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19: Sampling and the Bootstrap



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[Lecture Discussion on Ed](#)



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



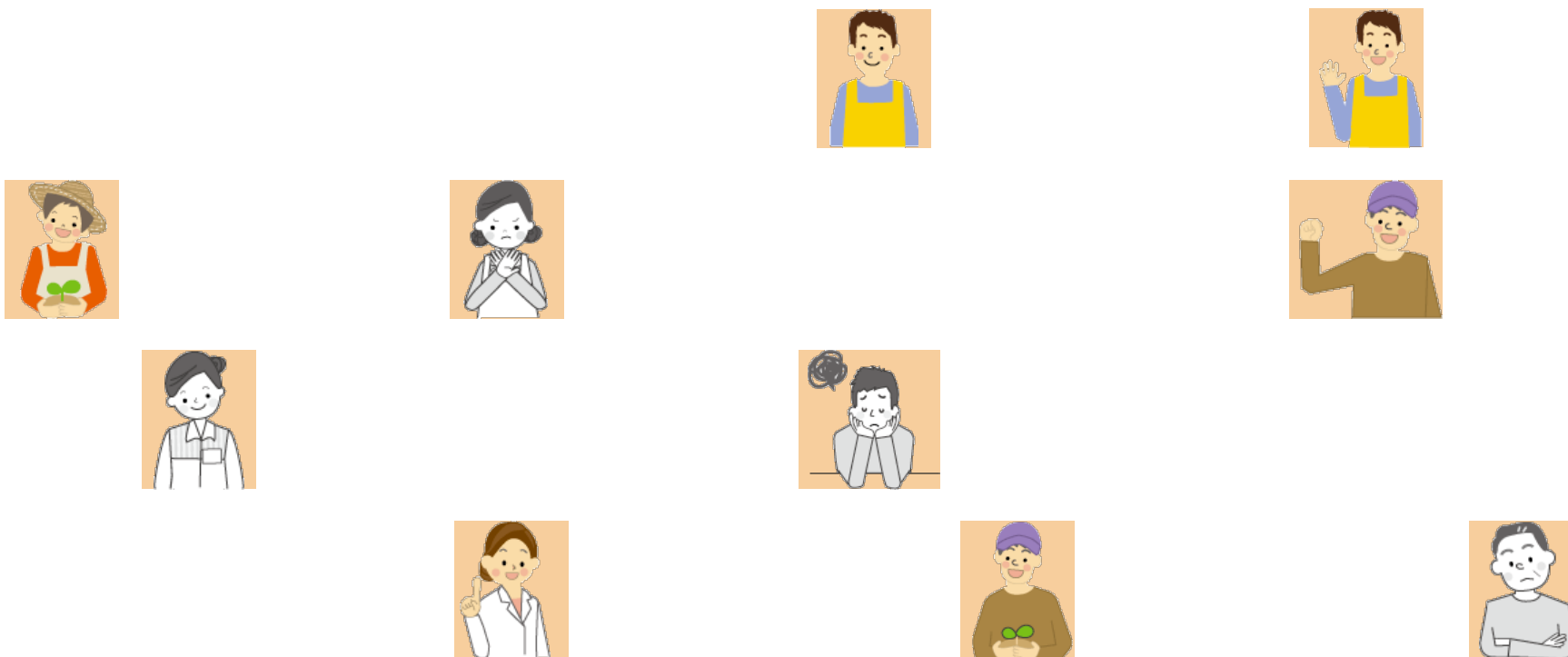
This is a **population**.

Sample



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

Sample



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

Reasonable Questions Starting Out

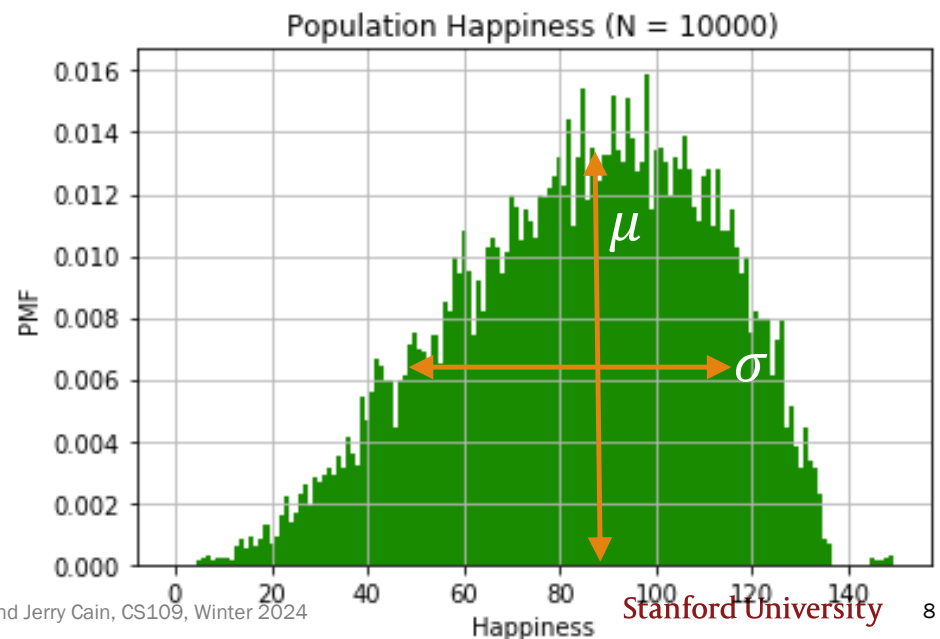
1. In situations where we can't observe the entire population, what can we safely infer by polling a sample drawn from that population?
2. How large does your sample need to be before your conclusions become trustworthy, and how do we express confidence in what we conclude.
3. Are there alternative ways to infer population statistics without polling entire populations?

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 (iid)
- X_i all have same distribution function F (the **underlying distribution**), where $E[X_i] = \mu$, $\text{Var}(X_i) = \sigma^2$



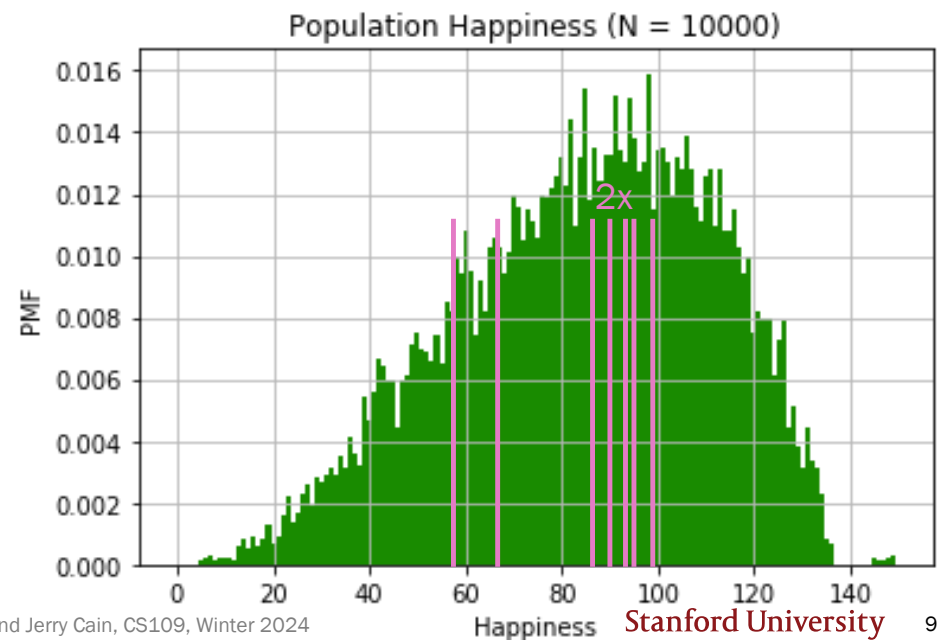
A sample, mathematically

A sample of **size** 8:

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

The **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—or rather, a sample.

Today: If we only have a single sample,

- How do we report **estimated** statistics?
 - We're careful to call them estimated mean and estimated variance, since they're based on samples (i.e., experiments)
- How do we report estimated errors on these estimates?
- How do we perform something called **hypothesis testing**? Oh, and what is it?



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 happiness.

But we only have 200 people (a sample).

These population-level statistics are unknown:

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

A single sample



A happy
Bhutanese person

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

But we only have 200 people (a sample).

- From these 200 people, what is our best estimate of the **population mean** and the **population variance**?
- How exactly 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 (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 .

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

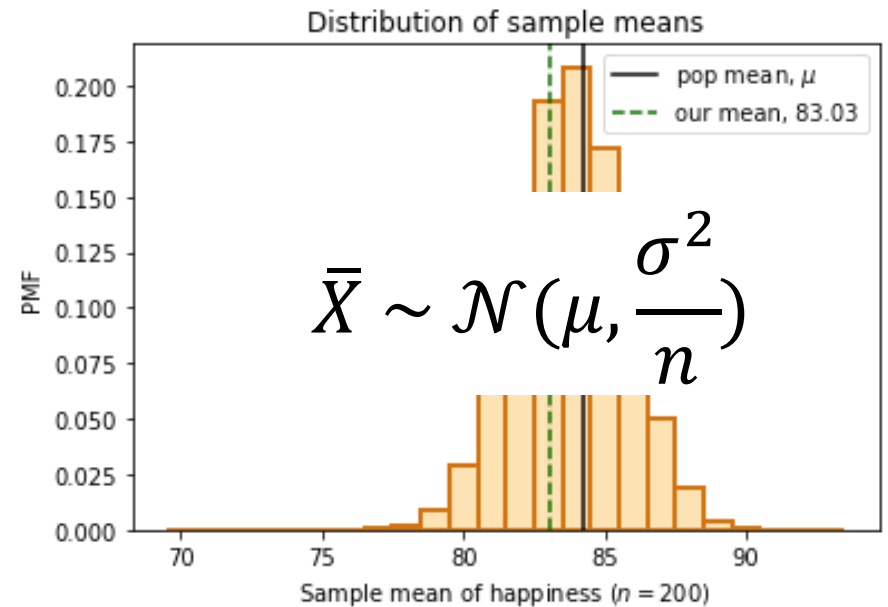
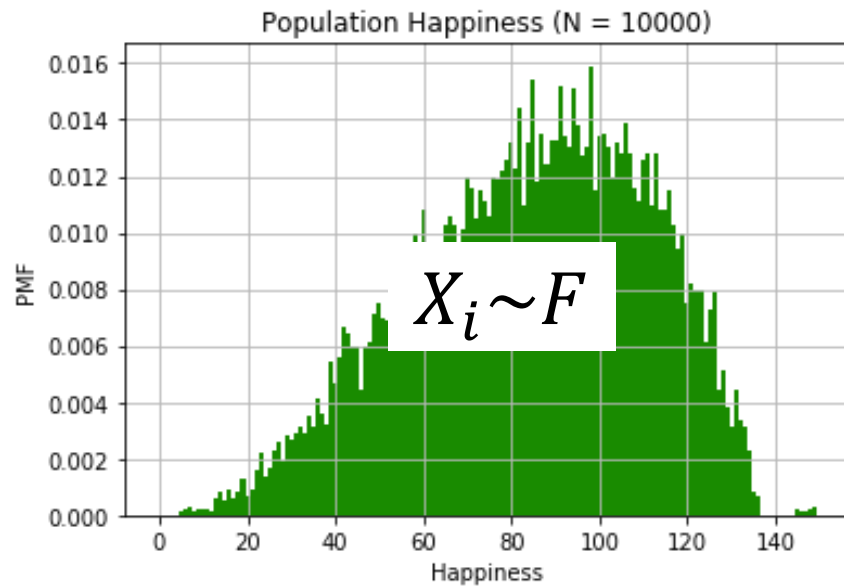
Intuition: By the CLT, $\bar{X} \sim \mathcal{N}\left(\mu, \frac{\sigma^2}{n}\right)$



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 one 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

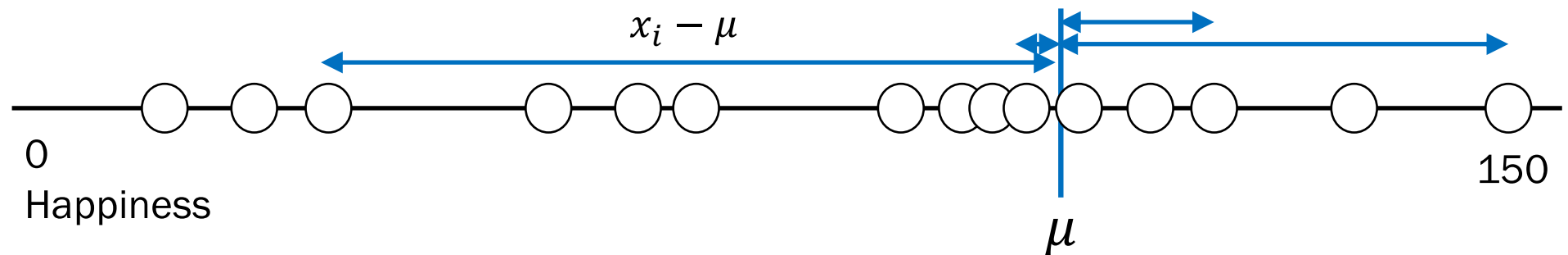
Intuition about the sample variance, S^2

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

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

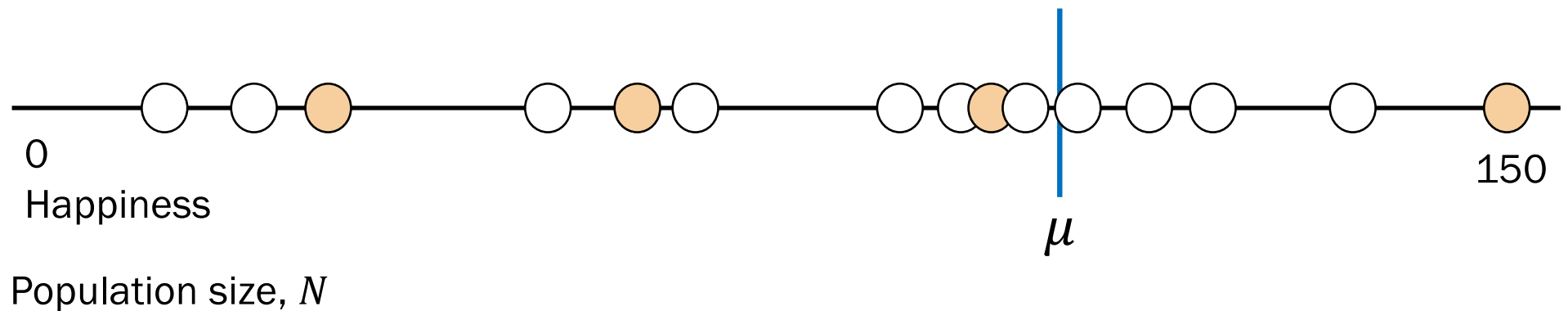
population mean

Estimate, S^2

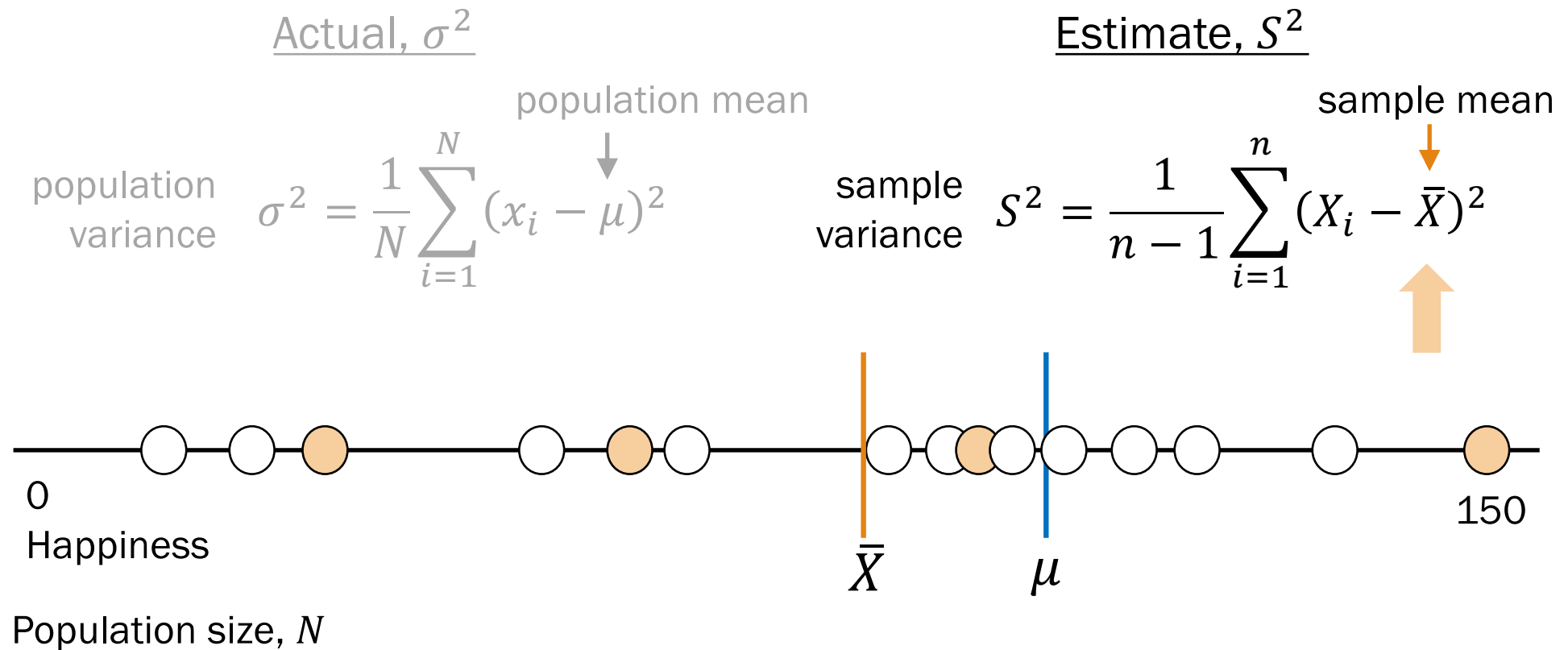
sample variance

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

sample mean



Intuition about the sample variance, S^2



Intuition about the sample variance, S^2

Actual, σ^2

population variance

population mean

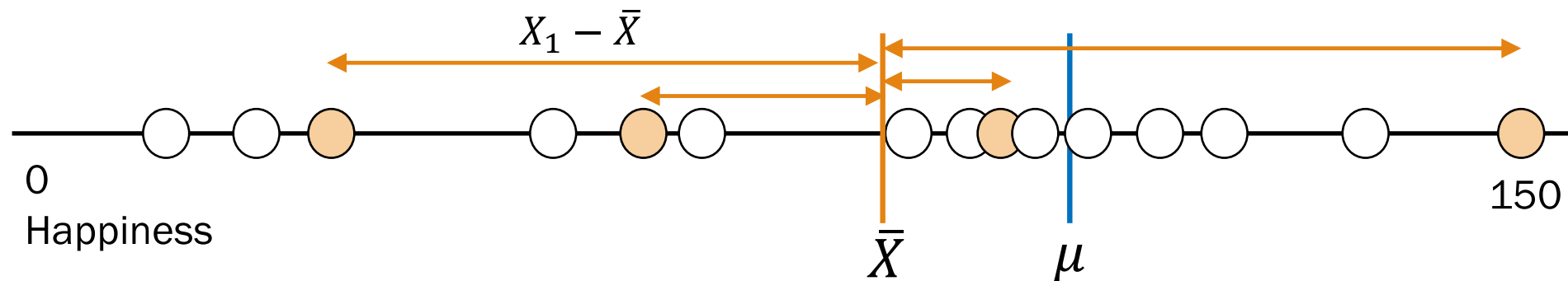
$$\sigma^2 = \frac{1}{N} \sum_{i=1}^N (x_i - \mu)^2$$

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

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



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

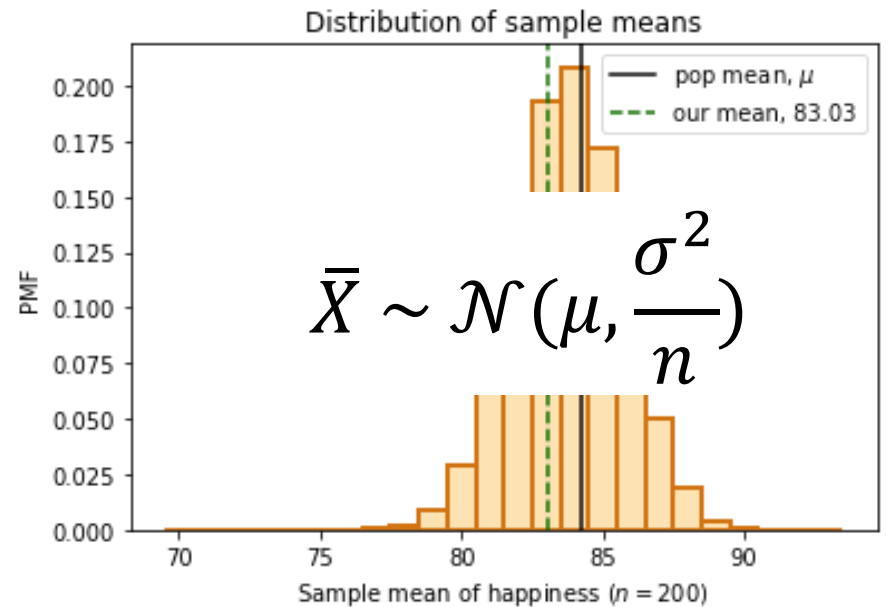
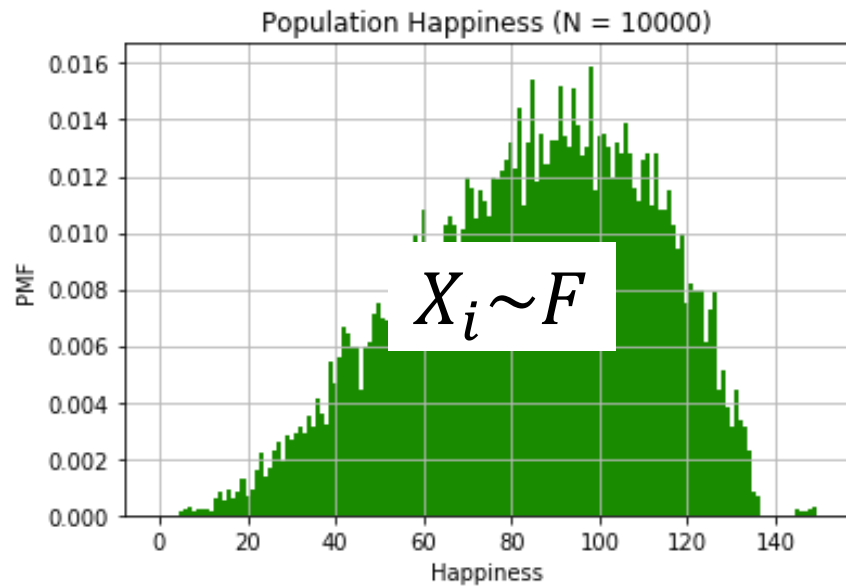
$\bar{X} = 83$

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

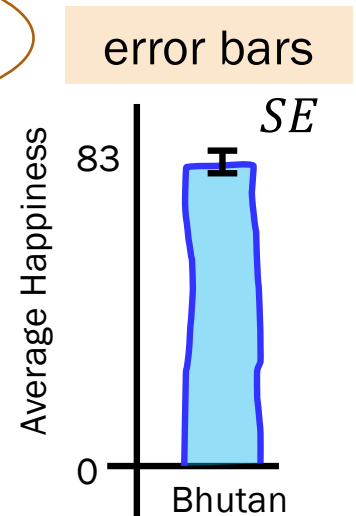
1. Mean happiness:

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



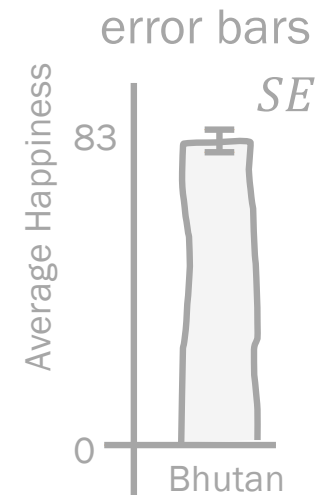
These 2 statistics give a sense of how \bar{X} —that is, the sample mean random variable—behaves.

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!



Bootstrap: Sample mean

Bootstrap

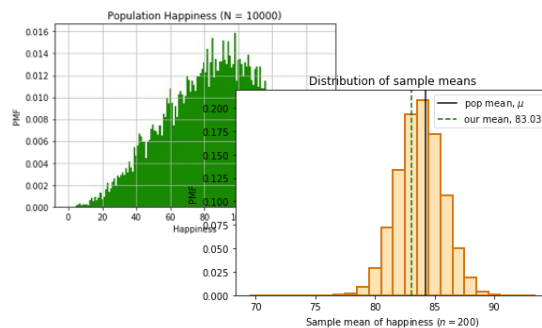
The Bootstrap:

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)



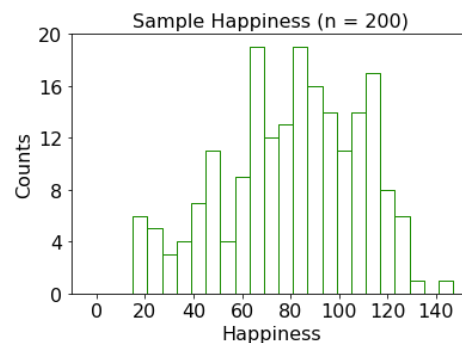
$$\frac{\sigma}{\sqrt{n}} = 1.886$$

Exact statistic
(we don't have this)

$$1.869$$

Simulated statistic
(we don't have this)

Sample
distribution
(we do have this)



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

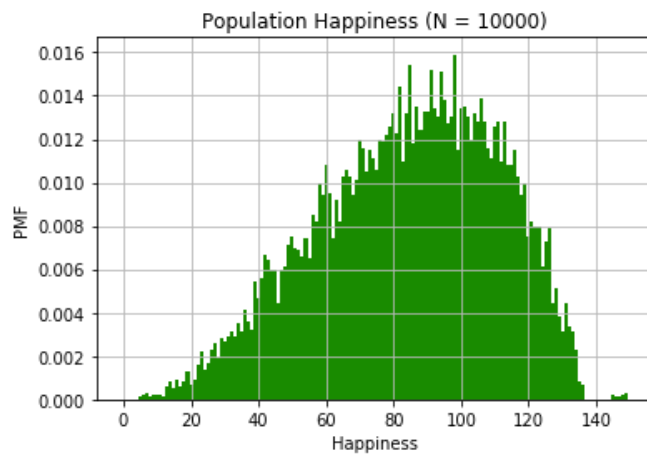
Estimated statistic,
by formula,
standard error

???

Simulated
estimated statistic

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

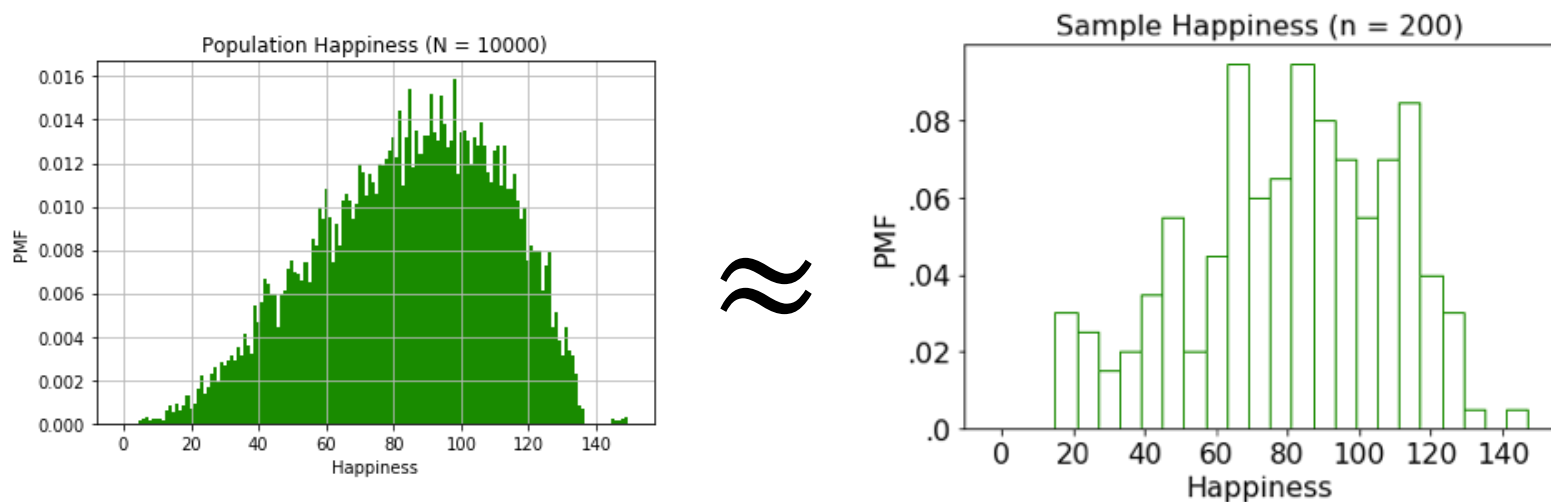
Bootstrap insight 1: Estimate the true distribution



\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!

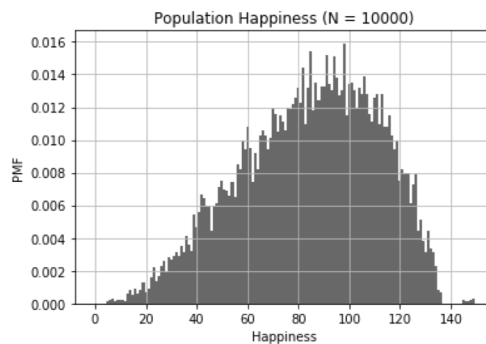
Bootstrap insight 2: Simulate a distribution

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

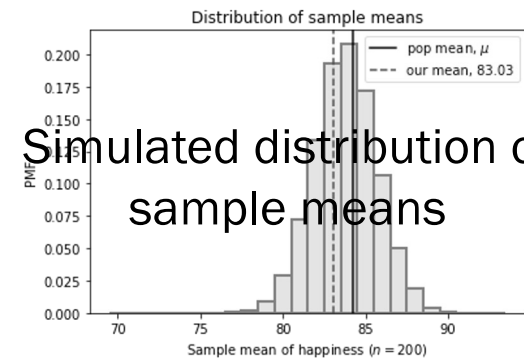
Population
distribution
(we don't have this)



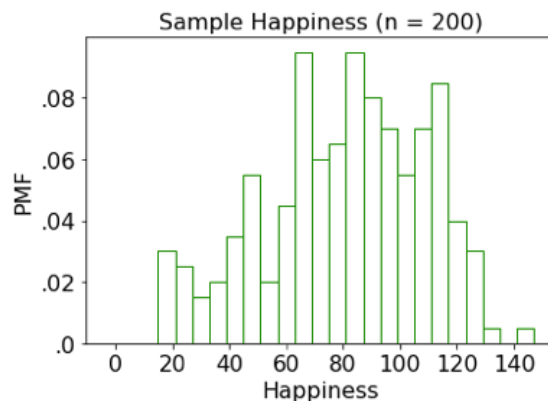
Sample
distribution
(we do have this)



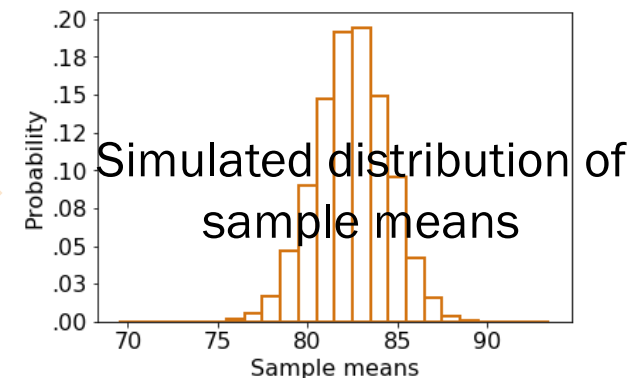
Distribution
of \bar{X}



Simulated distribution of
sample means



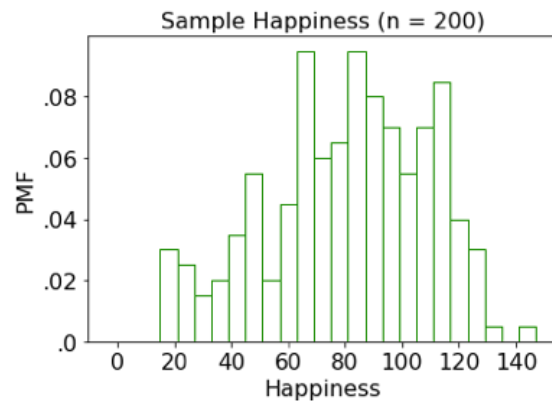
Bootstrap
means



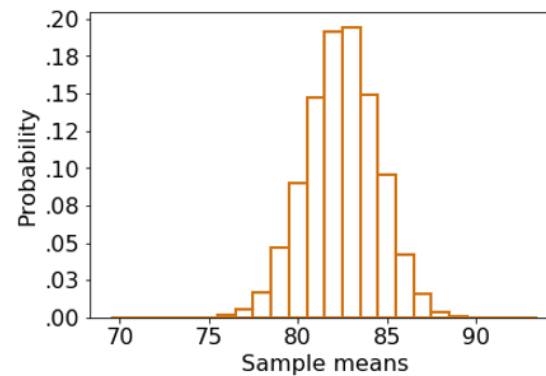
Simulated distribution of
sample means

Bootstrapped sample means

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



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



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

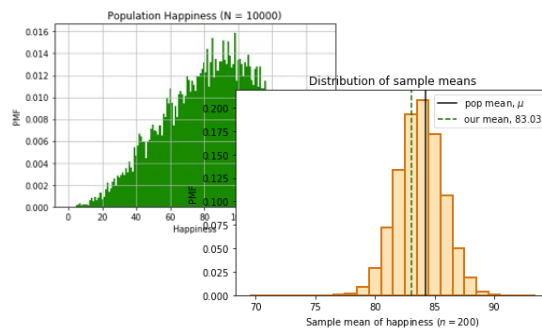
`np.std(means)`
2.003

...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)



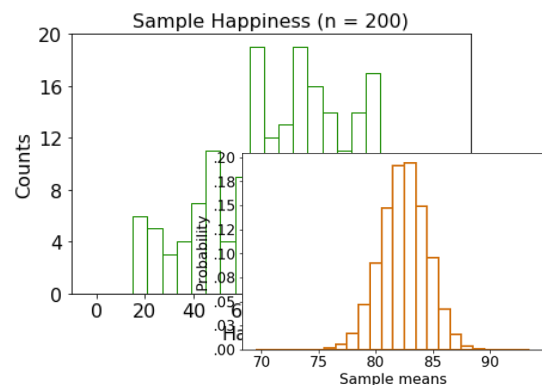
$$\frac{\sigma}{\sqrt{n}} = 1.886$$

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 with a demo!

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!



Bootstrap: Sample variance

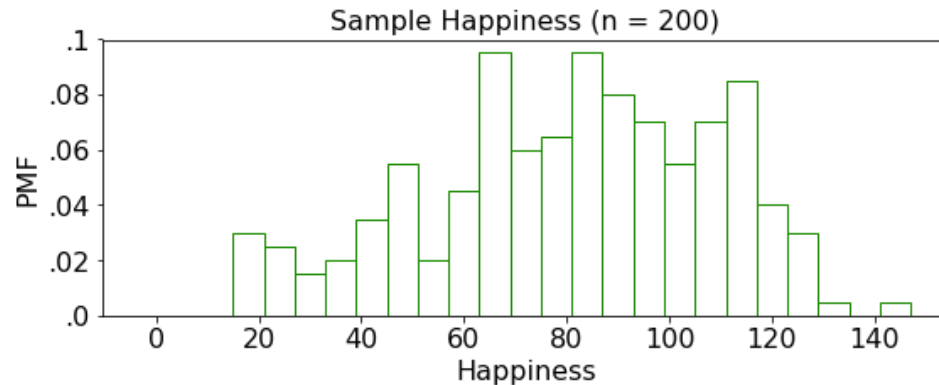
Bootstrapped sample variance

Bootstrap Algorithm (sample):

1. Estimate the **PMF** using the sample
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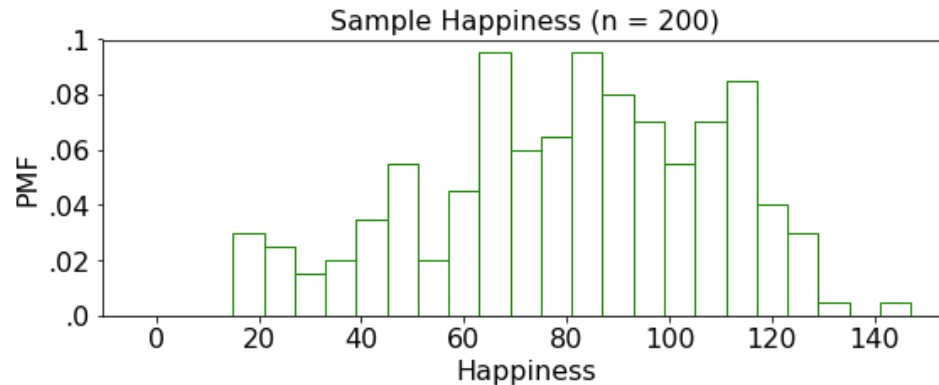
Goal What is the distribution of your **sample variance**?

Bootstrapped variance



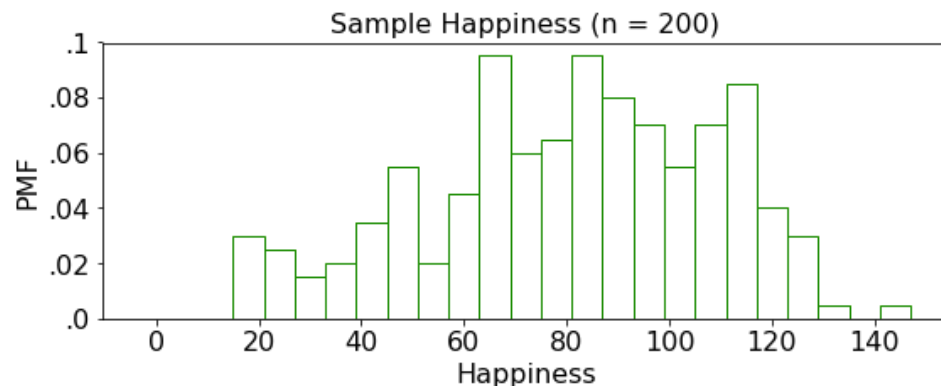
- ➔ 1. Estimate the **PMF** using the sample
2. Repeat **10,000** times:
 - a. Resample `sample.size()` from PMF
 - b. Recalculate the **sample variance** on the resample
3. You now have a **distribution of your sample variance**

Bootstrapped variance

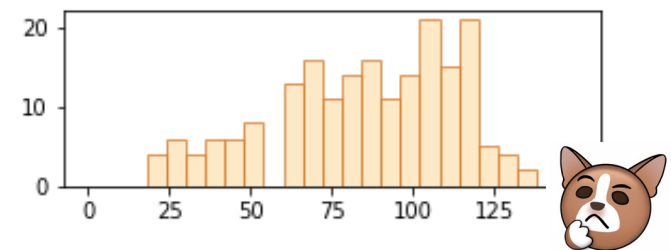


1. Estimate the **PMF** using the sample
- ➔ 2. Repeat **10,000** times:
 - a. Resample **sample.size()** from PMF
 - b. Recalculate the **sample variance** on the resample
3. You now have a **distribution of your sample variance**

Bootstrapped variance



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



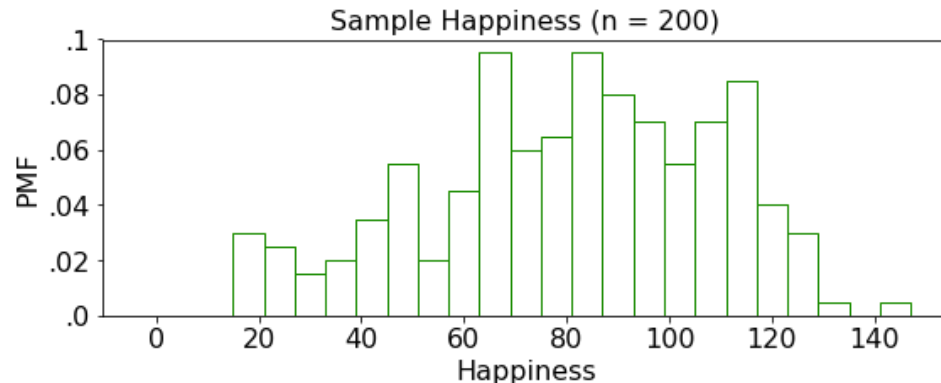
1. Estimate the **PMF** using the sample
2. Repeat **10,000** times:
 - a. Resample `sample.size()` from PMF
 - b. Recalculate the **sample variance** on the resample
3. You now have a **distribution of your**



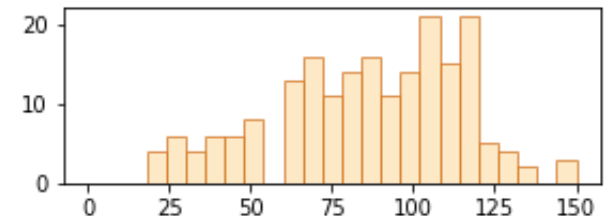
Why are these samples different?

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

Bootstrapped variance



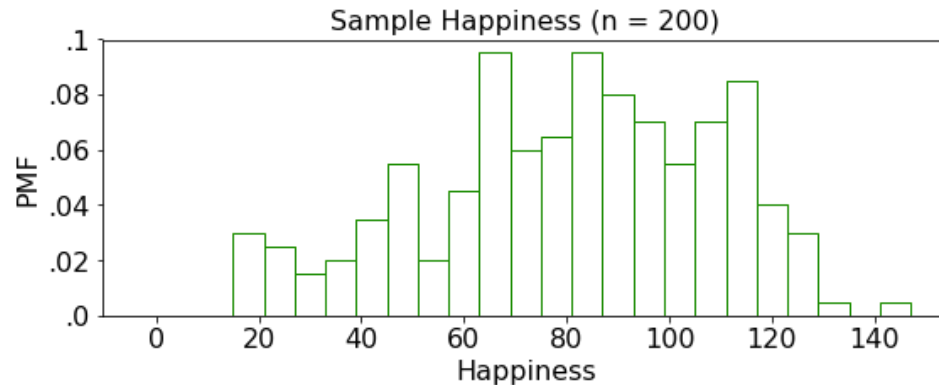
[52, 38, 98, 107, ..., 94]



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:

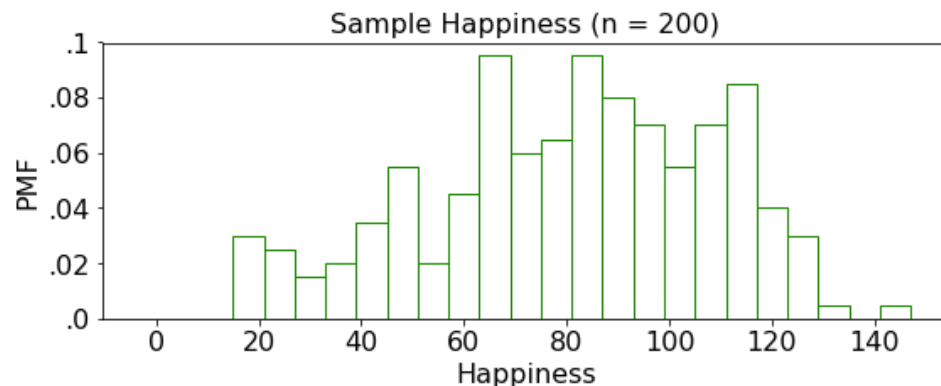
a. Resample **sample.size()** from PMF

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

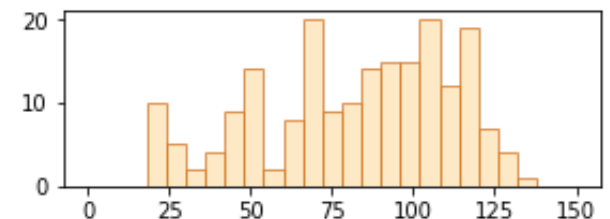
3. You now have a **distribution of your sample variance**

variances = [827.4]

Bootstrapped variance



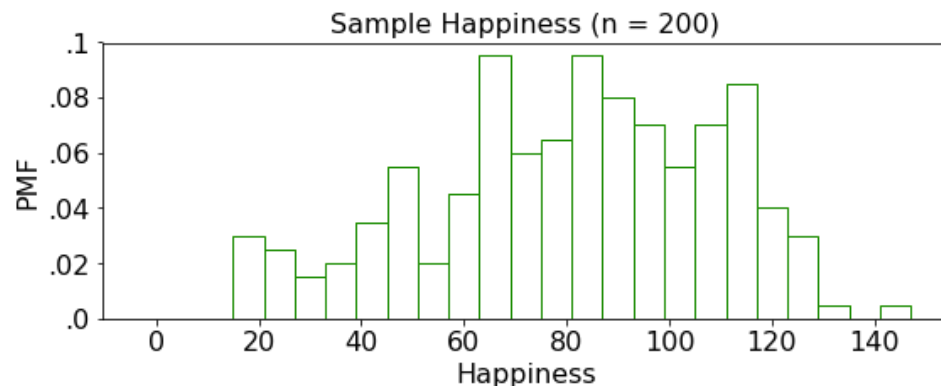
[116, 76, 132, 85, ..., 78]



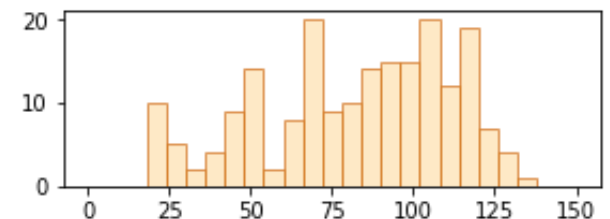
1. Estimate the **PMF** using the sample
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 - a. Resample **sample.size()** from PMF
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3. You now have a **distribution of your sample variance**

variances = [827.4]

Bootstrapped variance



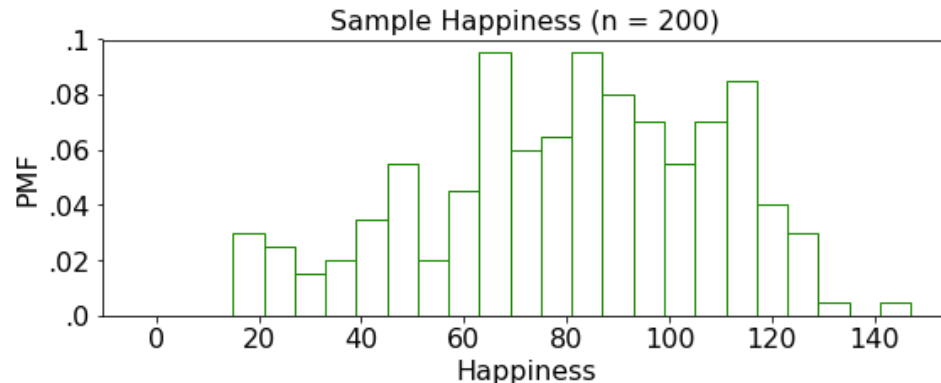
[116, 76, 132, 85, ..., 78]



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**

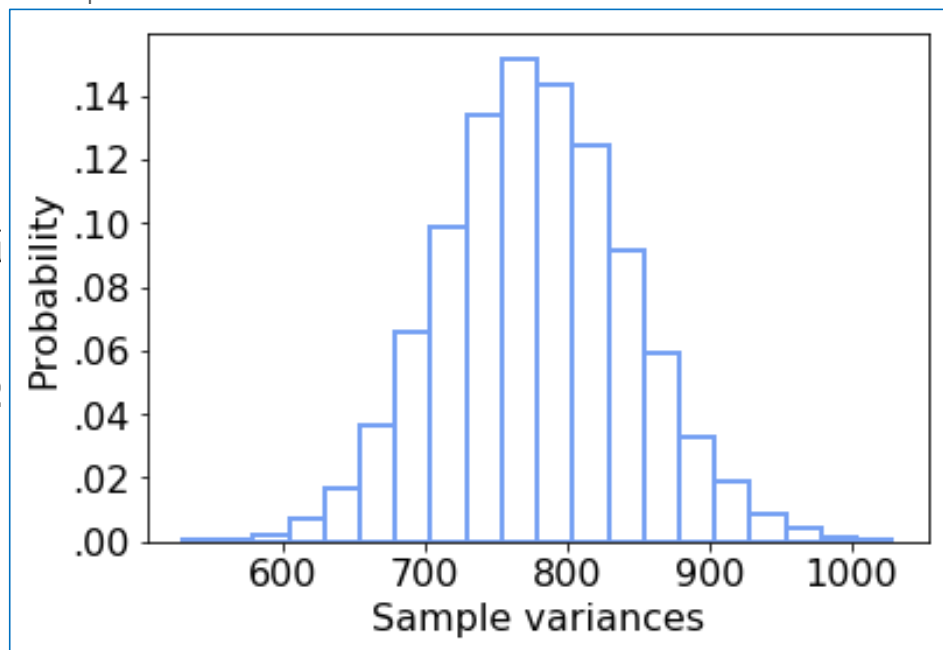
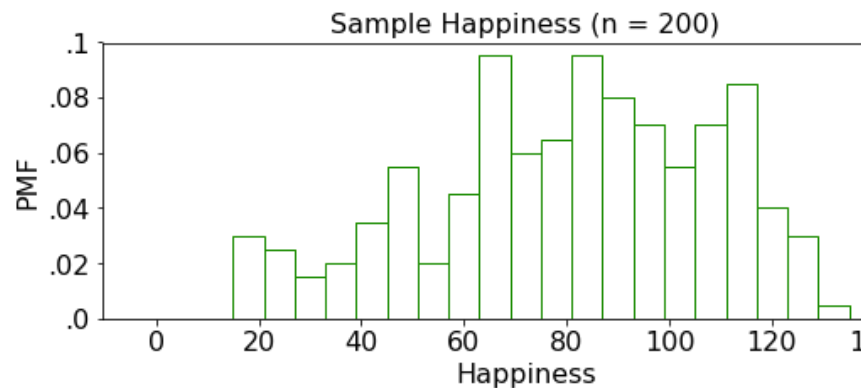
variances = [827.4, 846.1]

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

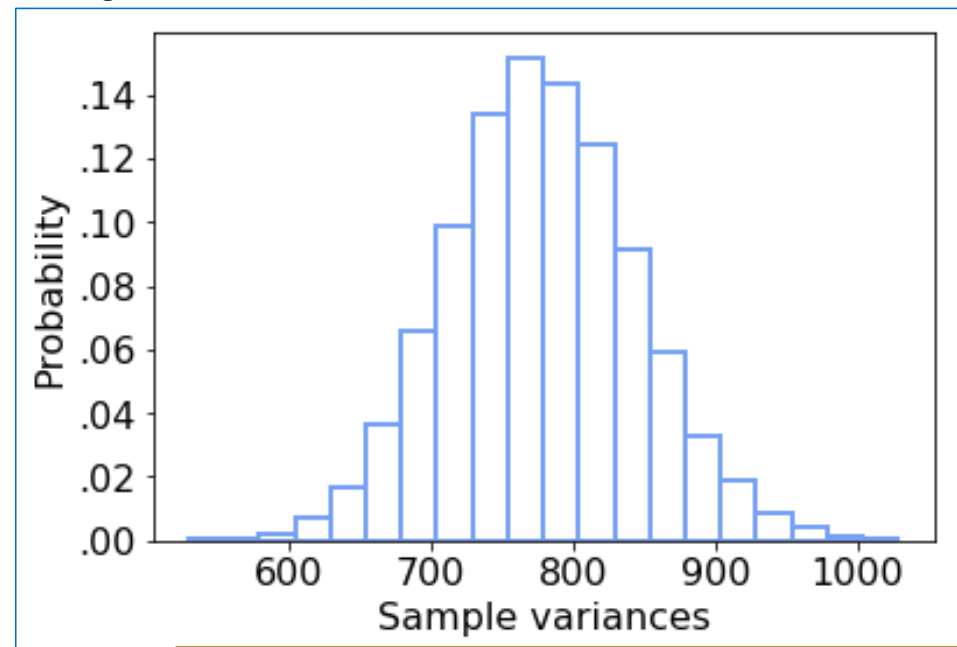
3. You now have a **distribution of your sample variance**

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

What is the bootstrapped standard error?

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

Bootstrapped standard error: 66.16



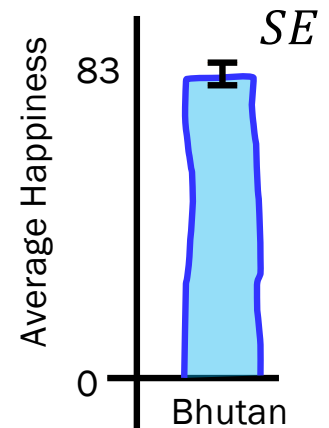
- Simulate a distribution of sample variances
- Compute standard deviation

Standard error

1. Mean happiness:

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

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



2. Variance of happiness:

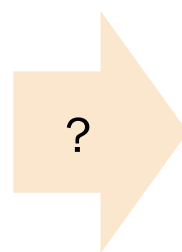
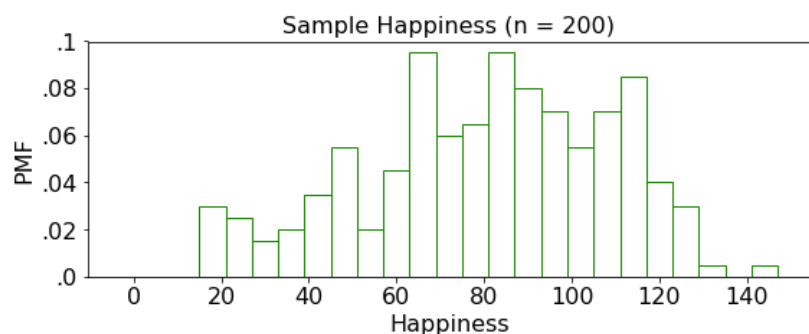
Claim: The variance of happiness of Bhutan is 793, with a **bootstrapped standard error of 66.16**.

S^2 is our best estimate of σ^2

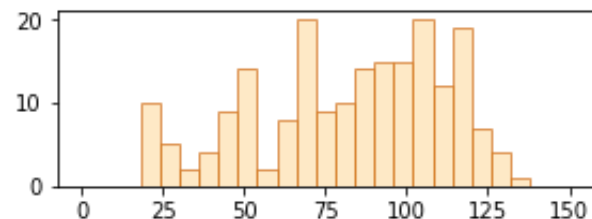
this is how close we are, calculated by bootstrapping

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**



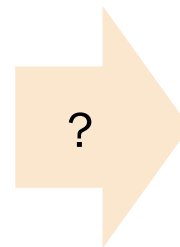
[116, 76, 132, 85, ..., 78]



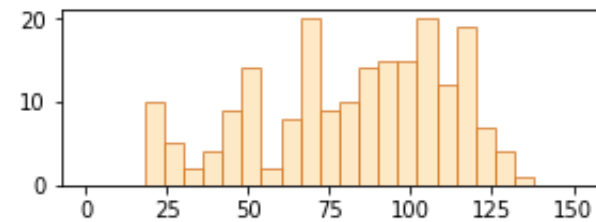
$$P(X = k) = \frac{\text{\# values in sample equal to } k}{n}$$

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.

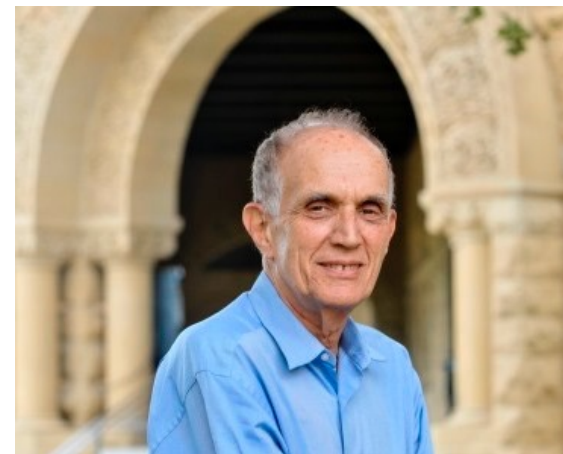
Bootstrapping works for any statistic*

*as long as your sample is iid and the underlying distribution does not have a long tail

Google colab notebook [link](#)

Bradley Efron

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



Inventor of Efron's dice: 4 dice A, B, C, D where:

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



Bootstrap: p-value



Null hypothesis test

Nepal
Happiness

4.45

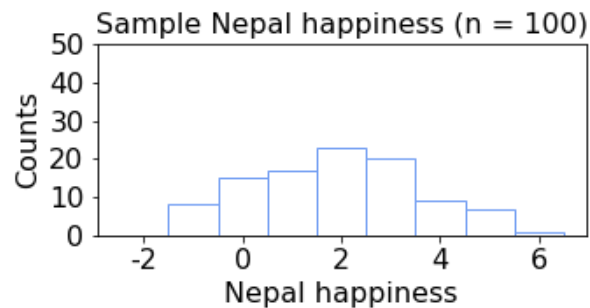
2.45

6.37

2.07

...

1.63



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

Bhutan
Happiness

0.91

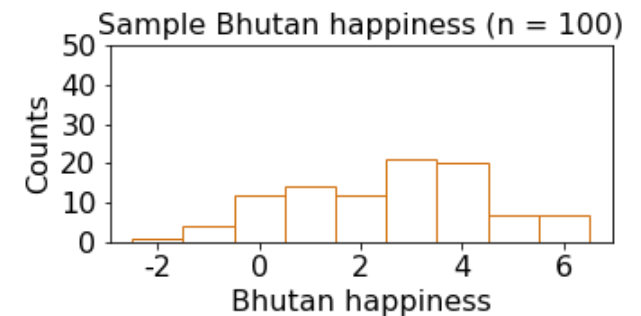
0.34

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 statistically 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

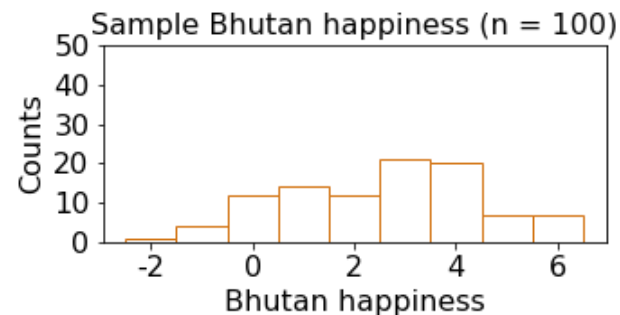
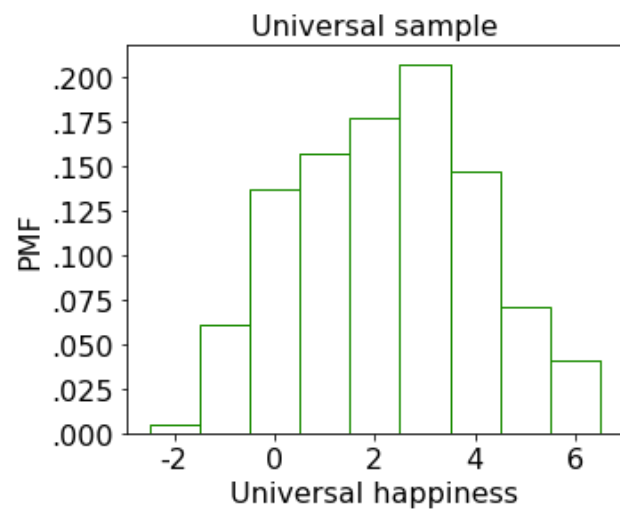
} **p-value**

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

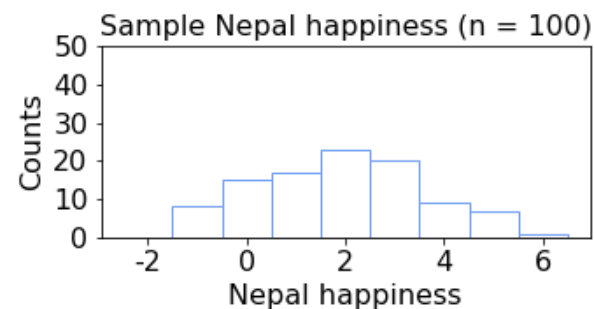
hypothesis.

Universal sample

(this is what the null hypothesis assumes)



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



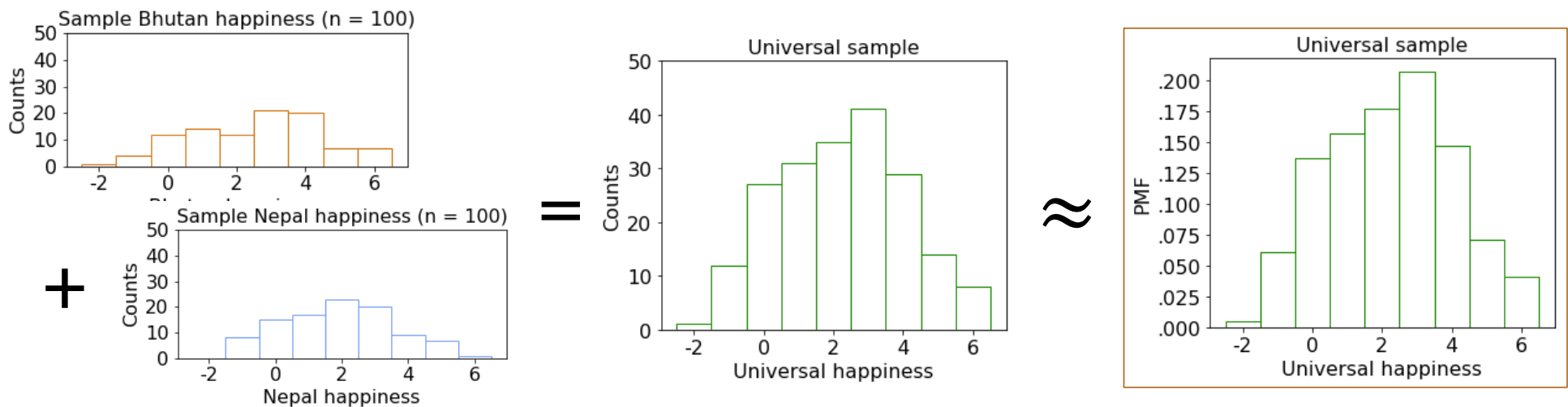
$$\bar{X}_2 = 2.4$$

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

Bootstrap for p-values

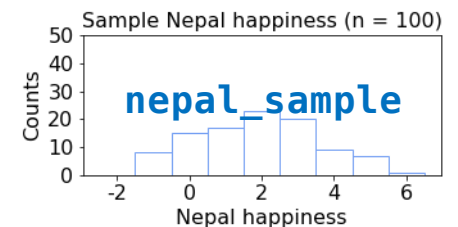
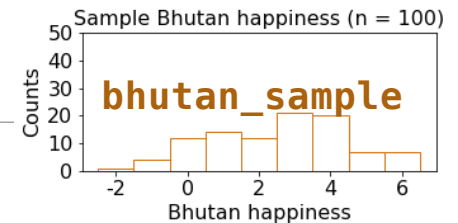
1. Create a **universal sample** using your two samples

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:  
        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

```
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
```

with replacement!

Bootstrap



Let's try it!

Google colab notebook [link](#)

Null hypothesis test

Nepal
Happiness

4.45

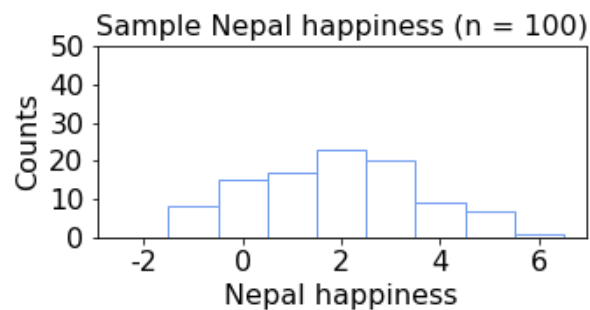
2.45

6.37

2.07

...

1.63



Bhutan
Happiness

0.91

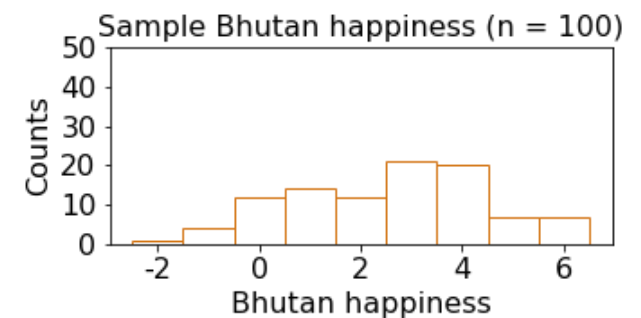
0.34

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 statistically significant ($p < 0.05$).