

# Expected values

Brian Caffo, Jeff Leek, Roger Peng

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# Expected values

- ▶ Expected values are useful for characterizing a distribution
- ▶ The mean is a characterization of its center
- ▶ The variance and standard deviation are characterizations of how spread out it is
- ▶ Our sample expected values (the sample mean and variance) will estimate the population versions

# The population mean

- ▶ The **expected value** or **mean** of a random variable is the center of its distribution
- ▶ For discrete random variable  $X$  with PMF  $p(x)$ , it is defined as follows

$$E[X] = \sum_x xp(x).$$

where the sum is taken over the possible values of  $x$

- ▶  $E[X]$  represents the center of mass of a collection of locations and weights,  $\{x, p(x)\}$

# The sample mean

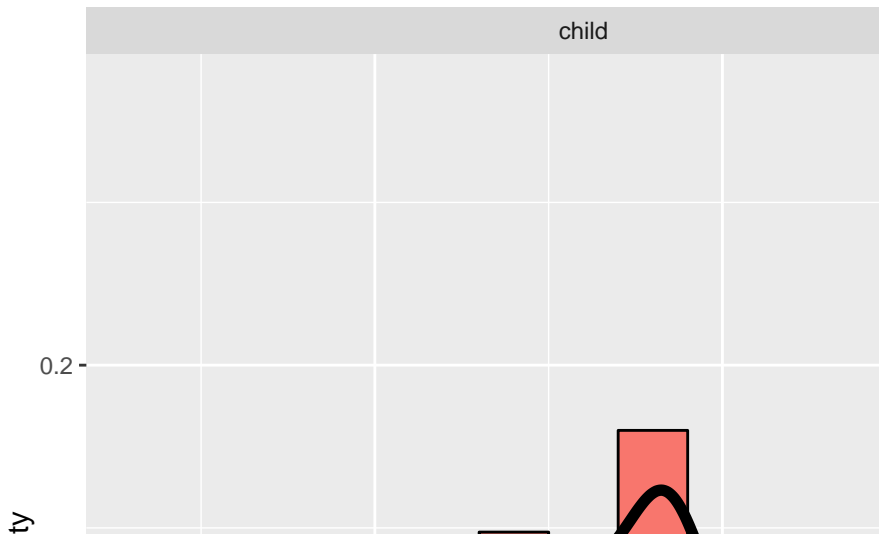
- ▶ The sample mean estimates this population mean
- ▶ The center of mass of the data is the empirical mean

$$\bar{X} = \sum_{i=1}^n x_i p(x_i)$$

where  $p(x_i) = 1/n$

## Example

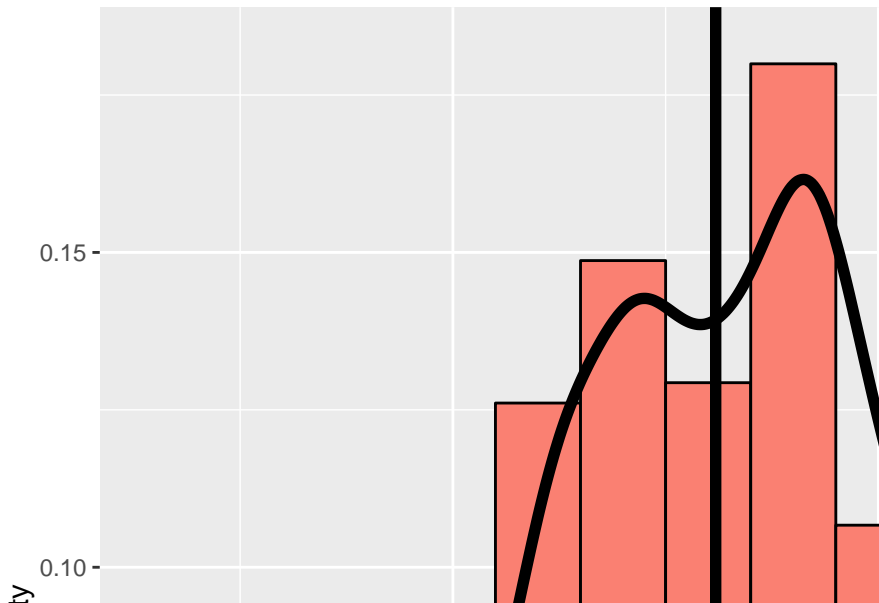
Find the center of mass of the bars



## Using manipulate

```
library(manipulate)
myHist <- function(mu){
  g <- ggplot(galton, aes(x = child))
  g <- g + geom_histogram(fill = "salmon",
    binwidth=1, aes(y = ..density..), colour = "black")
  g <- g + geom_density(size = 2)
  g <- g + geom_vline(xintercept = mu, size = 2)
  mse <- round(mean((galton$child - mu)^2), 3)
  g <- g + labs(title = paste('mu = ', mu, ' MSE = ', mse))
  g
}
manipulate(myHist(mu), mu = slider(62, 74, step = 0.5))
```

The center of mass is the empirical mean

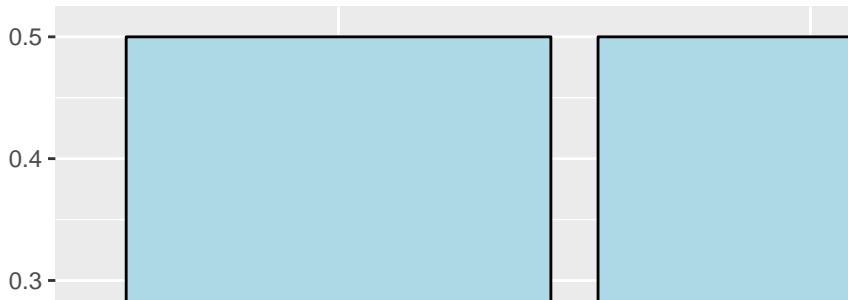


## Example of a population mean

- ▶ Suppose a coin is flipped and  $X$  is declared 0 or 1 corresponding to a head or a tail, respectively
- ▶ What is the expected value of  $X$ ?

$$E[X] = .5 \times 0 + .5 \times 1 = .5$$

- ▶ Note, if thought about geometrically, this answer is obvious; if two equal weights are spaced at 0 and 1, the center of mass will be .5





## What about a biased coin?

- ▶ Suppose that a random variable,  $X$ , is so that  $P(X = 1) = p$  and  $P(X = 0) = (1 - p)$
- ▶ (This is a biased coin when  $p \neq 0.5$ )
- ▶ What is its expected value?

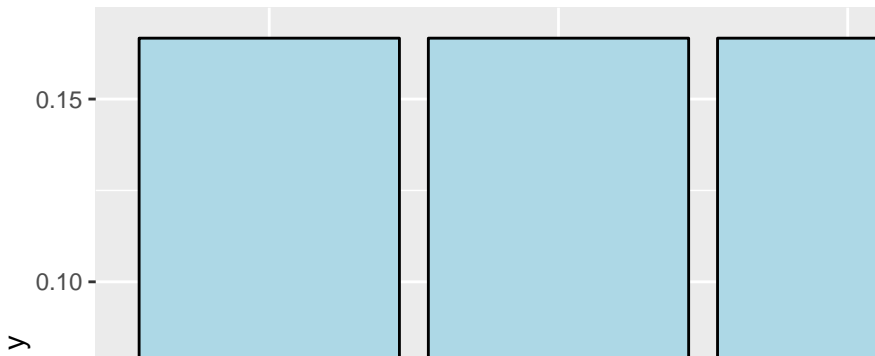
$$E[X] = 0 * (1 - p) + 1 * p = p$$

## Example

- ▶ Suppose that a die is rolled and  $X$  is the number face up
- ▶ What is the expected value of  $X$ ?

$$E[X] = 1 \times \frac{1}{6} + 2 \times \frac{1}{6} + 3 \times \frac{1}{6} + 4 \times \frac{1}{6} + 5 \times \frac{1}{6} + 6 \times \frac{1}{6} = 3.5$$

- ▶ Again, the geometric argument makes this answer obvious without calculation.

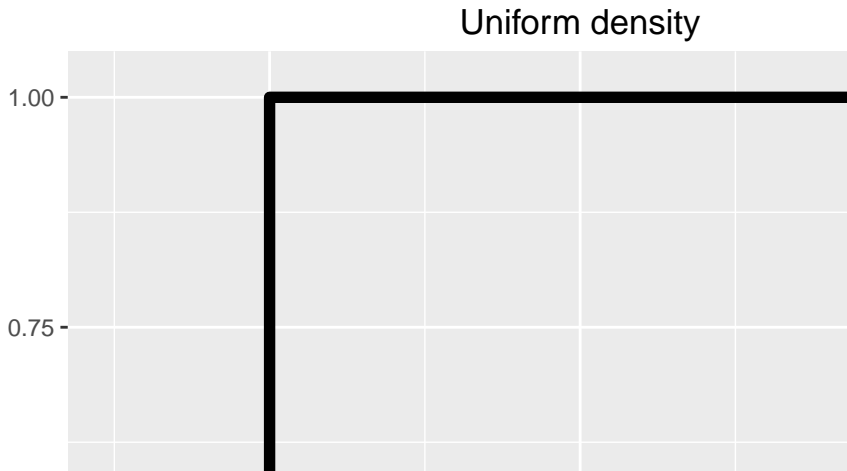


# Continuous random variables

- ▶ For a continuous random variable,  $X$ , with density,  $f$ , the expected value is again exactly the center of mass of the density

## Example

- ▶ Consider a density where  $f(x) = 1$  for  $x$  between zero and one
- ▶ (Is this a valid density?)
- ▶ Suppose that  $X$  follows this density; what is its expected value?

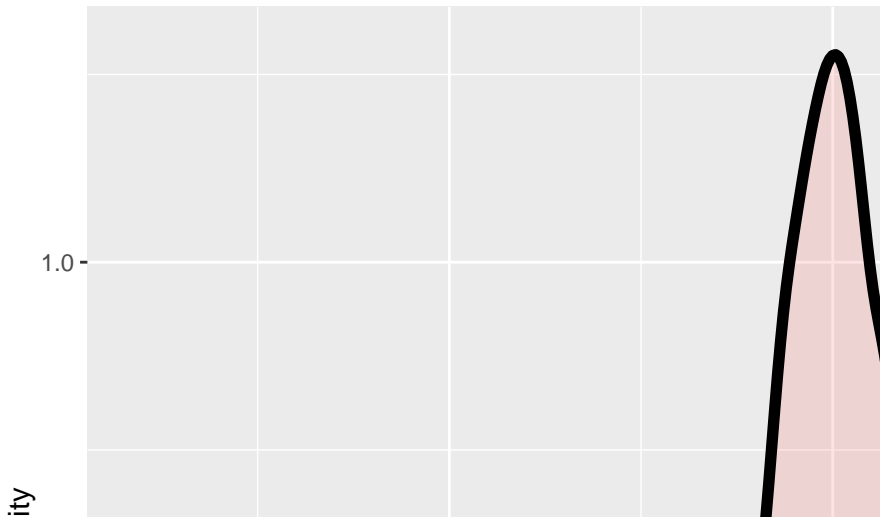


# Facts about expected values

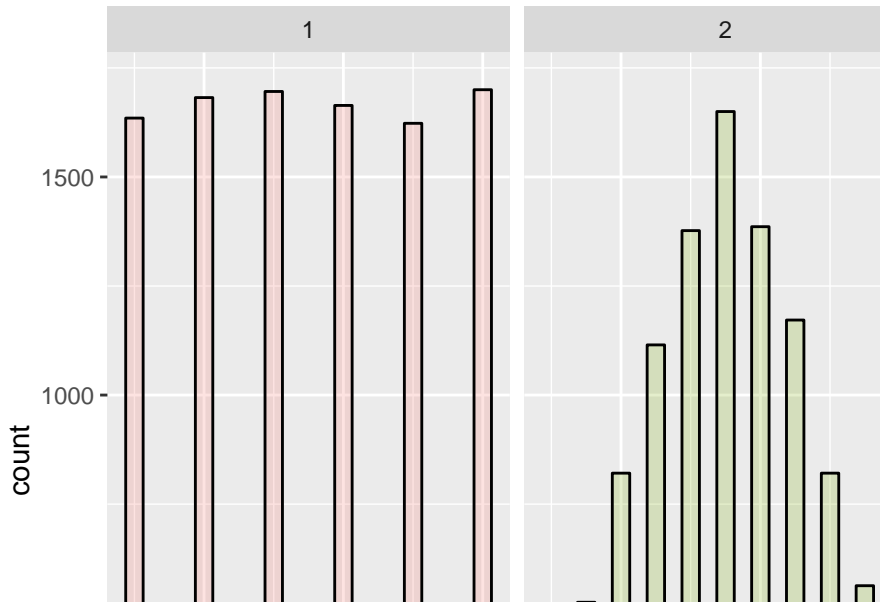
- ▶ Recall that expected values are properties of distributions
- ▶ Note the average of random variables is itself a random variable and its associated distribution has an expected value
- ▶ The center of this distribution is the same as that of the original distribution
- ▶ Therefore, the expected value of the **sample mean** is the population mean that it's trying to estimate
- ▶ When the expected value of an estimator is what its trying to estimate, we say that the estimator is **unbiased**
- ▶ Let's try a simulation experiment

## Simulation experiment

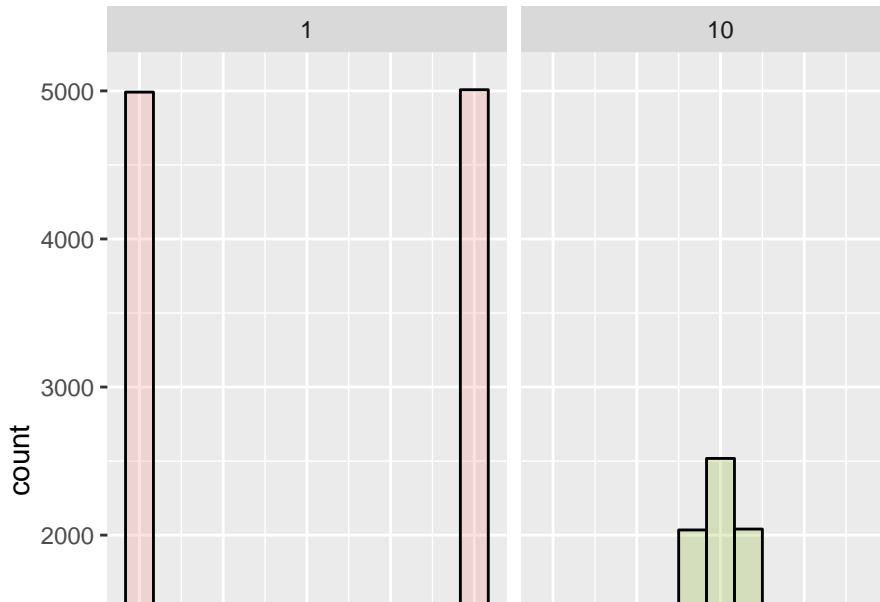
Simulating normals with mean 0 and variance 1 versus averages of 10 normals from the same population



## Averages of x die rolls



## Averages of $x$ coin flips





# Sumarizing what we know

- ▶ Expected values are properties of distributions
- ▶ The population mean is the center of mass of population
- ▶ The sample mean is the center of mass of the observed data
- ▶ The sample mean is an estimate of the population mean
- ▶ The sample mean is unbiased
- ▶ The population mean of its distribution is the mean that it's trying to estimate
- ▶ The more data that goes into the sample mean, the more concentrated its density / mass function is around the population mean