Inference about a Population Proportion (π) AAO unit 28: Baldi & Moore, Ch 19

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Proportion/Count in a Sample

Binomial Model for Sampling Variability of

It is the n + 1 probabilities p₀, p₁, ..., p_y, ..., p_n of observing 0, 1, 2, ..., n "positives" in n independent realizations of a Bernoulli random variable Y:

$$Y = \begin{cases} 1 & P(Y = 1) = \pi \\ 0 & P(Y = 0) = 1 - \pi \end{cases}$$

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The number is the sum of n i.i.d. Bernoulli random variables. (such as in SRS of n individuals)

- Each of the *n* observed elements is binary (0 or 1)
- There are 2^n possible sequences ... but only n+1 possible values, i.e. 0/n, 1/n, ..., n/n (can think of y as sum of n Bernoulli random variables)
- Note: it is better to work in same scale as the parameter, i.e., in [0,1]. Not the [0,n] count scale.

-

- Apart from (n), the probabilities p_0 to p_n depend on only 1 parameter:
 - the probability that a selected individual will be "positive" i.e.,
 - the proportion of "positive" individuals in sampled population
- lacksquare Usually denote this (un-knowable) proportion by π

Author	Parameter	Statistic
Clayton & Hills	π	p = D/N
Hanley et al.	π	p = y/n
M&M, Baldi & Moore	р	$\hat{p} = y/n$
Miettinen	Р	p = y/n

■ Shorthand: $Y \sim \text{Binomial}(n, \pi)$.

Example

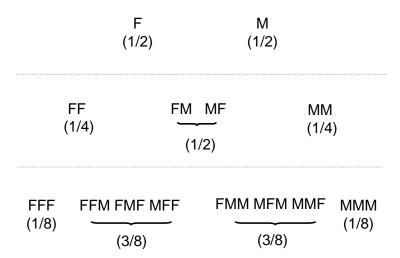
- Suppose a woman plans to have 3 children.
- Suppose at each birth,

$$P(\text{female child}) = 1/2$$

and the sex of the child at each birth is independent of the sex at any previous birth.

What is the probability of having all daughters?

The binomial distribution



The binomial distribution

Let Y be the number of daughters a woman will have, n the number of children she will have, and p the probability of a daughter at any birth. Then:

$$P(Y = k) = \frac{n!}{(n-k)!k!} p^k (1-p)^{(n-k)}$$

where $n! = 1 \times 2 \times 3 \times ... \times (n-1) \times n$, and 0! = 1.

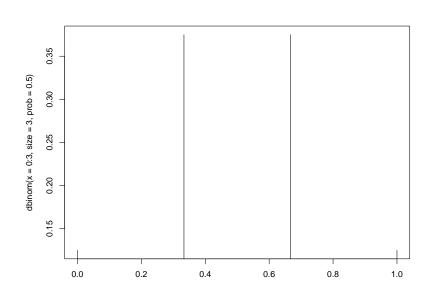
Calculating binomial probabilities in R

$$P(Y=3) = \frac{3!}{0!3!} 0.5^3 (1-0.5)^0$$
 which can be solved in R using: stats::dbinom(x = 3, size = 3, prob = 0.5)

[1] 0.125

The probability mass function (pmf)

```
plot(0:3/3, dbinom(x = 0:3, size = 3, prob = 0.5), type = "h")
```



What do we use it for?

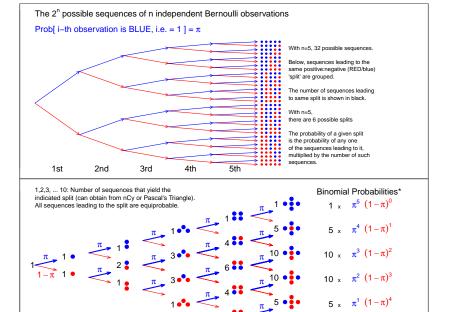
• to make inferences about π from observed proportion p = y/n.

What do we use it for?

- to make inferences about π from observed proportion p = y/n.
- to make inferences in more complex situations, e.g.
 - Prevalence Difference: $\pi_1 \pi_0$
 - ▶ Risk Difference (RD): $\pi_1 \pi_0$
 - **>** Risk Ratio, or its synonym Relative Risk (RR): π_1 / π_0
 - ▶ Odds Ratio (OR): $[\pi_1/(1-\pi_1)]/[\pi_0/(1-\pi_0)]$
 - ▶ Trend in several π 's

Requirements for y to have a Binomial (n, π) distribution

- 1. Fixed sample size *n*.
- 2. Elements selected at random (i.e. same probability of being sampled) and independent of each other;
- 3. Each element in "population" is 0 or 1, but we are only interested in estimating proportion (π) of 1's; we are not interested in individuals.
- 4. Denote by y_i the value of the *i*-th sampled element. $P(y_i = 1)$ is constant (it is π) across *i*.



 $1 \times \pi^0 (1-\pi)^5$

* in R: dbinom(0:5,size=5,prob=0.xx)

Does the Binomial Distribution Apply if...?

Interested in	π	the proportion of 16 year old girls in Québec protected against rubella
Choose	n = 100	girls: 20 at random from each of 5 randomly selected schools ['cluster' sample]
Count	у	how many of the $n=100$ are protected
• Is $y \sim \text{Binom}$	nial(n = 100)	$(0,\pi)$?
"SMAC"	π	P(abnormal Healthy) =0.03 for each chemistry in Auto-analyzer with $n=18$ channels
Count	у	How many of $n = 18$ give abnormal result.
• Is $y \sim \text{Binom}$	nial(n = 18,	$\pi=0.03$)? (cf. Ingelfinger: Clin. Biostatistics)

Does the Binomial Distribution Apply if...?

π_e expt'l. exercise classes who 'stay the co	urse'		
Randomly 4 classes of			
Allocate <u>25</u> students each to usual course			
$n_u = 100$			
4 classes of			
<u>25</u> students each to experimental course			
$n_e = 100$			
Count y_u how many of the $n_u = 100$ complete cou	ırse		
y_e how many of the $n_e=100$ complete cou			
• Is $y_u \sim \text{Binomial}(n_u = 100, \pi_u)$? Is $y_e \sim \text{Binomial}(n_e = 100, \pi_e)$?			

Does the Binomial Distribution Apply if...?

Sex Ratio	n = 4 y	children in each family number of girls in family			
• Is variation of y across families Binomial (n = 4, π = 0.49)?					
Pilot		To estimate proportion π of population that			
Study		is eligible & willing to participate in long-term			
		research study, keep recruiting until obtain			
	y = 5	who are. Have to approach <i>n</i> to get <i>y</i> .			
• Can we treat $y \sim \operatorname{Binomial}(n, \pi)$?					

Calculating Binomial probabilities - Exactly

- probability mass function (pmf): $P(Y = k) = \frac{n!}{(n-k)!k!} p^k (1-p)^{(n-k)}$
- in R: dbinom(), pbinom(), qbinom(): probability mass, distribution/cdf, and quantile functions.

Calculating Binomial probabilities - Using an approximation

- Poisson Distribution (n large; small π)
- Normal (Gaussian) Distribution (*n* large or midrange π) ¹
 - Have to specify scale. Say n = 10, whether summary is a r.v. e.g. E SD

count:
$$y$$
 2 $n \times \pi$ $\{n \times \pi \times (1-\pi)\}^{1/2}$ $n^{1/2} \times \sigma_{Bernoulli}$ proportion: $p = y/n$ 0.2 π $\{\pi \times (1-\pi)/n\}^{1/2}$ $\sigma_{Bernoulli}/n^{1/2}$

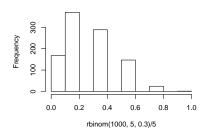
percentage: 100p% 20% $100 \times \pi$ $100 \times SD[p]$

 \blacktriangleright same core calculation for all 3 [only the *scale* changes]. JH prefers (0,1), the same scale as π .

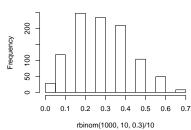
¹For when you don't have access to software or Tables, e.g, on a plane

Normal approximation to binomial is the CLT in action

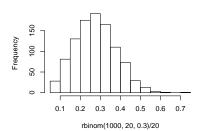




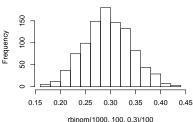
Histogram of rbinom(1000, 10, 0.3)/10



Histogram of rbinom(1000, 20, 0.3)/20

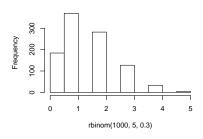


Histogram of rbinom(1000, 100, 0.3)/100

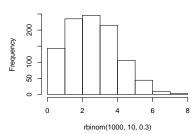


Normal approximation to binomial is the CLT in action

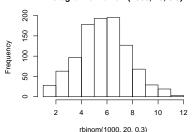




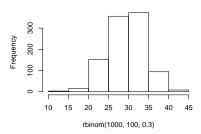
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Histogram of rbinom(1000, 100, 0.3)



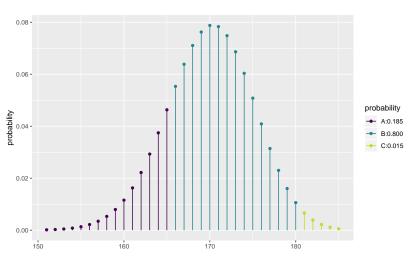
Example 1 from AAO Unit 21

A drug manufacturer claims that its flu vaccine is 85% effective; in other words, each person who is vaccinated stands an 85% chance of developing immunity. Suppose that 200 randomly selected people are vaccinated. Let Y be the number that develops immunity.

- 1. What is the distribution of Y?
- 2. What is the mean and standard deviation for Y?
- 3. What is the probability that between 165 and 180 of the 200 people who were vaccinated develop immunity? (Hint: Use a normal distribution to approximate the distribution of Y)

Example 1 from AAO Unit 21 - Exact Method

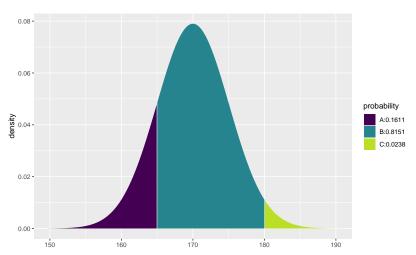
```
mosaic::xpbinom(q = c(165, 180), size = 200, prob = 0.85)
```



[1] 0.1850410 0.9851197

Example from AAO Unit 21- Normal Approximation

```
mosaic::xpnorm(q = c(165,180), mean = 200 * 0.85,
sd = sqrt(200*0.85*0.15))
```



[1] 0.1610510 0.9761648

Example 2 from AAO Unit 21

People with O- blood are called universal donors because most people can receive an O-blood transfusion. The probability of having blood type O- is 0.066. Suppose a random sample of five people show up during a blood drive to donate blood. Let Y be the number of people with blood type O-.

- 1. What is the probability that none of the five people has blood type O-?
- 2. What is the probability that exactly one of the five has blood type O-?
- 3. What is the probability that no more than one of the five people has blood type O-?
- 4. What is the probability that at least one of the five has blood type O-?

1. What is the probability that none of the five people has blood type O-?

$$P(Y=0) = {5 \choose 0} 0.066^0 (1 - 0.066)^5$$

```
stats::dbinom(x = 0, size = 5, prob = 0.066)
## [1] 0.7107787

(1-0.066)^5
## [1] 0.7107787
```

2. What is the probability that exactly one of the five has blood type O-?

$$P(Y=1) = {5 \choose 1} 0.066^{1} (1 - 0.066)^{4}$$

```
stats::dbinom(x = 1, size = 5, prob = 0.066)
## [1] 0.2511316
```

3. What is the probability that no more than one of the five people has blood type O-?

$$P(Y \le 1) = P(Y = 0) + P(Y = 1)$$

$$= \binom{5}{0} 0.066^{0} (1 - 0.066)^{5} + \binom{5}{1} 0.066^{1} (1 - 0.066)^{4}$$

$$stats:: \frac{\text{dbinom}(x = 0, \text{ size = 5, prob = 0.066})}{\text{stats::dbinom}(x = 1, \text{ size = 5, prob = 0.066})}$$
[1] 0.9619103
$$stats:: \frac{\text{pbinom}(q = 1, \text{ size = 5, prob = 0.066})}{\text{## [1] 0.9619103}}$$

$$mosaic:: xpbinom(q = 1, \text{ size = 5, prob = 0.066})$$

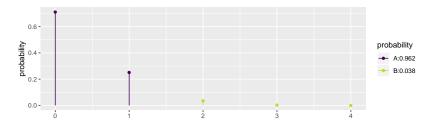
$$\sum_{\substack{\text{probability = 0.4 \\ \text{odd} 0.2 \\ \text{odd} 0.038}$$

4. What is the probability that more than one of the five has blood type O-?

$$P(Y > 1) = P(Y = 2) + P(Y = 3) + P(Y = 4) + P(Y = 5)$$

= 1 - P(Y \le 1)

```
1 - stats::pbinom(q = 1, size = 5, prob = 0.066)
## [1] 0.03808969
mosaic::xpbinom(q = 1, size = 5, prob = 0.066, lower.tail = FALSE)
```



[1] 0.03808969

Inference concerning a proportion π , based on s.r.s. of size n

Examples

The **Parameter** π of interest: the proportion ...

- with undiagnosed hypertension / seeing MD during a 1-year span
- who would respond to a specific therapy
- still breast-feeding at 6 months
- of pairs where response on treatment > response on placebo
- of Earth's surface covered by water
- who would enrol in a long-term study or answer a questionnaire
- of twin pairs where left-handed twin dies first
- able to tell imported from domestic beer in a "triangle taste test"

Inference via **Statistic**: the number (y) or proportion p = y/n 'positive' in an s.r.s. of size n.

Frequentist vs. Bayesian Inference

Frequentist (§2.1)

- based on prob[data $|\theta$], i.e.
- probability statements about data

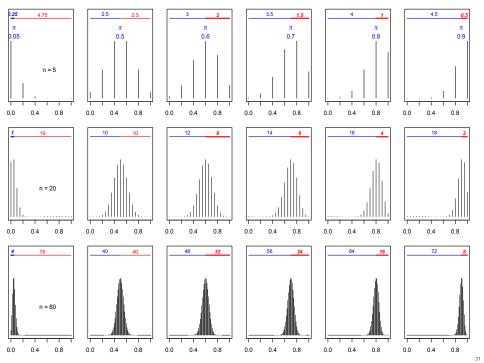
Evidence (P-value) against H_0 : $\pi = \pi_0$

Test of H_0 : Is P-value < (preset) α ?

CI: interval estmate

Bayesian ($\S 2.2$)

- based on prob[θ |data], i.e.,
- probability statements about π
- point estimate: mean/median/mode of posterior distribution of π
- (credible) interval



Justification for the $n \times \pi$ and $n \times (1 - \pi) \ge 10$

notes on the Figure from the previous slide:

- Binomial distributions, on (0,1) scale (rather than 0:n). Bigger expected numbers of 'positives' and 'negatives' imply less probability mass at the extreme(s) and thus help to approximate the (binomial) sampling distribution by a Gaussian distribution with mean π and $\sigma = \frac{\{\pi(1-\pi)\}^{1/2}}{\sqrt{n}}$.
- The amount of space needed at each extreme in order to accommodate a Gaussian distribution that does not spill over beyond the (0,1) boundaries is just another way to explain the ('taught but not explained') rule-of-thumb that the expected numbers, $n \times \pi$ and $n \times (1 \pi)$ should should exceed 10 (or 5, or 8, depending on the textbook, and the edition!)

Frequentist Confidence Interval for π , based on an observed proportion p = y/n stats::binom.test and mosaic::binom.test in R

It is sad that even today, with more emphasis on Cl's and less on p-values and tests, we have to go through the '.test' to get to the Cl. It is also of note that the procedure mentions the model (binomial) rather than the target parameter, the proportion π .

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- The base stats::binom.test function in R has just one method, the Clopper-Pearson one. The mosaic::binom.test one has it and four others, and these allow us to appreciate why different ones might be used in different circumstances. We will start with the most familiar of them, the so-called 'Wald' CI, which, because of its 'point estimate ±Margin.Of.Error' form, is symmetric.

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- The mosaic::binom.test allows for a vector of individual 0's and 1's, rather than the tallies of 1's and 0's required for stats::binom.test
- In practice, CIs for proportions, and functions thereof, will come from regression models.

CI based on Gaussian approximation to sampling distribution of the sample proportion p – the 'Wald' method in mosaic::binom.test

■ The Wald CI has been taught as having the form:

$$p \pm z^\star \times \mathrm{SE}[p]$$

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If the population sampled from has an (unknown) proportion π of 1's and an (unknown) proportion $1-\pi$ of 0's, then the theoretical SD of all of the 1's and 0's sampled from is $\sigma_{0/1} = \sqrt{\pi(1-\pi)}$.

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■ The Wald CI has been taught as having the form:

$$p \pm z^* \times SE[p]$$

- If the population sampled from has an (unknown) proportion π of 1's and an (unknown) proportion $1-\pi$ of 0's, then the theoretical SD of all of the 1's and 0's sampled from is $\sigma_{0/1} = \sqrt{\pi(1-\pi)}$.
- Since we don't know the true value of $\sigma_{0/1} = \sqrt{\pi(1-\pi)}$, we replace it with a version where we <u>substitute</u> p for π , i.e. the estimated SD of all of the 1's and 0's sampled from is $\widehat{\sigma_{0/1}} = \sqrt{p(1-p)}$.

CI based on Normal approximation to sampling distribution of the sample proportion p – the 'Wald' method in mosaic::binom.test

■ Dividing this $\widehat{\sigma_{0/1}}$ by the square root of n, we get the standard error, our best estimate of the spread of the sampling distribution of a sample proportion, i.e.,

$$SE[p] = \frac{\sqrt{p(1-p)}}{\sqrt{n}} = \frac{\widehat{\sigma_{0/1}}}{\sqrt{n}}.$$

CI based on Normal approximation to sampling distribution of the sample proportion p – the 'Wald' method in mosaic::binom.test

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$$p \pm z^* \times \sqrt{\frac{p(1-p)}{n}}.$$

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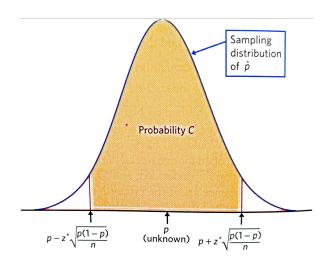
■ So, as it is traditionally presented, the CI becomes

$$p \pm z^{\star} \times \sqrt{\frac{p(1-p)}{n}}.$$

 As we will see below, now that we seldom calculate a CI 'from scratch,' today the Wald CI is better presented in the R-computational form

qnorm(p=c(0.025,0.975), mean= p, sd = sqrt(p*(1-p))/sqrt(n)).

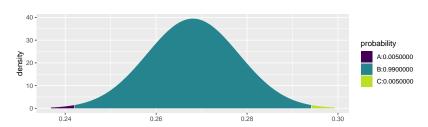
Sampling distribution of p



Example 1: Assessing the prevalence of HPV infections

NHANES found that 515 of a sample of 1921 women aged 14 to 59 years currently tested positive for HPV. Provide a 99% confidence interval for HPV prevalence.

```
n <- 1921
number_infected <- 515
p <- number_infected / n
s <- sqrt(p * (1 - p))
SEP <- s / sqrt(n)
mosaic::xqnorm(p=c(0.005,0.995), mean = p, sd = SEP)</pre>
```



[1] 0.2420567 0.2941224

Example 2: Assessing the prevalence of HPV infections

```
mosaic::binom.test(x = 515, n = 1921, ci.method=c("wald"), conf.level=0.99)

##

## ^^IExact binomial test (with Wald CI)

## data: 515 out of 1921

## number of successes = 515, number of trials = 1921, p-value <
## 2.2e-16

## 2.2e-16

## 3lternative hypothesis: true probability of success is not equal to 0.5

## 99 percent confidence interval:

## 0.2420567 0.2941224

## sample estimates:

## probability of success

## 0.2680895</pre>
```

Note: HPV positive ≠ 'success' !!

Suppose our observed proportion of 'water' locations was p=4/5, or 80%.

```
mosaic::binom.test(x = 4, n = 5, ci.method=c("wald"), conf.level=0.95)

##

## ^^IExact binomial test (with Wald CI)

## data: 4 out of 5

## number of successes = 4, number of trials = 5, p-value = 0.375

## alternative hypothesis: true probability of success is not equal to 0.5

## 95 percent confidence interval:

## 0.449391 1.150609

## sample estimates:

## probability of success

## 0.8
```

Suppose our observed proportion of 'water' locations was p=4/5, or 80%.

```
mosaic::binom.test(x = 4, n = 5, ci.method=c("wald"), conf.level=0.95)

##
## ^^IExact binomial test (with Wald CI)
##
## data: 4 out of 5
## number of successes = 4, number of trials = 5, p-value = 0.375
## alternative hypothesis: true probability of success is not equal to 0.5
## 95 percent confidence interval:
## 0.449391 1.150609
## sample estimates:
## probability of success
## 0.8
```

Clearly the proportion or percentage of the Earth's surface covered by water cannot be <u>1.15</u> or <u>115</u>%.

```
stats::qnorm(p=c(0.025,0.975), mean = 0, sd = sqrt(0 * 1 / 5))
## [1] 0 0
stats::qnorm(p=c(0.025,0.975), mean = 0.2, sd = sqrt(0.2 * 0.8 / 5))
## [1] -0.150609 0.550609
stats::qnorm(p=c(0.025,0.975), mean = 0.4, sd = sqrt(0.4 * 0.6 / 5))
## [1] -0.02940659 0.82940659
stats::qnorm(p=c(0.025,0.975), mean = 0.6, sd = sqrt(0.6 * 0.4 / 5))
## [1] 0.1705934 1.0294066
stats::qnorm(p=c(0.025,0.975), mean = 0.8, sd = sqrt(0.8 * 0.2 / 5))
## [1] 0.449391 1.150609
stats::qnorm(p=c(0.025,0.975), mean = 1, sd = sqrt(1 * 0 / 5))
## [1] 1 1
```

Thus, whatever your result, the Wald 95% CI gives a nonsensical result. Using the Normal/Gaussian approximation to the Binomial sampling distribution does not work when n = 5.

■ Answer: use a non-symmetric one, and one that respects the (0,1) scale.

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- The other 4 methods in mosaic::binom.test do respect the (0,1) scale

- Answer: use a non-symmetric one, and one that respects the (0,1) scale.
- The other 4 methods in mosaic::binom.test do respect the (0,1) scale
- We can also switch to the $(-\infty, \infty)$ logit scale, computing the CI in this scale, and then back-transforming to the (0,1) scale \rightarrow logistic regression.

1. Asymmetric (Wilson and Clopper-Pearson) Methods

content...

