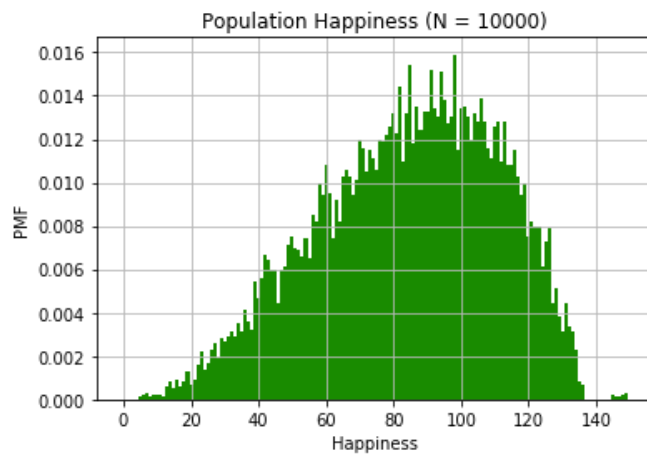
Estimated statistic,
by formula,
standard error

???

Simulated
estimated statistic

Note: We don't have access to the population.
But Lisa is sharing the exact statistic with you.

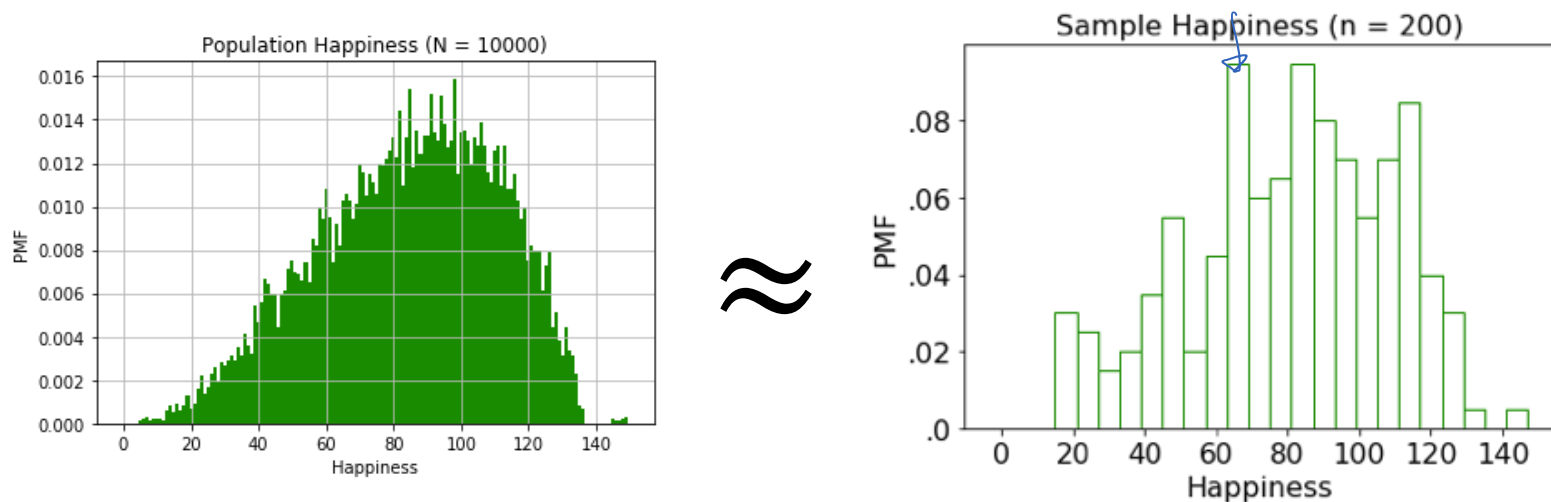
Bootstrap insight 1: Estimate the true distribution



\approx

Bootstrap insight 1: Estimate the true distribution

You can estimate the PMF of the underlying distribution, using your sample.*



The underlying
distribution



$$F \approx \hat{F}$$



the sample distribution
(aka the histogram of
your data)

*This is just a histogram of your data!

John, Chris Piech, Mehran Sahami, and Jerry Cain, CS109, Spring 2021

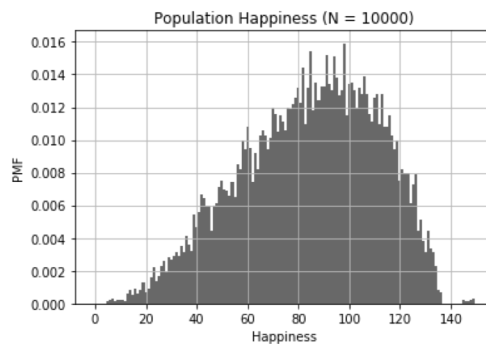
Bootstrap insight 2: Simulate a distribution

Approximate the procedure of simulating a distribution of a statistic, e.g., \bar{X} .

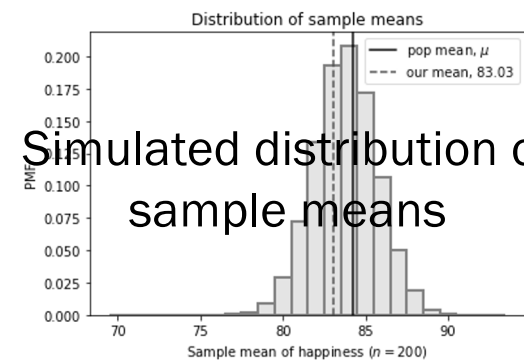
Population
distribution
(we don't have this)

\approx

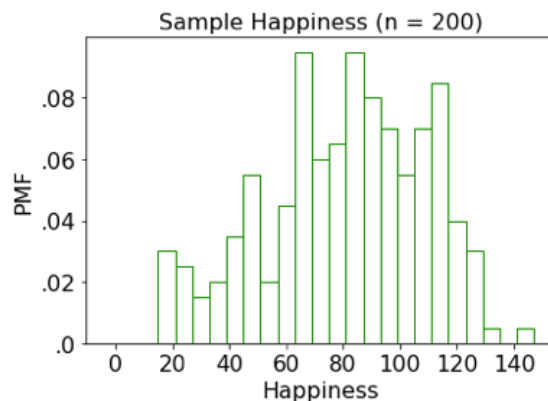
Sample
distribution
(we do have this)



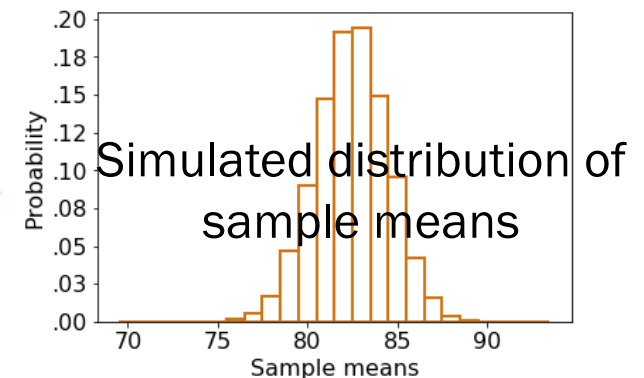
Distribution
of \bar{X}



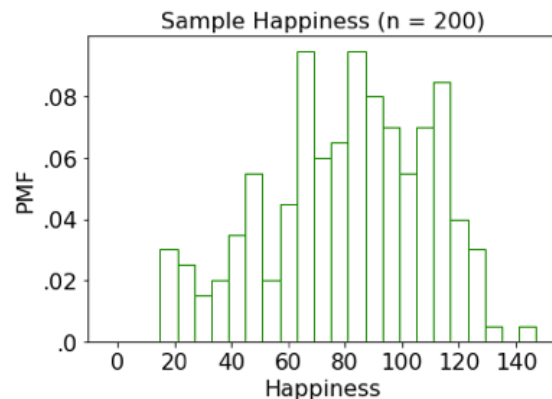
Simulated distribution of
sample means



Bootstrap
means

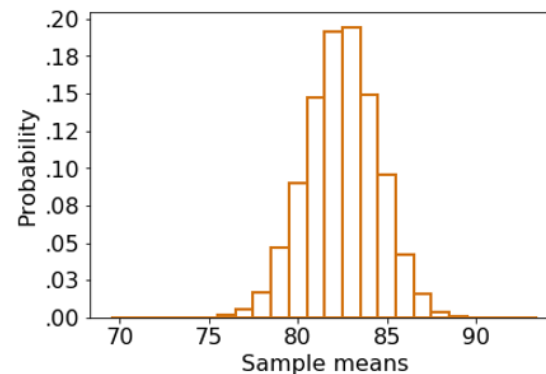


Bootstrapped sample means



Estimate the true PMF
using our “PMF” (histogram)
of our sample.

`means = [84.7,
83.9, 80.6, 79.8,
90.3, ..., 85.2]`



...generate a whole
bunch of sample means
of this estimated distribution...

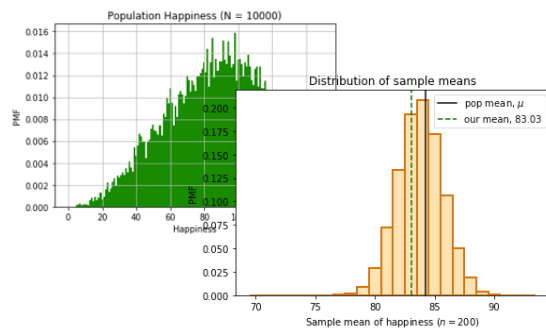
`np.std(means)`
2.003

...and compute the
standard deviation
of this distribution.

Computing statistic of sample mean

What is the standard deviation of the sample mean \bar{X} ? (sample size $n = 200$)

Population
distribution
(we don't have this)



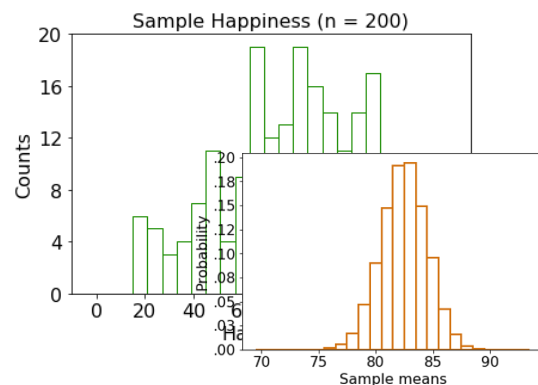
$$\frac{\sigma}{\sqrt{n}} = 1.886$$

Exact statistic
(we don't have this)

1.869

Simulated statistic
(we don't have this)

Sample
distribution
(we do have this)



$$SE = \frac{S}{\sqrt{n}} = 1.992$$

Estimated statistic,
by formula,
standard error

2.003

Simulated estimated
statistic, **bootstrapped
standard error**

Bootstrap algorithm

Bootstrap Algorithm (sample):

1. Estimate the **PMF** using the sample
2. Repeat **10,000** times:
 - a. Resample **sample.size()** from PMF
 - b. Recalculate the **sample mean** on the resample
3. You now have a **distribution of your sample mean**

What is the distribution of your **sample mean**?

We'll talk about this algorithm in detail during live lecture!

Bootstrap algorithm

Bootstrap Algorithm (sample):

1. Estimate the **PMF** using the sample
2. Repeat **10,000** times:
 - a. Resample **sample.size()** from PMF
 - b. Recalculate the **statistic** on the resample
3. You now have a **distribution of your statistic**

What is the distribution of your **statistic**?

Bootstrapped sample variance

Bootstrap Algorithm (sample):

1. Estimate the **PMF** using the sample
2. Repeat **10,000** times:
 - a. Resample **sample.size()** from PMF
 - b. Recalculate the **sample variance** on the resample
3. You now have a **distribution of your sample variance**

What is the distribution of your **sample variance**?

Even if we don't have a closed form equation,
we estimate statistics of sample variance with bootstrapping!

19: Sampling and the Bootstrap (live)

Jerry Cain
May 10, 2021

Think

Slide 42 has a question to go over by yourself.

Post any clarifications here or in Zoom chat!

<https://edstem.org/us/courses/5090/discussion/428950>

Think by yourself: 2 min



Quick check

1. μ , the population mean unknown value B
2. $(X_1, X_2, X_3, X_4, X_5, X_6, X_7, X_8)$, a sample A (8-D)
3. σ^2 , the population variance ??? B
4. \bar{X} , the sample mean A EV $\bar{X} = \frac{1}{n} \sum_{i=1}^n X_i$
5. $\bar{X} = 83$ C
6. $(X_1 = 59, X_2 = 87, X_3 = 94, X_4 = 99,$
 $X_5 = 87, X_6 = 78, X_7 = 69, X_8 = 91)$ C

- A. Random variable(s)
- B. Value
- C. Event



Quick check

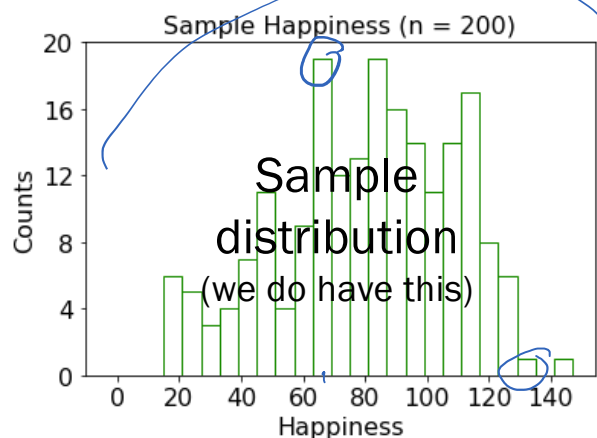
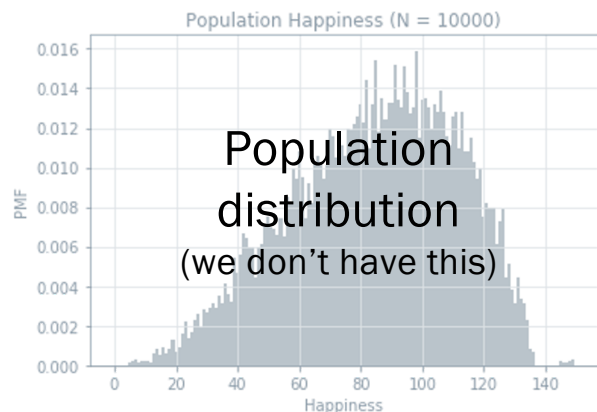
1. μ , the population mean
2. $(X_1, X_2, X_3, X_4, X_5, X_6, X_7, X_8)$, a sample
3. σ^2 , the population variance
4. \bar{X} , the sample mean
5. $\bar{X} = 83$
6. $(X_1 = 59, X_2 = 87, X_3 = 94, X_4 = 99,$
 $X_5 = 87, X_6 = 78, X_7 = 69, X_8 = 91)$

- A. Random variable(s)
- B. Value
- C. Event

These are outcomes
from your collected
data.

Today: Crash course on (bootstrapped) statistics

Review



If we only have a single sample of RVs generated i.i.d. from the same unknown distribution, how can we perform statistical analysis?

- What is the probability that a Bhutanese peep is just straight up loving life?
- What is a good estimate of the population mean (and how “close” is the estimate)?
- What is a good estimate of the population variance (and how “close” is the estimate)?

Standard error

Review

1. Mean happiness:

Claim: The average happiness of Bhutan is 83, with a standard error of 1.99.

this is our best estimate of μ

$$\bar{X} = \frac{1}{n} \sum_{i=1}^n X_i$$

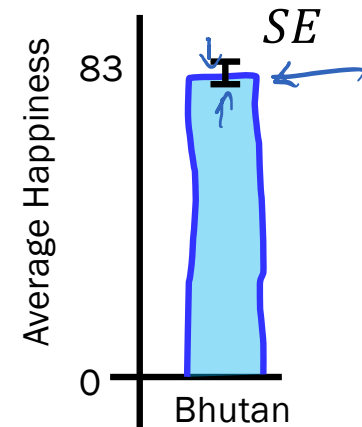
Closed form:

$$SE = \sqrt{\frac{S^2}{n}}$$

this is how close we are



Verified via bootstrap:
`np.std(means)`
`= 2.003`



Standard error

Review

1. Mean happiness:

Claim: The average happiness of Bhutan is 83,
with a standard error of 1.99.

Closed
form: $SE = \sqrt{\frac{S^2}{n}}$

2. Variance of happiness:

this is our best
estimate of σ^2

$$S^2 = \frac{1}{n-1} \sum_{i=1}^n (X_i - \bar{X})^2$$

Claim: The variance of happiness of Bhutan is 793.

Closed Not covered
form: in CS109

But how close
are we?

We can bootstrap for standard
error of sample variance—
a statistic of a statistic.

The Bootstrap:

Probability for Computer Scientists

Allows you to do the following:

- Calculate distributions over statistics
- Calculate p values *null hypotheses*

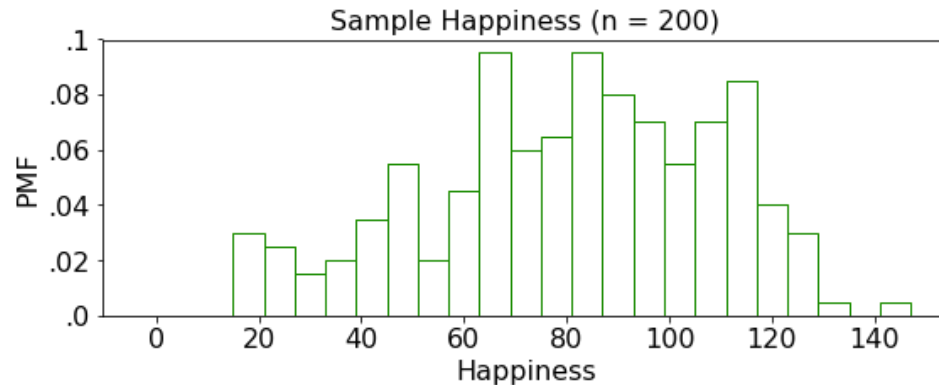
Bootstrapped sample variance

Bootstrap Algorithm (sample): $n=200$

1. Estimate the **PMF** using the sample
2. Repeat **10,000** times:
 - a. Resample **sample.size()** from PMF
 - b. Recalculate the **sample variance** on the resample
3. You now have a **distribution of your sample variance**

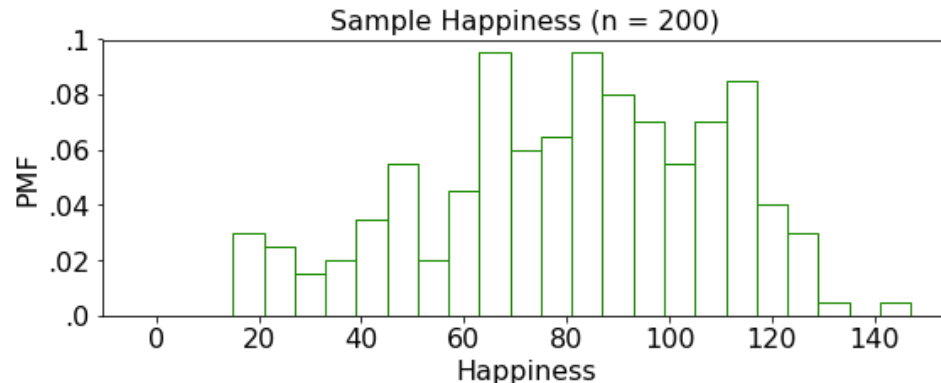
Goal What is the distribution of your **sample variance**?

Bootstrapped variance



- ➔ 1. Estimate the **PMF** using the sample
2. Repeat **10,000** times:
 - a. Resample `sample.size()` from PMF
 - b. Recalculate the **sample variance** on the resample
3. You now have a **distribution of your sample variance**

Bootstrapped variance



1. Estimate the **PMF** using the sample



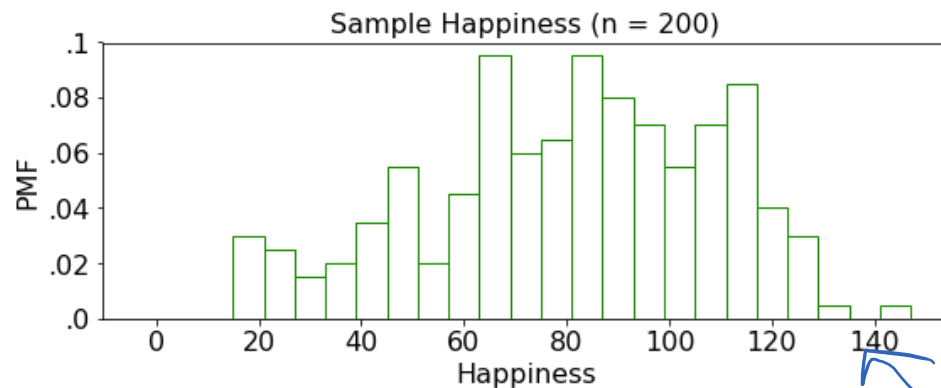
2. Repeat **10,000** times:

a. Resample **sample.size()** from PMF

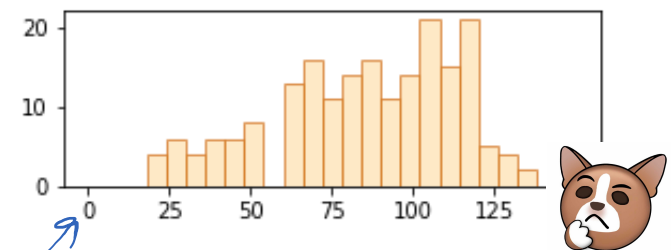
b. Recalculate the **sample variance** on the resample

3. You now have a **distribution of your sample variance**

Bootstrapped variance



[52, 38, 98, 107, ..., 94]



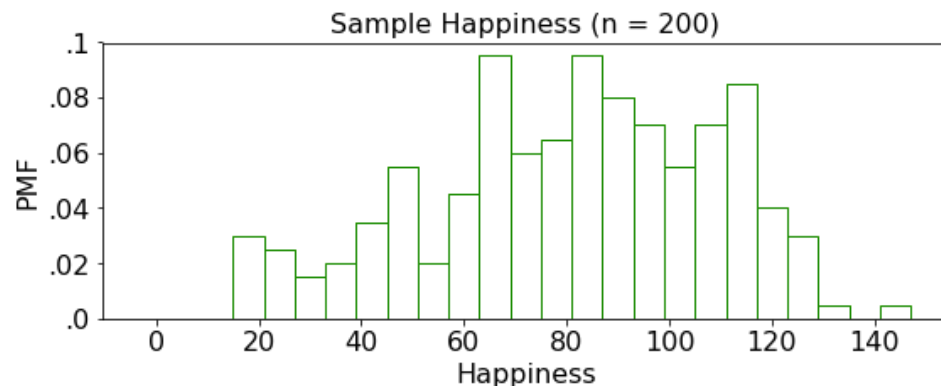
1. Estimate the **PMF** using the sample
2. Repeat **10,000** times:
 - a. Resample **sample.size()** from PMF
 - b. Recalculate the **sample variance** on the resample
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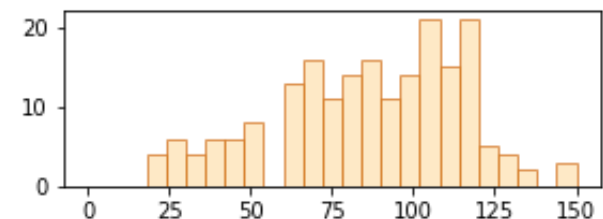
Why are these samples different?

This resampled sample is generated **with replacement**.

Bootstrapped variance



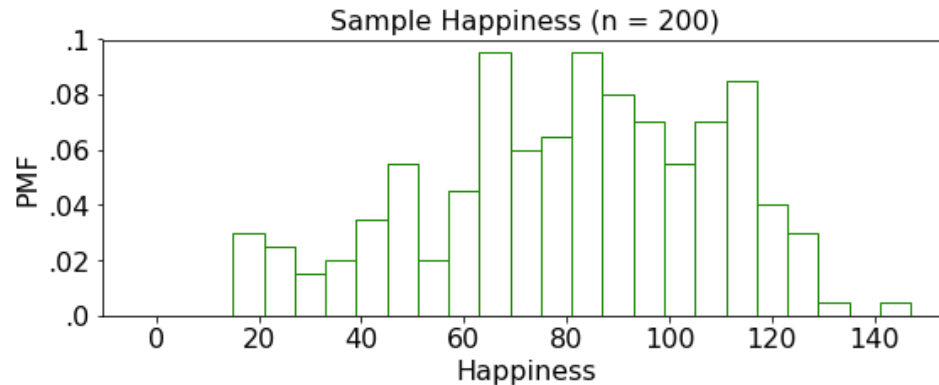
[52, 38, 98, 107, ..., 94]



1. Estimate the **PMF** using the sample
2. Repeat **10,000** times:
 - a. Resample **sample.size()** from PMF
 - ➡ b. Recalculate the **sample variance** on the resample
3. You now have a **distribution of your sample variance**

variances = [827.4]

Bootstrapped variance



1. Estimate the **PMF** using the sample



2. Repeat **10,000** times:

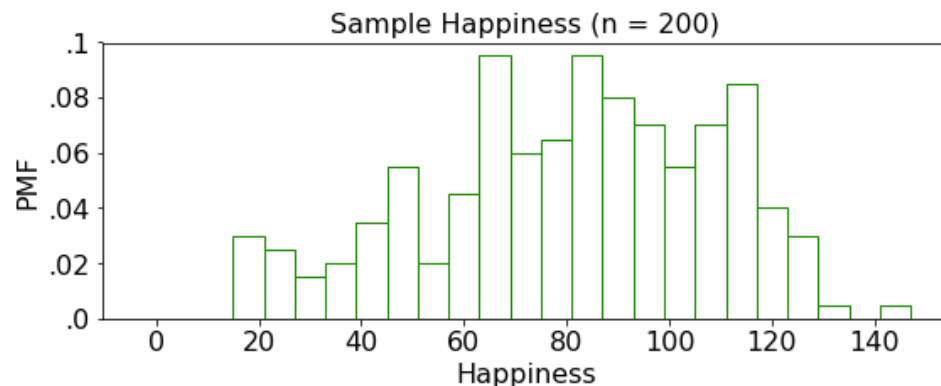
a. Resample **sample.size()** from PMF

b. Recalculate the **sample variance** on the resample

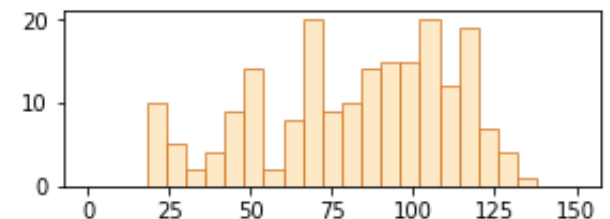
3. You now have a **distribution of your sample variance**

variances = [827.4]

Bootstrapped variance



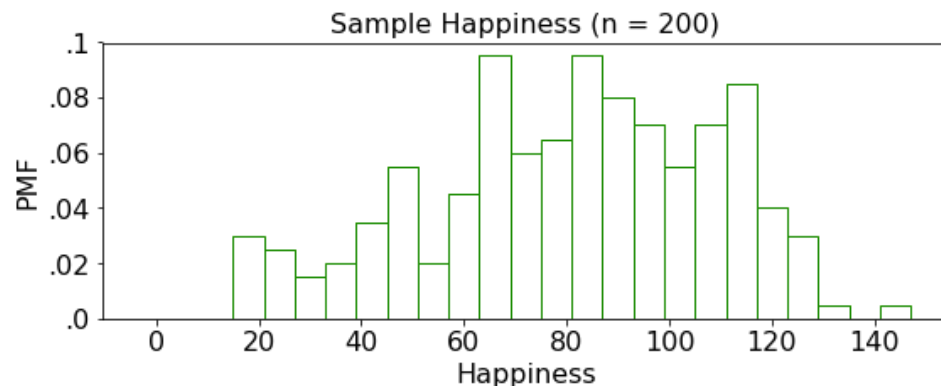
[116, 76, 132, 85, ..., 78]



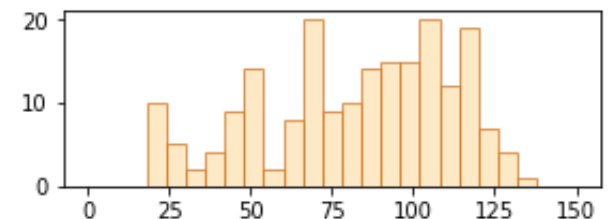
1. Estimate the **PMF** using the sample
2. Repeat **10,000** times:
 - a. Resample **sample.size()** from PMF
 - b. Recalculate the **sample variance** on the resample
3. You now have a **distribution of your sample variance**

variances = [827.4]

Bootstrapped variance



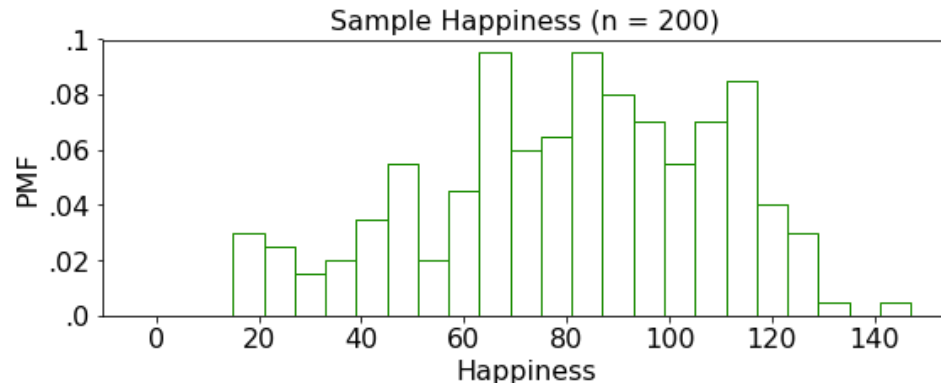
[116, 76, 132, 85, ..., 78]



1. Estimate the **PMF** using the sample
2. Repeat **10,000** times:
 - a. Resample **sample.size()** from PMF
 - ➡ b. Recalculate the **sample variance** on the resample
3. You now have a **distribution of your sample variance**

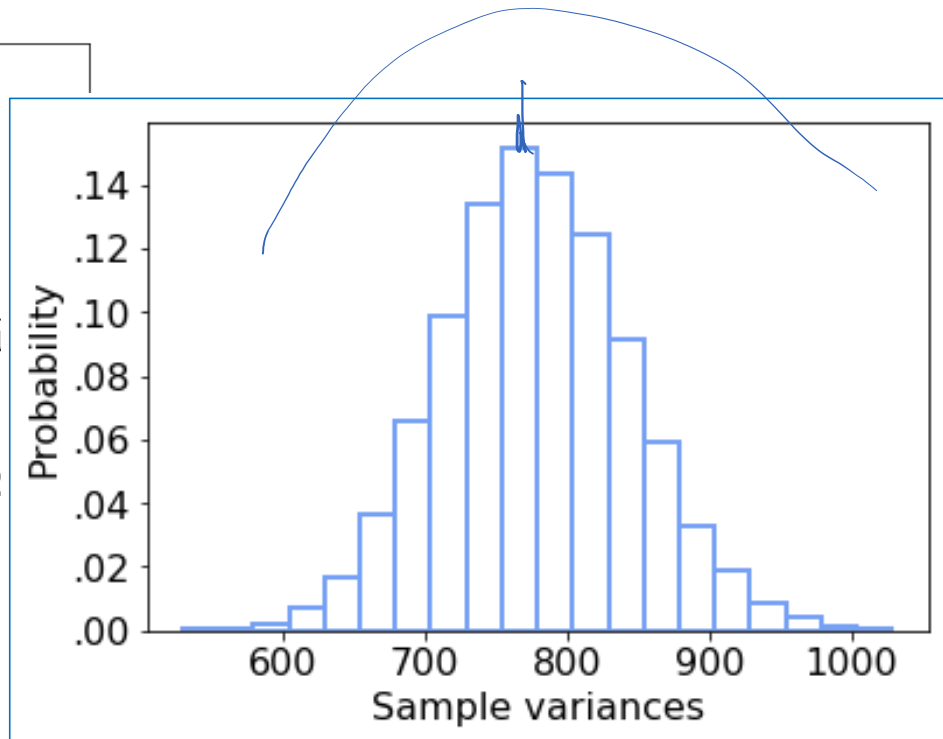
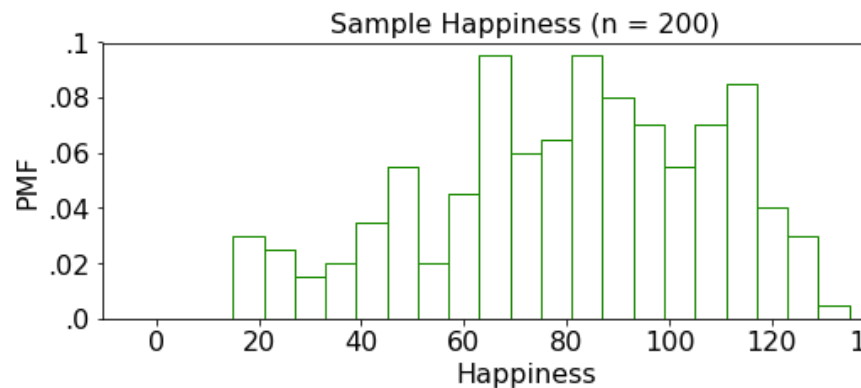
variances = [827.4, 846.1]

Bootstrapped variance



1. Estimate the **PMF** using the sample
 2. Repeat **10,000** times:
 - a. Resample **sample.size()** from PMF
 - b. Recalculate the **sample variance** on the resample
 3. You now have a **distribution of your sample variance**
- variances = [827.4, 846.1]**

Bootstrapped variance



1. Estimate the **PMF** using the
2. Repeat **10,000** times:
 - a. Resample **sample.size()**
 - b. Recalculate the **sample**
3. You now have a **distribution of your sample variance**

variances = [827.4, 846.1, 726.0, ..., 860.7]

Bootstrapped variance

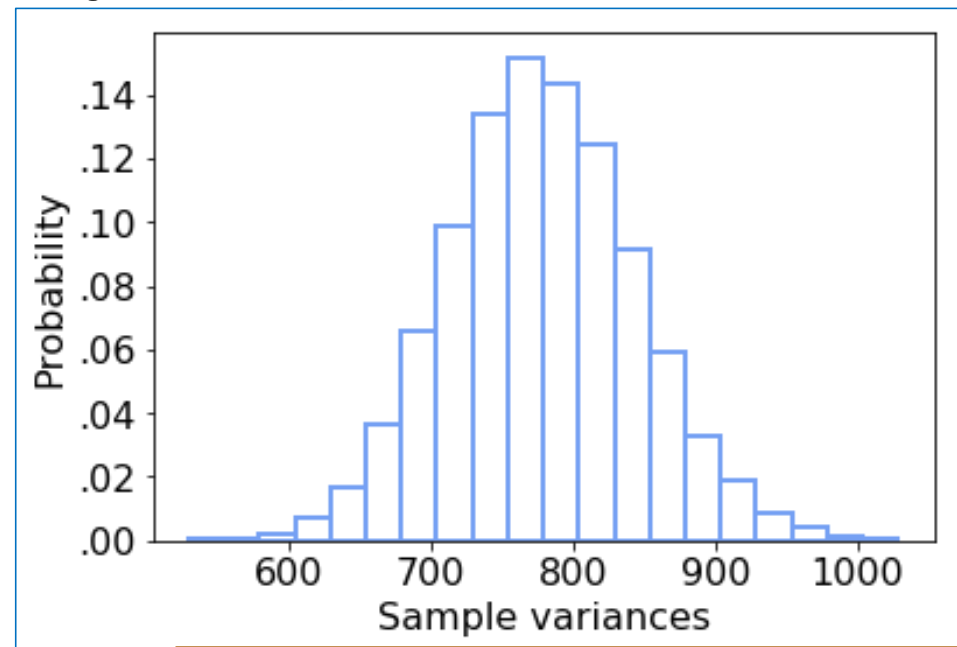
3. You now have a distribution of your **sample variance**

```
variances = [827.4,  
             846.1, 726.0, ...,  
             860.7]
```

What is the bootstrapped standard error?

```
( np.std(variances) )
```

Bootstrapped standard error: 66.16



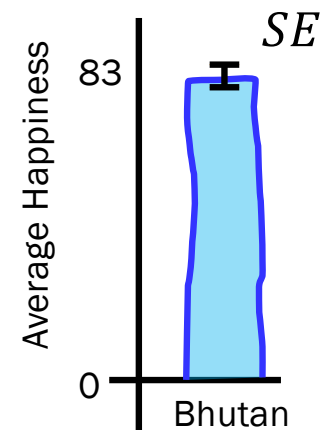
- Simulate a distribution of sample variances
- Compute standard deviation

Standard error

1. Mean happiness:

Claim: The average happiness of Bhutan is 83, with a standard error of 1.99.

Closed form: $SE = \sqrt{\frac{S^2}{n}}$



S^2 is our best estimate of σ^2

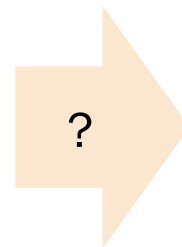
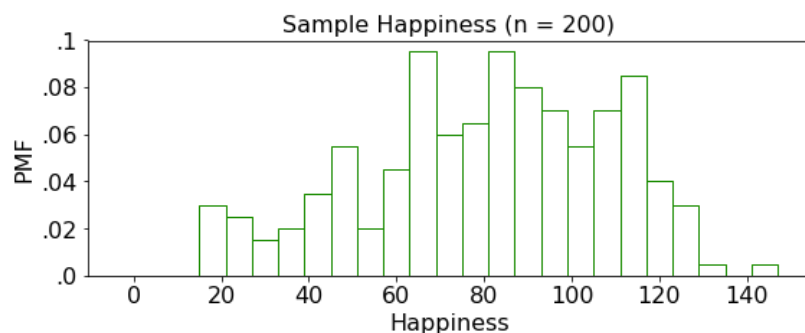
2. Variance of happiness:

Claim: The variance of happiness of Bhutan is 793, with a **bootstrapped standard error of 66.16**.

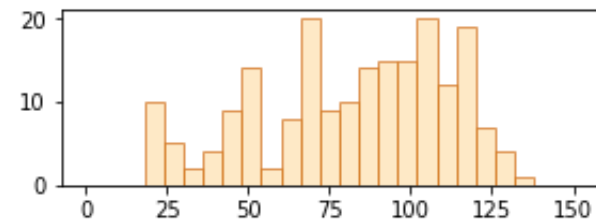
this is how close we are, calculated by bootstrapping

Algorithm in practice: Resampling

1. Estimate the **PMF** using the sample
2. Repeat **10,000** times:
 - a. Resample **sample.size()** from PMF
 - b. Recalculate the **statistic** on the resample
3. You now have a **distribution of your statistic**



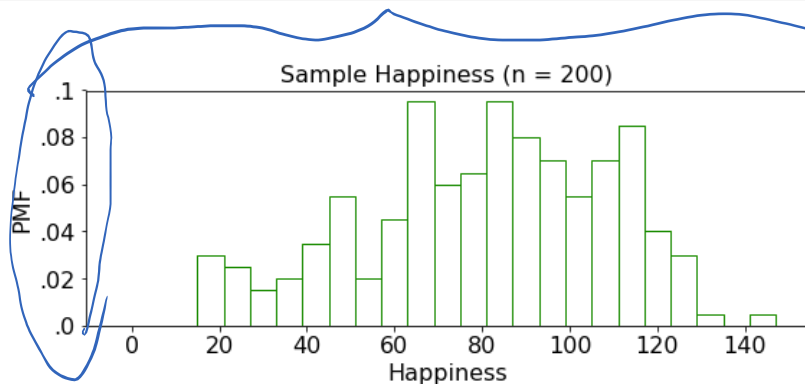
[116, 76, 132, 85, ..., 78]



$$P(X = k) = \frac{\text{\# values in sample equal to } k}{n}$$

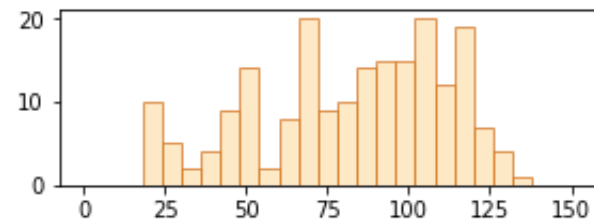
Algorithm in practice: Resampling

```
def resample(sample, n):  
    # estimate the PMF using the sample  
    # draw n new samples from the PMF  
    return np.random.choice(sample, n, replace=True)
```



?

[116, 76, 132, 85, ..., 78]



$$P(X = k) = \frac{\text{\# values in sample equal to } k}{n}$$

This resampled sample is generated **with replacement**.

To the code!

Bootstrap provides a way to calculate probabilities of statistics using code.

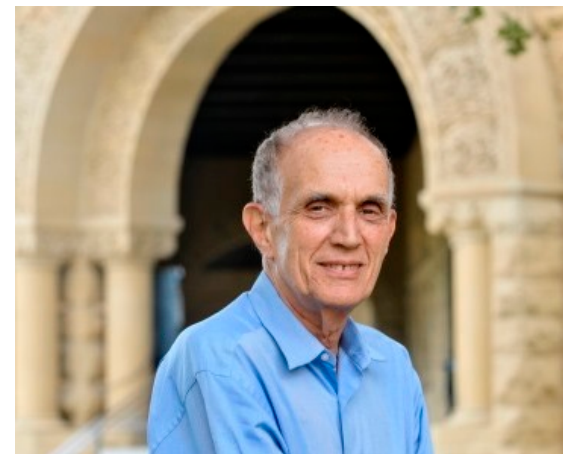
Bootstrapping works for any statistic*

*as long as your sample is i.i.d. and the underlying distribution does not have a long tail

Google colab notebook [link](#)
(we will use this in Breakout rooms)

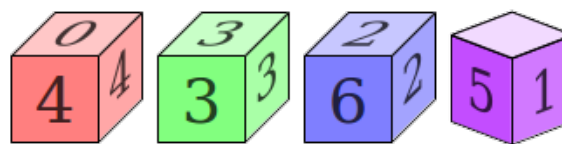
Bradley Efron

- Invented bootstrapping in 1979
- Still a professor at Stanford
- Won a National Science Medal



Efron's dice: 4 dice A, B, C, D such that

$$P(A > B) = P(B > C) = P(C > D) = P(D > A) = \frac{2}{3}$$



Interlude for announcements

Announcements

Problem Set 5

Out: now
Due: Friday 5/21 10:00am
Covers: Up to and including today

Quiz #2

Time frame: This Wednesday 5/12 11:00am – Friday 5/14 10:00am PT
Covers: Up to end of Week 5 (including Lecture 15). PS3+PS4
Emma's Review session: Tonight at 7pm PT (and will be recorded)
Info and practice: <http://web.stanford.edu/class/cs109/quizzes/>

LIVE

Bootstrap: p-value



Null hypothesis test

Nepal
Happiness

4.45

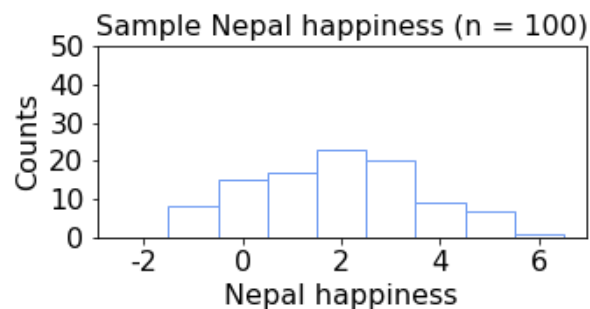
2.45

6.37

2.07

...

1.63



$$\bar{X}_1 = 3.1$$

Bhutan
Happiness

0.91

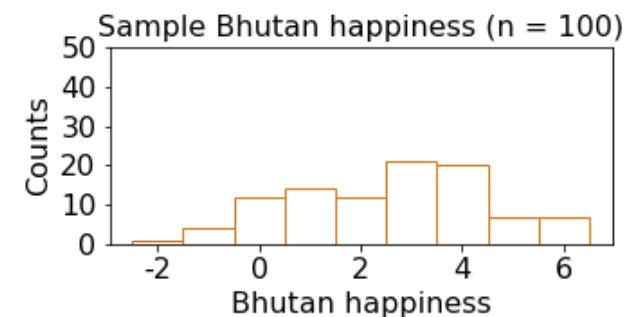
0.34

1.91

1.61

...

1.08



$$\bar{X}_2 = 2.4$$

Claim: The difference in mean happiness between Nepal and Bhutan is 0.7 happiness points, and **this is significant**.

Null hypothesis test

def **null hypothesis** – Even if there is no pattern (i.e., the two samples are from identical distributions), your claim might have arisen by chance.

def **p-value** – What is the probability that the observed difference occurs under the null hypothesis?

Example:

- Flip some coin 100 times.
- Flip the same coin another 150 times.
- Compute fraction of heads in both groups.
- There is a possibility we'll see the observed difference in these fractions even if we used the same coin

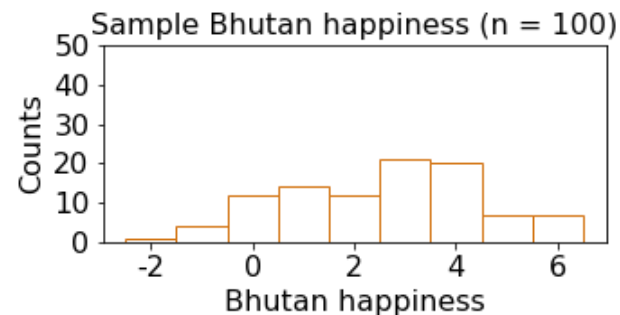
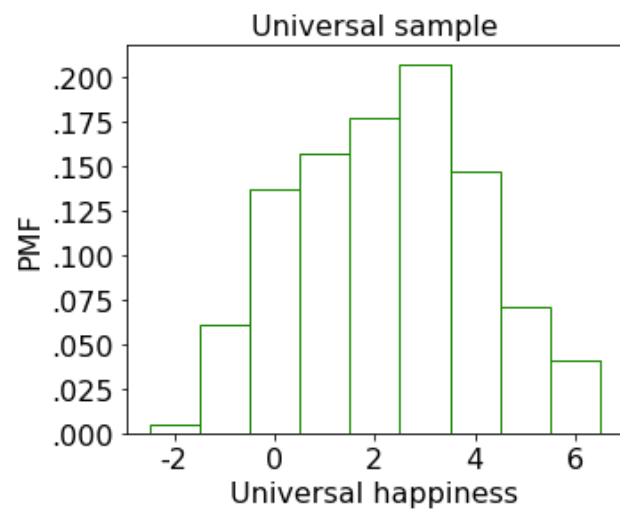
} **Null hypothesis** assumes we use the same coin

} **p-value**

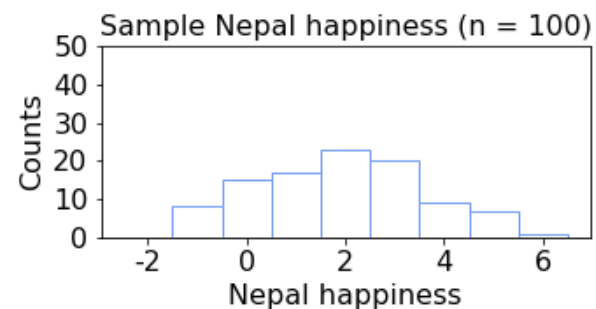
A **significant** p-value (< 0.05) means we reject the null hypothesis.

Universal sample

(this is what the null hypothesis assumes)



$$\bar{X}_1 = 3.1$$



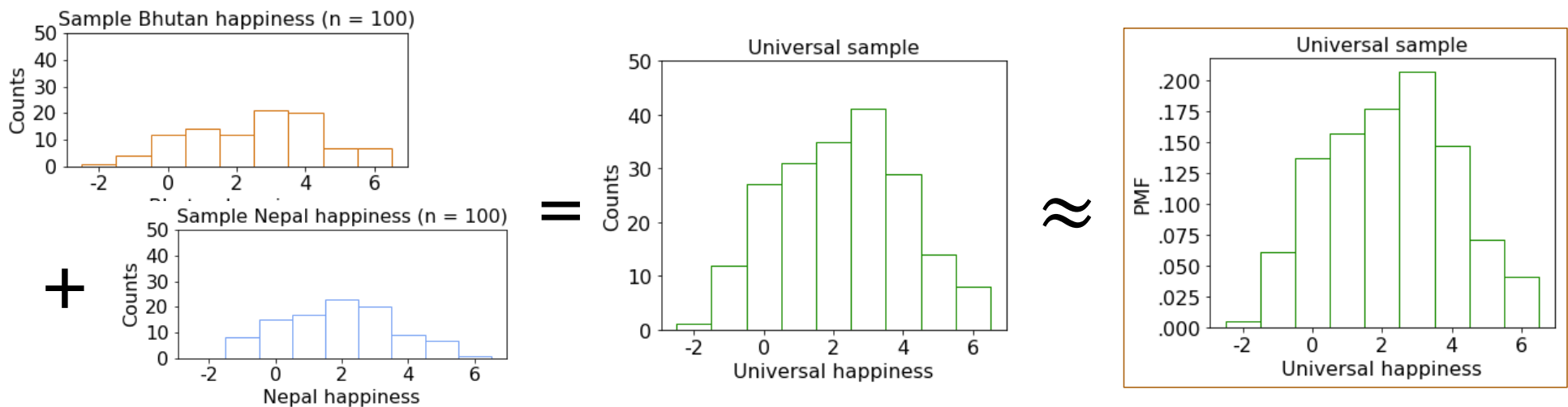
$$\bar{X}_2 = 2.4$$

Want **p-value**: probability $|\bar{X}_1 - \bar{X}_2| \geq |3.1 - 2.4|$ happens under null hypothesis

Bootstrap for p-values

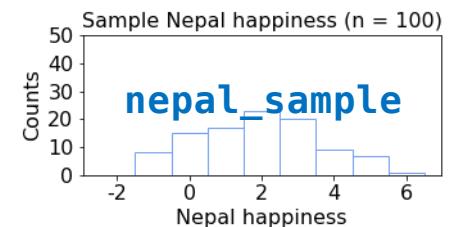
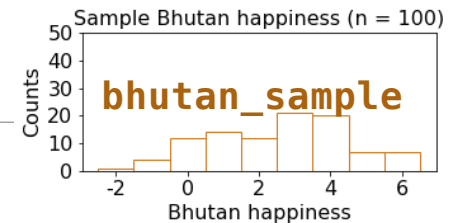
1. Create a **universal sample** using your two samples

i.e., recreate the null hypothesis



Bootstrap for p-values

1. Create a **universal sample** using your two samples
2. Repeat **10,000** times:
 - a. Resample **both samples**
 - b. Recalculate the **mean difference** between the resamples
3. **p-value** =
$$\frac{\# (\text{mean diffs} \geq \text{observed diff})}{n}$$



Probability
that observed
difference arose
by chance

Bootstrap for p-values

```
def pvalue_boot(bhutan_sample, nepal_sample):  
    N = size of the bhutan_sample  
    M = size of the nepal_sample  
    observed_diff = |mean of bhutan_sample - mean of nepal_sample|  
    uni_sample = combine bhutan_sample and nepal_sample  
    count = 0  
  
    repeat 10,000 times:  
        bhutan_resample = draw N resamples from the uni_sample  
        nepal_resample = draw M resamples from the uni_sample  
        muBhutan = sample mean of the bhutan_resample  
        muNepal = sample mean of the nepal_resample  
        diff = |muNepal - muBhutan|  
        if diff >= observed_diff:  
            count += 1
```

pValue = count / 10,000

Bootstrap for p-values

1. Create a universal sample using your two samples

```
def pvalue_boot(bhutan_sample, nepal_sample):  
    N = size of the bhutan_sample  
    M = size of the nepal_sample  
    observed_diff = |mean of bhutan_sample – mean of nepal_sample|  
  
    uni_sample = combine bhutan_sample and nepal_sample  
    count = 0
```

repeat 10,000 times:

```
    bhutan_resample = draw N resamples from the uni_sample  
    nepal_resample = draw M resamples from the uni_sample  
    muBhutan = sample mean of the bhutan_resample  
    muNepal = sample mean of the nepal_resample  
    diff = |muNepal – muBhutan|  
    if diff >= observed_diff:  
        count += 1
```

pValue = count / 10,000

Bootstrap for p-values

2. a. Resample both samples

```
def pvalue_boot(bhutan_sample, nepal_sample):  
    N = size of the bhutan_sample  
    M = size of the nepal_sample  
    observed_diff = |mean of bhutan_sample – mean of nepal_sample|  
  
    uni_sample = combine bhutan_sample and nepal_sample  
    count = 0  
  
    repeat 10,000 times:  
        bhutan_resample = draw N resamples from the uni_sample  
        nepal_resample = draw M resamples from the uni_sample  
        muBhutan = sample mean of the bhutan_resample  
        muNepal = sample mean of the nepal_resample  
        diff = |muNepal – muBhutan|  
        if diff >= observed_diff:  
            count += 1  
  
    pValue = count / 10,000
```

Bootstrap for p-values

2. b. Recalculate the **mean difference** b/t resamples

```
def pvalue_boot(bhutan_sample, nepal_sample):  
    N = size of the bhutan_sample  
    M = size of the nepal_sample  
    observed_diff = |mean of bhutan_sample – mean of nepal_sample|  
  
    uni_sample = combine bhutan_sample and nepal_sample  
    count = 0  
  
    repeat 10,000 times:  
        bhutan_resample = draw N resamples from the uni_sample  
        nepal_resample = draw M resamples from the uni_sample  
        muBhutan = sample mean of the bhutan_resample  
        muNepal = sample mean of the nepal_resample  
        diff = |muNepal – muBhutan|  
        if diff >= observed_diff:  
            count += 1
```

pValue = count / 10,000

Bootstrap for p-values

$$3. \text{ p-value} = \frac{\# (\text{mean diffs} > \text{observed diff})}{n}$$

```
def pvalue_boot(bhutan_sample, nepal_sample):  
    N = size of the bhutan_sample  
    M = size of the nepal_sample  
    observed_diff = |mean of bhutan_sample - mean of nepal_sample|  
  
    uni_sample = combine bhutan_sample and nepal_sample  
    count = 0  
  
    repeat 10,000 times:  
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        muBhutan = sample mean of the bhutan_resample  
        muNepal = sample mean of the nepal_resample  
        diff = |muNepal - muBhutan|  
        if diff >= observed_diff:  
            count += 1
```

pValue = count / 10,000

Lisa Yan, Chris Piech, Mehran Sahami, and Jerry Cain, CS109, Spring 2021

Bootstrap for p-values

```
def pvalue_boot(bhutan_sample, nepal_sample):  
    N = size of the bhutan_sample  
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        muBhutan = sample mean of the bhutan_resample  
        muNepal = sample mean of the nepal_resample  
        diff = |muNepal – muBhutan|  
        if diff >= observed_diff:  
            count += 1  
  
pValue = count / 10,000
```

with replacement!

Bootstrap



Let's try it!

Google colab notebook [link](#)

Null hypothesis test

Nepal
Happiness

4.45

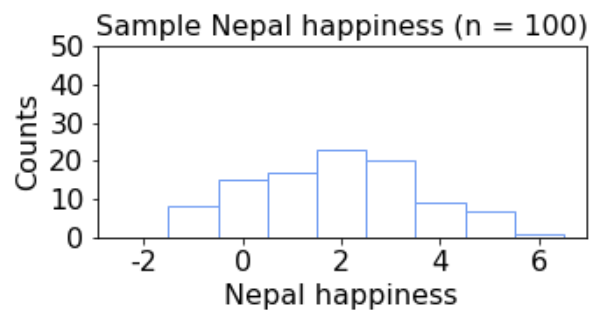
2.45

6.37

2.07

...

1.63



Bhutan
Happiness

0.91

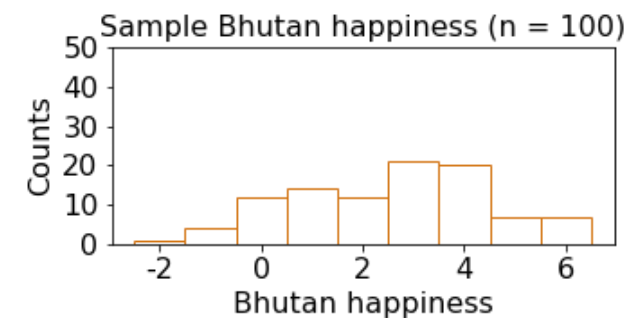
0.34

1.91

1.61

...

1.08



$$\bar{X}_1 = 3.1$$

$$\bar{X}_2 = 2.4$$

Claim: The happiness of Nepal and Bhutan have a 0.7 difference of means, and this is significant ($p < 0.05$).