High-Throughput Sequencing Course Count Models for RNA-Seq

Biostatistics and Bioinformatics



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TWO APPROACHES FOR ANALYSIS OF RNA-SEQ

- ► Two-stage method: Convert counts to "Expression" and then use statistical methods for microarrays (e.g., t-test)
- ► One-stage method: Relate the counts directly to the phenotype
- ► This is done through using statistical methods for modeling counts
- ▶ We generally promote the latter approach for data analysis

DESEQ FOR RNA-SEQ

- ► The goal is to provide sufficient background to understand the DESeq method
- \blacktriangleright We are not suggesting that DESeq is the best approach for analysis of RNA-Seq data
- ► We are considering it in this course as one, of many other methods, that adhere to the one-stage approach principle
- ► Added bonus: Nicely written R extension package (important feature for teaching)
- ► DESeq has many limitations (e.g., it cannot directly deal with quantitative and censored outcomes)
- ► Also some of the theoretical details (e.g., the effect of using plugin estimates for nuisance parameters) have seemingly not been fully fleshed out

THREE DISTRIBUTIONS FOR COUNT DATA

- ► RNA-Seq data are counts (not continuous measurements)
- ► To properly model RNA-Seq data, we need to consider distributions to model counts
- ▶ We will consider three important distributions for counts:
 - ► Binomial
 - ► Poisson
 - ► Negative Binomial
- ► There are many other distributions for counts (e.g., geometric distribution) that will not be discussed
- ▶ Brief notes on multinomial distribution

DISTRIBUTION FOR COUNTS: SUPPORT

- ► A count is a non-negative (zero or positive) integer
- ► When considering a distribution of a count variable, we first have to determine its *support*
- ► The support of the distribution consists of the values that could occur with positive probability
- ▶ For example, if we toss a coin once and we count the number of heads, the support is $\{0,1\}$
- ▶ If we flip it twice, the support is $\{0, 1, 2\}$
- ▶ Why is 3 not in the support? How about -1?
- ightharpoonup These values are not possible (they have zero probability)
- ► The probability to observed three heads among two tosses is zero.

DISTRIBUTION FOR COUNTS: PROBABILITY MASS FUNCTION

 \blacktriangleright Example: we toss a fair coin once and we count the number of heads (call it K)

$$P(K=0) = \frac{1}{2} \text{ and } P(K=1) = \frac{1}{2}$$

and

$$P(K=k) = 0$$

if k is not 0 or 1

- ▶ The probability mass function (PMF) determines the probability that K assumes value k in the support
- ► Sometimes we use the terms "distribution" and "PMF" interchangeably

DISTRIBUTION FOR COUNTS: PROBABILITY MASS FUNCTION

ightharpoonup Example: we toss a fair coin twice and we count the number of heads (call it K)

$$P(K = 0) = \frac{1}{4}$$
 and $P(K = 1) = \frac{1}{2}$ and $P(K = 2) = \frac{1}{4}$

- ► Why?
- ▶ Note that if once adds up P(K = k) over all k in the support the sum should be one

$$\sum_{k} P(K = k) = 1$$

EXERCISE: SUPPORT AND PMF

- \blacktriangleright we toss a biased coin twice and we count the number of heads (call it K)
- ▶ the probability that any toss lands a head is $\pi = \frac{1}{3}$
- ▶ What is the support of the distribution
- ► What is the PMF
- ▶ Repeat the last steps if π is any arbitrary number (between 0 and 1 of course)

EXERCISE: SUPPORT AND PMF

- the support is as in the previous example $\{0, 1, 2\}$
- ► Why is it unchanged

$$P(K = 0) = \frac{4}{9}$$
 and $P(K = 1) = \frac{4}{9}$ and $P(K = 2) = \frac{1}{9}$

► More generally

$$P(K = 0) = (1 - \pi)^{2}$$
$$P(K = 1) = 2\pi(1 - \pi)$$

$$P(K=2)=\pi^2$$

FLIPPING THE COIN

- ► Throughout this discussing we will consider flipping a coin
- ▶ The coin lands a head with probability π (could be biased) or tail with probability $1-\pi$
- ▶ For convenience, we will recode H as 1 and T as 0
- \blacktriangleright We will flip it n times.
- ► Notation:
 - ightharpoonup n is to denote the number of trials
 - ► On any trial (or flip), if we land an H we will call it an event (or success)
 - ▶ or if we land a T we will call it a failure
- ► RNA-seq connection: You can think of a read mapping to a gene to be an event

THREE VARIANTS OF THE COIN TOSSING EXPERIMENT

- 1. Fix the number of trials (n) upfront and then toss the coin n times
 - ightharpoonup The number of events (among n trials) is random
- Toss the coin a large number of times and assume that each one of these many trials has a small probability of being an event
 - ▶ Here n is large and π is small (close to 0)
- 3. Fix the number of desired events upfront, then toss the coin repeatedly to achieve that number ${\bf r}$
 - \blacktriangleright Here the number of trials n is random

OUTLINE FOR TODAY (AND MAYBE THUR)

- Provide an overview of the properties of the three distributions
- ► PMF. Mean and Variance
- ▶ Discuss relationship between the three distributions
- ► Need to introduce some notation (unfortunately)
- ► The goal is develop a regression model for counts
- \blacktriangleright We motivate this first using linear regression
- \blacktriangleright And then through logistic regression
- ▶ Before moving on to dicussing a regression model based on negative binomial distribution
- ▶ Provide some insight on how these models are estimated

Example: Fixed n

- ▶ We flip the coin n = 6 times
- ▶ Observed sequence: TTHTTH
- ▶ We recode this as 001001
- ► This corresponds to
 - ightharpoonup n = 6 trials
 - ► 2 events (or successes)
 - ▶ or equivalently 4 failures

Number of Possible Outcomes

- ▶ Example 1: Suppose that n = 2
 - \blacktriangleright 4 possible outcomes: $\{00,10,01,11\}$
 - $4 = 2 \times 2 = 2^2$
- ▶ Example 2: Suppose that n = 3
 - ► Eight possible outcomes: {000, 100, 101, 001, 110, 011, 101, 111}
 - $8 = 2 \times 2 \times = 2^3$
 - Example 3: n = 6
 - $64 = 2^6$ outcomes
- ▶ The number of possible outcomes based on n trials is 2^n
- \blacktriangleright But we are not interested in counting outcomes
- ▶ We want to count the number of outcomes corresponding to K=0, K=1,...,K=n

Number of Successes

- ▶ Example 1: Suppose that n = 2
 - \blacktriangleright 4 possible outcomes: $\{00,10,01,11\}$
 - ▶ Number outcomes corresponding to K = 0 is 1
 - ▶ Number outcomes corresponding to K = 1 is 2
 - ▶ Number outcomes corresponding to K = 2 is 1
- \blacktriangleright Example 2: Suppose that n=3
 - ► Eight possible outcomes: {000, 100, 010, 001, 110, 011, 101, 111}
 - ▶ Number outcomes corresponding to K = 0 is 1
 - ▶ Number outcomes corresponding to K = 1 is 3
 - Number outcomes corresponding to K = 2 is 3
 - ▶ Number outcomes corresponding to K = 3 is 1
- \blacktriangleright What does this look like for a general n?
- \blacktriangleright If you toss the coin n times, how many outcomes correspond to k events?

FACTORIAL FUNCTION

- ▶ Integers are "whole" numbers ..., -2, -1, 0, 1, 2, ...
- \blacktriangleright Consider a non-negative integer k (0, 1, 2, ...)
- 0! = 1
- **▶** 1! = 1
- $ightharpoonup 2! = 2 \times 1 = 2$
- ▶ $3! = 3 \times 2 \times 1 = 6$
- $ightharpoonup 4! = 4 \times 3 \times 2 = 24$
- **.** . .
- $\blacktriangleright k! = k \times (k-1) \times (k-2) \times \dots \times \times \times \times 1$

NUMBER OF COMBINATIONS

ightharpoonup The number of possible combinations on the basis of k events among n trials

$$\binom{n}{k} = \frac{n!}{k!(n-k)!}$$

▶ Example 1: Suppose that n = 3 and k = 1

$$\binom{3}{1} = \frac{3!}{1!(2-1)!} = \frac{3 \times 2 \times 1}{1 \times 2 \times 1} = 3$$

choose(3, 1)

▶ Example 2: Suppose that n = 4 and k = 2

$$\binom{4}{2} = \frac{4!}{2!(4-2)!} = \frac{4 \times 3 \times 2 \times 1}{2 \times 1 \times 2 \times 1} = \frac{24}{4} = 6$$

choose(4, 2)
[1] 6

Toss the coin n times

- ightharpoonup Toss the coin n times
- ▶ Number of possible outcomes: 2^n
- ▶ Number outcomes corresponding to K = 0 is 1.
- ▶ Number outcomes corresponding to K = 1 is n.
- ▶ Number outcomes corresponding to K = n 1 is n.
- ▶ Number outcomes corresponding to K = n is 1.
- ▶ Number outcomes corresponding to K = k is

$$\binom{n}{k} = \frac{n!}{k!(n-k)!}$$

for $k = 0, 1, 2, \dots$

- ▶ Do the results for K = 0, 1, n 1, K = n agree with the formula?
- ▶ Related to the Pascal Triangle

PASCAL TRIANGLE

BERNOULLI DISTRIBUTION

- ► Suppose that we toss the coin just once
- ▶ In other words n = 1
- ▶ We say that the number of events follows a Bernoulli distribution with parameter π
- ► The PMF is

$$P(K = k) = \pi^k (1 - \pi)^{1-k}, k = 0, 1$$

```
set.seed(12324)
# Simulate 10 Bernoulli random variables with parameter pi=0.5
rbinom(10, 1, 0.5)
## [1] 1 1 1 1 1 1 0 0 0 0 0
# Simulate 5 Bernoulli random variables with parameter pi=0.23
rbinom(5, 1, 0.23)
## [1] 0 0 0 0 0
```

BINOMIAL DISTRIBUTION

- ▶ For the Bernoulli distribution n = 1
- ▶ More generally (when $n \ge 1$) the number of events K is said to follow a Binomial distribution with parameters n and π
- \blacktriangleright The distribution is

$$P[K=k] = \binom{n}{k} \pi^k (1-\pi)^{n-k},$$

 $k = 0, 1, 2, \dots, n$

- ▶ Note that when n = 1 the Binomial reduces to a Bernoulli distribution . Why?
- ▶ Why is does this distribution have $\binom{n}{k}$?
- ▶ The average count for this distribution is $n\pi$
- ▶ The variance for this distribution is $n\pi(1-\pi)$

```
set.seed(12324)
# Simulate 10 Binomial random variables with parameter n=2 and pi=0.5
rbinom(10, 2, 0.5)
## [1] 1 2 2 1 2 0 0 1 1 1
```

Poisson Distribution

- ► The Poisson distribution is used to model the count of the occurrence of events
- ▶ Classical application: Model for earthquakes
- ► The PMF is

$$P(K = k) = \frac{e^{-\lambda} \lambda^k}{k!},$$

where k = 0, 1, 2, ...

- \triangleright λ is the average number of events for this distribution
- \triangleright λ is also the variance of this distribution

```
set.seed(13224)
# Simulate 10 Poisson variates with m
rpois(10, 0.1)
## [1] 0 1 0 0 0 0 1 0 0 0
```

RELATIONSHIP BETWEEN BINOMIAL AND POISSON DISTRIBUTION

- ► Consider tossing the coin a large number of times

 n = 1e+06
- ▶ Note that we have $n = 10^6$ trials with a low success probability of $p = 10^{-6}$
- ► The expected number of events among these 10^6 trials is $n \times p = 1$. Why?
- Now simulate 99999 numbers from this binomial distribution

```
set.sed(9988)
x <- rbinom(B9, n, p)
length(x)
## [1] 99999
```

▶ What is the expected number of events (i.e., the expected number of events (among n trials) across B = 99999 simulations)?

```
mean(x)
## [1] 1.00055
```

RELATIONSHIP BETWEEN BINOMIAL AND POISSON DISTRIBUTION

► Now compare the empirical distributions to the Poisson distributions

```
round(dpois(0:7, lambda = 1), 3)
## [1] 0.368 0.368 0.184 0.061 0.015 0.003 0.001 0.000
round(table(x)/B9, 3)
## x
## 0 1 2 3 4 5 6 7
## 0.367 0.369 0.183 0.061 0.016 0.003 0.000 0.000
```

NEGATIVE BINOMIAL DISTRIBUTION

- ▶ How many times do you have to flip a coin to get r > 0 events
- \blacktriangleright Model the number of random trials needed to get r events
- ► This distribution is called the negative binomial distribution
- ▶ The probability distribution is

$$P[K = k] = \binom{k+r-1}{r-1} \pi^r (1-\pi)^k,$$

where k = r, r + 1, r + 2, ...

set.seed(13224)
Simulate the number of trials needed to get k=5 events
rnbinom(10, 5, 0.1)
[1] 63 60 56 30 64 62 36 36 44 37

MEAN AND VARIANCE OF NEGATIVE BINOMIAL

- \blacktriangleright A negative binomial distribution can be parameterized in terms of
 - ightharpoonup r and p
 - or μ and σ^2
 - \blacktriangleright or μ and a dispersion parameter α (more on this later)
- ► The relationship between these two parametrizations is given by

$$\mu = r \frac{1-p}{p} \text{ and } \sigma^2 = r \frac{1-p}{p^2},$$

and

$$p = \frac{\mu}{\sigma^2}$$
 and $r = \frac{\mu^2}{\sigma^2 - \mu}$

- ▶ If you provide r and p, you can calculate μ and σ^2
- Or, if you provide μ and σ^2 , you can recover r and p.

Negative Binomial PMF in terms of μ and α

▶ The NB PMF parametrized in terms of p and r (the number of events) is

$$P[K = k] = \binom{k+r-1}{r-1} \pi^r (1-\pi)^k,$$

where k = r, r + 1, r + 2, ...

▶ The NB PMF parametrized in terms of the mean μ and the dispersion parameter α is

$$P[K=k] = \frac{\Gamma[k+\alpha^{-1}]}{\Gamma[\alpha^{-1}]\Gamma[k+1]} \left(\frac{1}{1+\mu\alpha}\right)^{\alpha^{-1}} \left(\frac{\mu}{\alpha^{-1}+\mu}\right)^k,$$

where k = 0, 1, ...

- ▶ The variance is $\mu(1 + \alpha\mu)$
- ▶ As α shrinks to 0 (no-dispersion), the distribution becomes Poisson

NEGATIVE BINOMIAL PMF FOR RNA-SEQ

 \blacktriangleright We will use the mean/dispersion parameter representation for RNA-Seq

$$P[K=k] = \frac{\Gamma[k+\alpha^{-1}]}{\Gamma[\alpha^{-1}]\Gamma[k+1]} \bigg(\frac{1}{1+\mu\alpha}\bigg)^{\alpha^{-1}} \bigg(\frac{\mu}{\alpha^{-1}+\mu}\bigg)^k,$$

where k = 0, 1, ...

- ▶ The variance is $\mu(1 + \alpha \mu)$
- ► IMPORTANT:
 - If $\alpha > 0$, then the variance is greater than the mean. Why?
 - \blacktriangleright As α shrinks to 0 (no-dispersion), the distribution becomes Poisson
- ► More on over-dispersion later

MEANS AND VARIANCES

Distribution	Support	Mean	Variance
Bernoulli (π)	0,1	π	$\pi(1-\pi)$
Binomial (n, π)	$0,1,\ldots,n$	$n\pi$	$n\pi(1-\pi)$
$Poisson(\lambda)$	$0,1,2,\ldots,$	λ	λ
NB(p,r)	$r, r+1, r+2, \ldots,$	$r^{\frac{1-p}{p}}$	$r^{\frac{1-p}{p^2}}$
$NB(\mu, \alpha)$	$0,1,\ldots,$	μ	$\mu(1+\alpha\mu)$

Multinomial Model

- ► Suppose that there are 3 urns
- \blacktriangleright n balls are to be randomly distributed among these M urns
- ▶ Let K_j denote the number of balls assigned to urns j = 1, 2 or 3
- ▶ Let $\pi_j = \mathbb{P}[X_i = j]$ denote the probability that ball i = 1, ..., n is assigned to urn i = 1, 2, 3
- ► Finally let K_j denote the number of balls, among n, assigned to urn j
- ▶ (K_1, K_2, K_3) is said to have multinomial (trinomial) distribution with parameter $(3, \pi_1, \pi_2, \pi_3)$

MULTINOMIAL MODEL: A CHECK LIST

- ▶ Note that $K_1 + K_2 + K_3 = n$
- ► Why?
- ▶ Note that $\pi_1 + \pi_2 + \pi_3 = 1$
- ► Why?
- ► The support of K_j , the number of balls assigned to urn j, is $\{0, 1, ..., n\}$
- ► Why?
- ▶ The support if X_i , the urn to which ball i is assigned, is $\{1, 2, 3\}$
- ► Why?

MULTINOMIAL MODEL: PROPERTIES

▶ The PMF of X_i is

$$\pi_1 = \mathbb{P}[X_i = 1], \pi_2 = \mathbb{P}[X_i = 2] \text{ and } \pi_3 = \mathbb{P}[X_i = 3]$$

► The PMF of

$$\mathbb{P}[K_1 = k_1, K_2 = k_2, K_3 = k_3] = \frac{n!}{k_1! k_2! k_3!} \pi_1^{k_1} \times \pi_2^{k_2} \times \pi_3^{k_3}$$

- ▶ The PMF of K_1 is binomial with parameter (n, π_1)
- ▶ The PMF of K_2 is binomial with parameter (n, π_2)
- ▶ The PMF of K_3 is binomial with parameter (n, π_3)

MULTINOMIAL: RELATIONSHIP TO RNA-SEQ

- ► Suppose that the genome has only three genes (the urns)
- \blacktriangleright n sequencing reads are to be mapped to these three genes
- ▶ K_1 is the number of reads mapped to gene 1
- \blacktriangleright K_2 is the number of reads mapped to gene 2
- ► $K_3 = n K_1 K_3$ is the number of reads mapped to gene 3
- ▶ The multinomial model provides a framework for thinking about the count model for (K_1, K_2, K_3)
- ► One can easily extend the multinomial distribution to arbitrarry number of urns (genes)
- ▶ Caveat: Marginal PMFs do not account for overdispersion

MULTINOMIAL SIMULATION

```
set.seed(3213)
### Number of balls
n <- 100
### The three urn probabilities
pik <- c(2, 1, 4)/7
pik

## [1] 0.2857143 0.1428571 0.5714286

### Simulate two replicates from trinomial distribution with parameter
### (100,2/7,1/7,1/7,1/7)
K <- rmultinom(2, 100, pik)
K

## [,1] [,2]
## [1,] 32 29
## [2,] 17 11
## [3,] 51 60

### Add the two columns to verify they add up to n=100
apply(K, 2, sum)

## [1] 100 100</pre>
```