Primer on Negative Binomial in the context of RNAseq analysis

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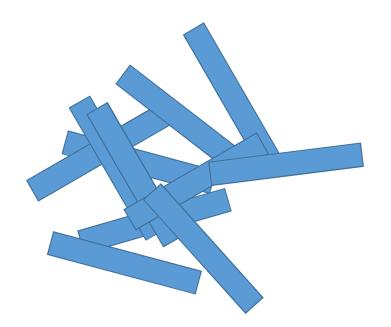
Outline

- Overview on analyzing count data
- Binomial -> Poisson -> Negative binomial (NB)
- Motivation for NB in RNA-seq analysis
- Derivation of NB from the marginalization of Gamma-Poisson mixture
- Mean / Variance relationship of Negative Binomial
- Why this framework works well





θ chance to see a head



n cDNA fragments (reads)

θ_{g} chance to map to the gene g

Gene g

Let's define

 $X = \underline{\text{the number of reads}} \text{ mapped to gene } g \text{ out of } \underline{n \text{ total reads}},$ where each read has $\underline{\text{the probability } \theta_g}$ then Let's define

X =<u>the number of reads</u> mapped to gene g out of <u>n total reads</u>, where each read has <u>the probability</u> θ_g then

$$X \sim Bin(n, \theta_g)$$

$$P(X = x) = {\binom{n}{x}} \theta_g^x (1 - \theta_g)^{n-x}$$

But what if you don't know *n*? (total number of reads)

When $n \to \infty$ $Bin(n,\theta) \rightarrow Pois(\lambda)$ Where $\lambda = n\theta$ $P(X = x) = \frac{\lambda^{x} e^{-\lambda}}{x!}$

Proof $P(X = x) = \binom{n}{x} \theta^{x} (1 - \theta)^{x}$ Define $\lambda = n\theta$ $\Rightarrow \theta = \frac{\lambda}{-}$ $=\frac{n!}{(n-x)!x!}\left(\frac{\lambda}{n}\right)^{x}\left(1-\frac{\lambda}{n}\right)^{n-x}$ $=\frac{n(n-1)\dots(n-x+1)(n-x)!}{(n-x)!x!}\left(\frac{\lambda}{n}\right)^{x}\left(1-\frac{\lambda}{n}\right)^{n-1}$ $=\frac{n(n-1)\dots(n-x+1)}{x!}\left(\frac{\lambda^{x}}{n^{x}}\right)\left(1-\frac{\lambda}{n}\right)^{n-x}$ $=\frac{n^{x}\left(\mathbf{1}-\frac{\mathbf{1}}{n}\right)...\left(\mathbf{1}-\frac{x}{n}+\frac{\mathbf{1}}{n}\right)}{x^{\mathbf{1}}}\left(\frac{\lambda^{x}}{n^{x}}\right)\left(\mathbf{1}-\frac{\lambda}{n}\right)^{n-x}$

$$= \frac{n^{x} \left(1 - \frac{1}{n}\right) \dots \left(1 - \frac{x}{n} + \frac{1}{n}\right)}{x!} \left(\frac{\lambda^{x}}{n^{x}}\right) \left(1 - \frac{\lambda}{n}\right)^{n-x}}$$

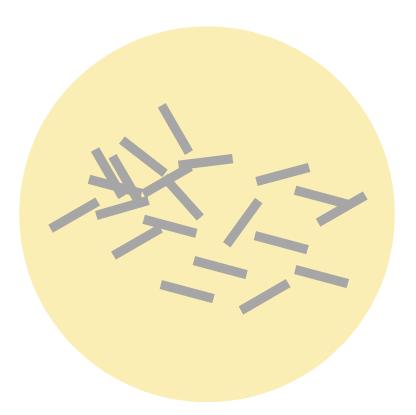
$$= \frac{\left(1 - \frac{1}{n}\right) \dots \left(1 - \frac{x}{n} + \frac{1}{n}\right)}{x!} \lambda^{x} \left(1 - \frac{\lambda}{n}\right)^{n-x}}$$

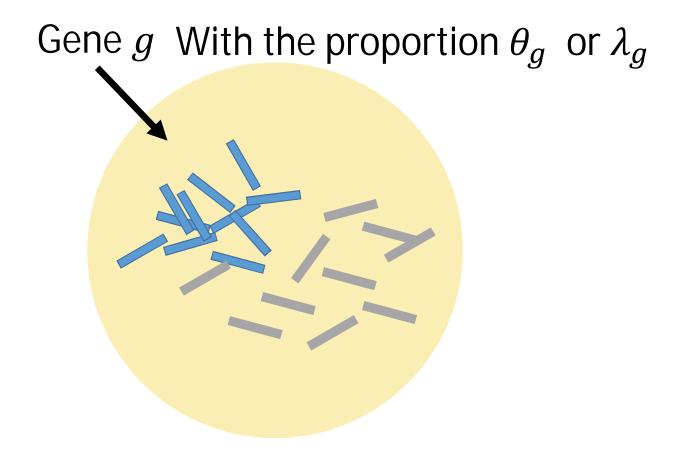
$$= \frac{\left(1 - \frac{1}{n}\right) \dots \left(1 - \frac{x}{n} + \frac{1}{n}\right)}{x!} \lambda^{x} \left(1 - \frac{\lambda}{n}\right)^{n} \left(1 - \frac{\lambda}{n}\right)^{-x}$$

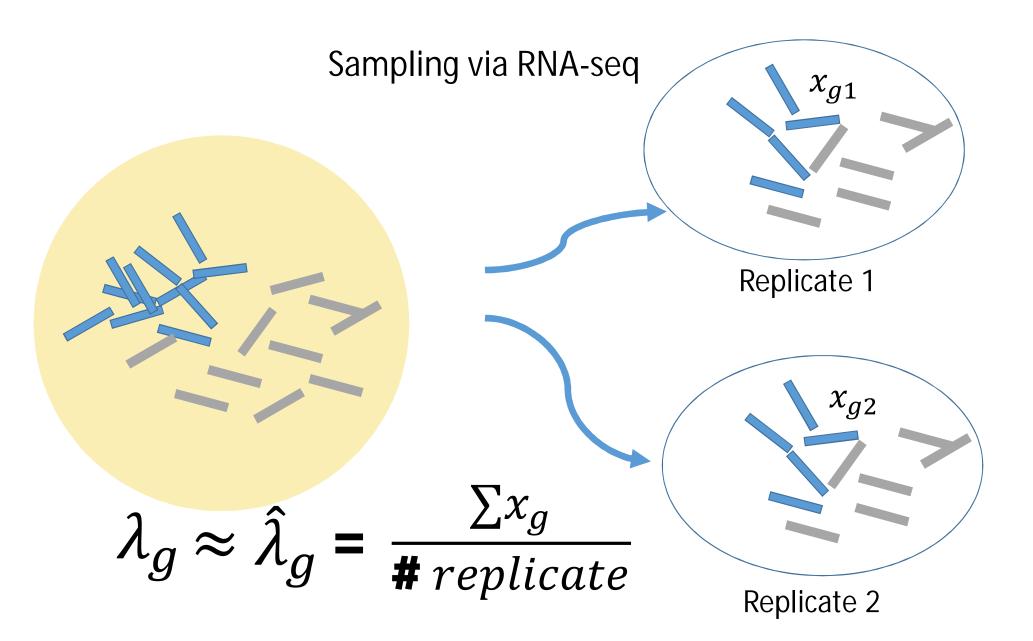
$$n \to \infty$$

$$e^{x} = \lim_{n \to \infty} \left(1 + \frac{x}{n}\right)^{n}$$

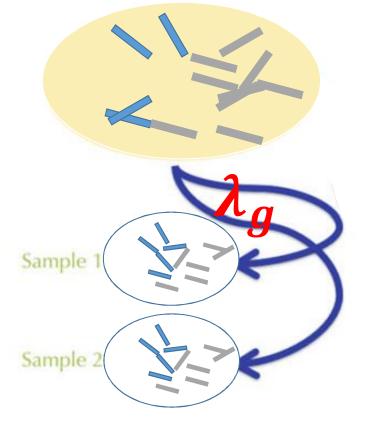
$$= \frac{1}{x!} \lambda^{x} e^{-\lambda}$$







cDNA in the sample *i* mappable to gene *g*



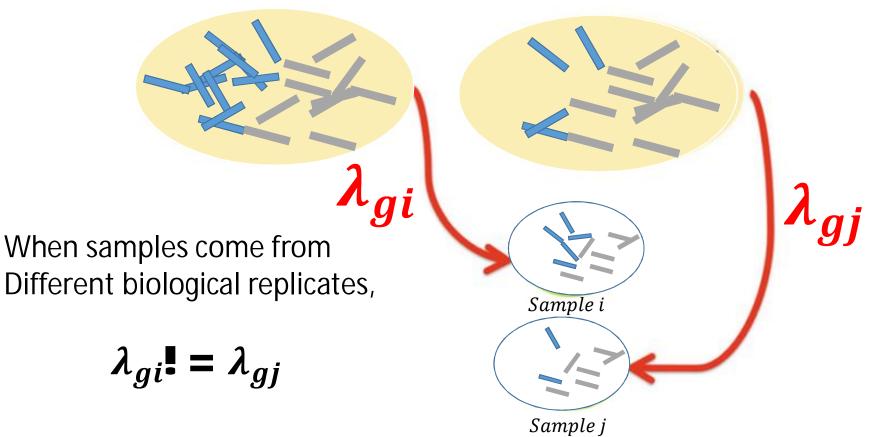
But this only reflects when θ_g (or λ_g) is constant

=> Technical replicates

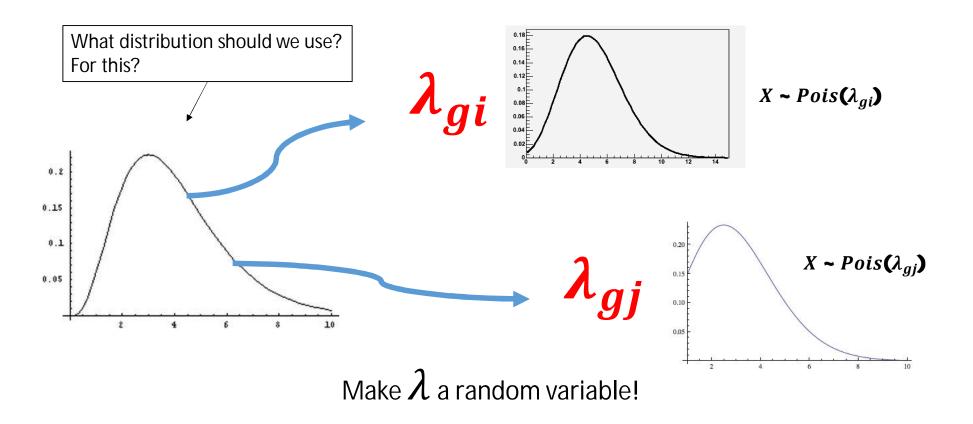
But..

Fragments in the sample *i* mappable to gene *g*

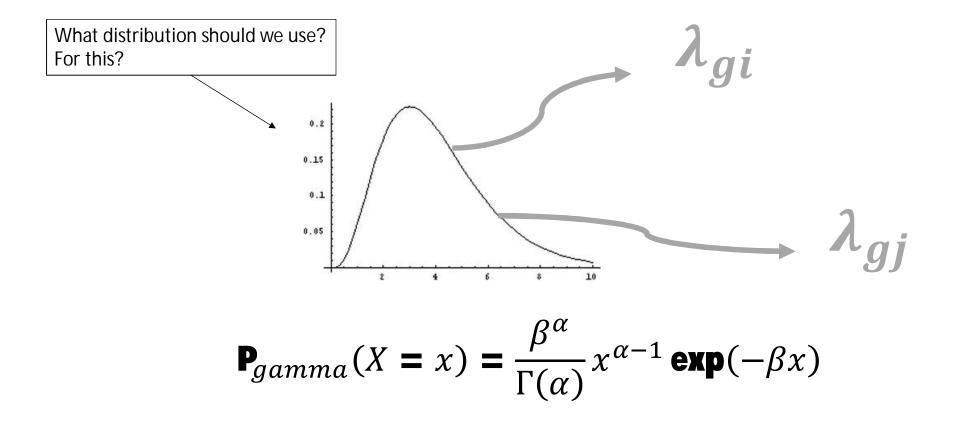
Fragments in the sample *j* mappable to gene *g*



What do you do?



We use gamma distribution



Why gamma?

• It's mathematically convenient

$$\frac{\beta^{\alpha}}{\Gamma(\alpha)} x^{\alpha-1} \exp(-\beta x)$$

Shape parameter α Rate parameter β

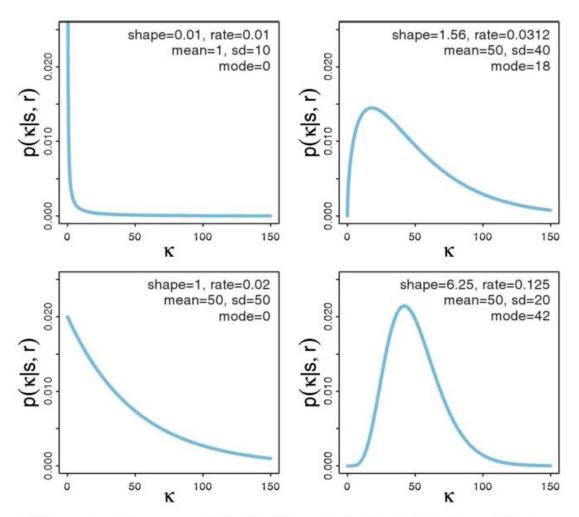
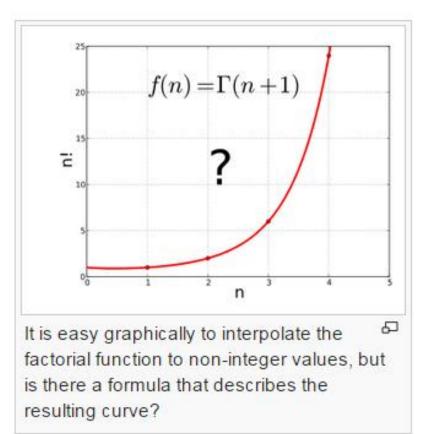


Figure 9.8 Examples of the gamma distribution. The vertical axis is $p(\kappa|s, r)$ where s is the shape and r is the rate, whose values are annotated in each panel. From Doing Bayesian Analysis 2^{nd} ed

What is Gamma function?
$$\Gamma(t)$$
 in $\frac{\beta^{\alpha}}{\Gamma(\alpha)}x^{\alpha-1}\exp(-\beta x)$

$$\Gamma(t) = (t-1)! \text{ if } t \in \mathbb{Z} \ \& t-1 \ge 0$$

but what if $t \in \mathbb{R}$ & $t-1 \geq \mathbf{0}$



 $\sim \sim$

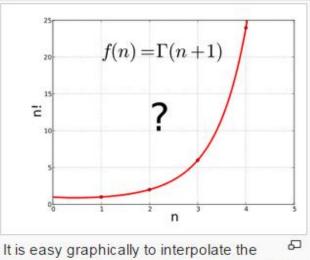
What is Gamma function? $\Gamma(t)$

$$\Gamma(t) = \int_0^\infty x^{t-1} e^{-x} dx$$

$$\frac{\Gamma(t)}{a^t} = \int_0^\infty x^{t-1} e^{-ax} dx$$

$$\int_{0}^{\infty} x^{t-1} e^{-(ax)} dx = a^{t-1} a^{1-t} \int_{0}^{\infty} x^{t-1} e^{-(ax)} dx$$
$$= a^{1-t} \int_{0}^{\infty} (ax)^{t-1} e^{-(ax)} dx$$

 $=a^{-t}\int_0^\infty u^{t-1}e^{-u}du = a^{-t}\Gamma(t)$



factorial function to non-integer values, but is there a formula that describes the resulting curve?

let u = axdu = a dx

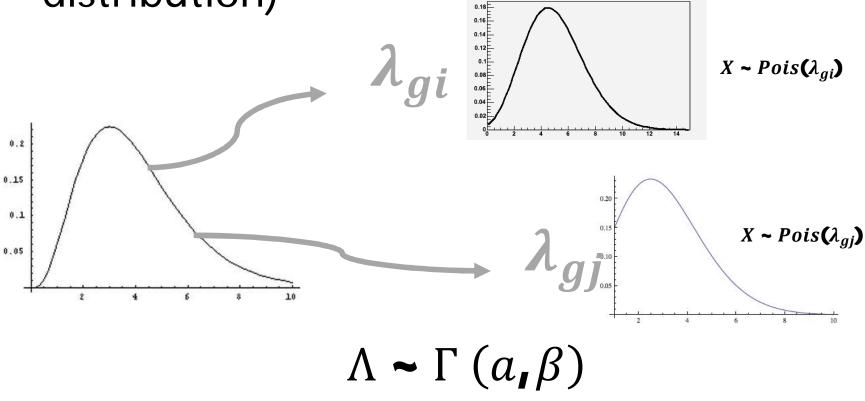
Why Gamma distribution is valid prob. Dist?

$$\mathbf{P}_{gamma}(X = x) = \frac{\beta^{\alpha}}{\Gamma(\alpha)} x^{\alpha-1} \exp(-\beta x)$$

$$\int \mathbf{P}_{gamma}(X = x) = \int \frac{\beta^{\alpha}}{\Gamma(\alpha)} x^{\alpha-1} \exp(-\beta x) dx$$

$$= \frac{\beta^{\alpha}}{\Gamma(\alpha)} \int x^{\alpha-1} \exp(-\beta x) dx$$
$$= \frac{\beta^{\alpha}}{\Gamma(\alpha)} \beta^{-\alpha} \Gamma(\alpha) = \mathbf{1}$$

Hierarchical modeling (or Mixture distribution)



 $X \mid \Lambda = \lambda \sim Pois(\lambda)$

Gamma-Poisson mixture

$$\Lambda \sim \Gamma(\alpha, \beta) \rightarrow P(\lambda)$$

$$(X \mid \Lambda = \lambda) \sim Pois(\lambda) \rightarrow P(x \mid \lambda)$$

$$P(X = x) =?$$

$$P(X = x) = \int_{0}^{\infty} P(\lambda, x) d\lambda \qquad \text{marginalization}$$

$$= \int_{0}^{\infty} P(x \mid \lambda) P(\lambda) d\lambda \qquad \text{Chain rule}$$

$$= \int_{0}^{\infty} \left(\frac{\lambda^{x} e^{-\lambda}}{x!}\right) \left(\frac{\beta^{\alpha}}{\Gamma(\alpha)} \lambda^{a-1} e^{(-\beta\lambda)}\right) d\lambda$$

$$= \frac{1}{x!} \frac{\beta^{\alpha}}{\Gamma(\alpha)} \int_{0}^{\infty} \lambda^{x} e^{-\lambda} \lambda^{\alpha-1} e^{-\beta\lambda} d\lambda$$

$$= \frac{1}{x!} \frac{\beta^{\alpha}}{\Gamma(\alpha)} \int_{0}^{\infty} \lambda^{(x+\alpha)-1} e^{-(\beta+1)\lambda} d\lambda$$

$$= \frac{1}{x!} \frac{\beta^{\alpha}}{\Gamma(\alpha)} \left(\frac{(\beta+1)^{x+\alpha}}{\Gamma(x+\alpha)} \right)^{-1} \left(\frac{(\beta+1)^{x+\alpha}}{\Gamma(x+\alpha)} \right) \int_{0}^{\infty} \lambda^{(x+\alpha)-1} e^{-(\beta+1)\lambda} d\lambda$$

$$= \frac{1}{x!} \frac{\beta^{\alpha}}{\Gamma(\alpha)} \left(\frac{(\beta+1)^{x+\alpha}}{\Gamma(x+\alpha)} \right)^{-1} \left(\frac{(\beta+1)^{x+\alpha}}{\Gamma(x+\alpha)} \right) \int_{0}^{\infty} \lambda^{(x+\alpha)-1} e^{-(\beta+1)\lambda} d\lambda$$
This is negative binomial!
$$= \frac{1}{x!} \frac{\beta^{\alpha}}{\Gamma(\alpha)} \left(\frac{\Gamma(x+\alpha)}{(\beta+1)^{x+\alpha}} \right) = \frac{\Gamma(x+\alpha)}{x!\Gamma(\alpha)} \left(\frac{1}{\beta+1} \right)^{x} \left(\frac{\beta}{\beta+1} \right)^{\alpha}$$

$$= \int_0^\infty \left(\frac{\lambda^x e^{-\lambda}}{x!}\right) \left(\frac{\beta^\alpha}{\Gamma(\alpha)} \lambda^{\alpha-1} e^{-\beta\lambda}\right) d\lambda$$

$$= \frac{\Gamma(x+a)}{x!\Gamma(a)} \left(\frac{1}{\beta+1}\right)^{x} \left(\frac{\beta}{\beta+1}\right)^{a}$$
Let's restrict that x+a is integer, then

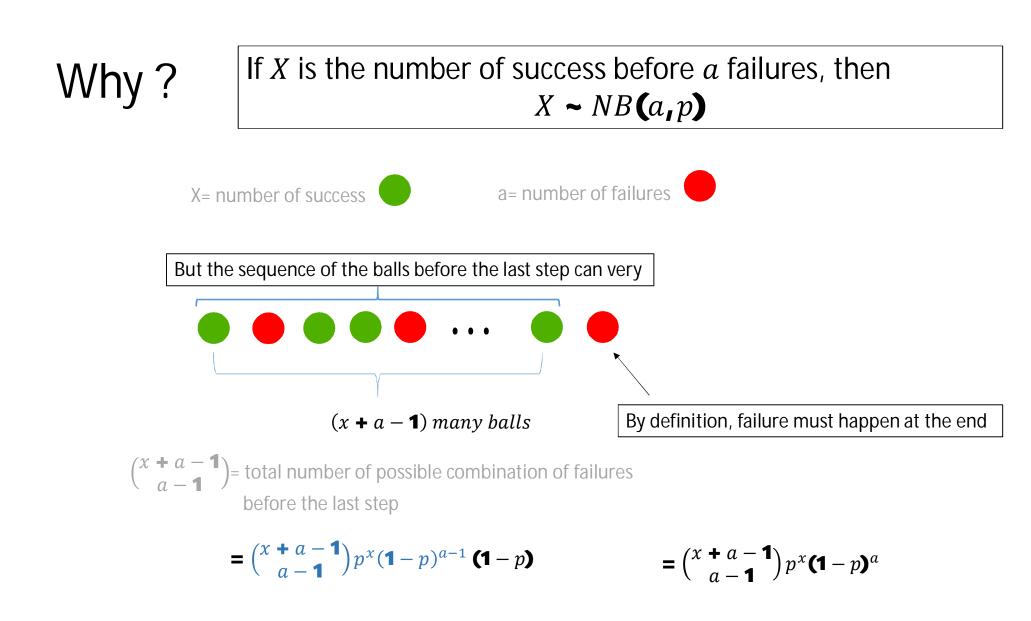
$$\Gamma(x) = (x-1)!$$

$$= \left(\frac{(x+a-1)!}{x!(a-1)!}\right) \left(\frac{1}{\beta+1}\right)^{x} \left(\frac{\beta}{\beta+1}\right)^{a}$$
Because $\binom{n}{k} = \frac{n!}{k!(n-k)!}$

$$= \left(\frac{x+a-1}{a-1}\right) \left(\frac{1}{\beta+1}\right)^{x} \left(\frac{\beta}{\beta+1}\right)^{a}$$
Define $p = \frac{1}{\beta+1} \implies 1-p = \frac{\beta}{\beta+1}$

$$= \left(\frac{x+a-1}{a-1}\right) p^{x} (1-p)^{a}$$

If X is the number of success before a failures, then $X \sim NB(a,p)$



We know that for Poisson : $E[X_{pois}] = \lambda$ $Var[X_{pois}] = \lambda$

But for NB, what's

$$E[X_{NB}] =?$$

 $Var[X_{NB}] =?$

Mean of negative binomial

 $\mathbf{E}[X] = \int \mathbf{x} P(X) dx$

$$= \int_0^\infty x \frac{\Gamma(x+r)}{x! \Gamma(r)} p^x (1-p)^r dx$$

NASTY !

Instead solve it via Moment generating function of NB

Moment generating function = magical formula that spits out things we need: mean,variance,skewness, kurtosis,... or what's collectively known as "moments" (shape of the distribution)

$$M_X(t) = E[e^{tX}]$$

$$M_X'(\mathbf{0}) = E[X]$$

$$M_X''(\mathbf{0}) = E[X^2]$$

$$Var[X] = E[X^2] - E[X]^2$$
Variance

"Secret of the magic trick"

$$e^{x} = \sum_{k=0}^{\infty} \frac{x^{k}}{k!} = \frac{x^{0}}{0!} + \frac{x^{1}}{1!} + \frac{x^{2}}{2!} + \frac{x^{3}}{3!} + \dots$$

$$e^{tX} = \frac{tX^{0}}{0!} + \frac{(tX)^{1}}{1!} + \frac{(tX)^{2}}{2!} + \frac{(tX)^{3}}{3!} + \dots$$

$$M_{X}(t) = \mathbb{E}[e^{tX}] = \mathbb{E}\left[\frac{(tX)^{0}}{0!} + \frac{(tX)^{1}}{1!} + \frac{(tX)^{2}}{2!} + \frac{(tX)^{3}}{3!} + \dots\right]$$

$$= \mathbb{E}\left[\frac{(tX)^{0}}{0!}\right] + \mathbb{E}\left[\frac{(tX)^{1}}{1!}\right] + \mathbb{E}\left[\frac{(tX)^{2}}{2!}\right] + \mathbb{E}\left[\frac{(tX)^{3}}{3!}\right] + \dots$$
Linearity of expectation
$$= 1 + tE[X] + \frac{t^{2}E[X^{2}]}{2!} + \frac{t^{3}E[X^{3}]}{3!} + \dots$$

$$M'_{X}(t) = 0 + E[X] + tE[X^{2}] + \frac{t^{2}E[X^{3}]}{2!} + \dots$$

$$M'_{X}(0) = \mathbb{E}[X]$$

Instead solve it via Moment generating function of NB

$$M_{X}(t) = E[e^{tX}]$$

= $\sum_{x=0}^{\infty} e^{tx} \left({x + r - 1 \atop x} \right) p^{x} (1-p)^{a}$
= $(1-p)^{a} \sum_{x=0}^{\infty} e^{tx} \left({x + r - 1 \atop x} \right) p^{x}$
= $(1-p)^{a} \sum_{x=0}^{\infty} \left({x + r - 1 \atop x} \right) (pe^{t})^{x}$
= $\frac{(1-p)^{a}}{(1-pe^{t})^{a}}$

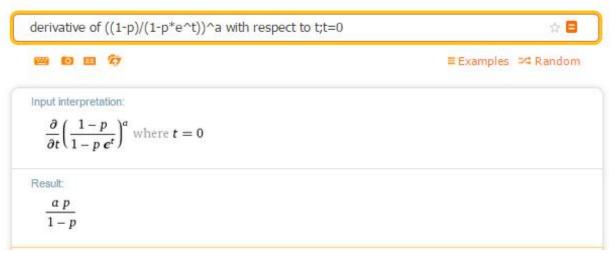
$$M_{X_{NB}}(t) = \left(\frac{1-p}{1-pe^t}\right)^r$$

$$\sum_{x=0}^{\infty} \binom{x+a-1}{x} (pe^t)^x = \frac{1}{(1-pe^t)^a}$$

Mean of NB

$$M_X(t) = \left(\frac{1-p}{1-pe^t}\right)^r$$

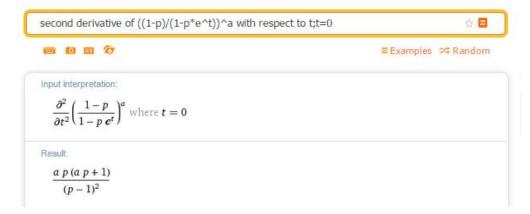




$$M'_X(\mathbf{0}) = \frac{ap}{\mathbf{1} - p} = E[X]$$

Variance of NB

WolframAlpha computational...



$$M_X''(\mathbf{0}) = \frac{ap(ap + \mathbf{1})}{(p - \mathbf{1})^2} = E[X^2]$$

$$Var(X) = E[X^{2}] - E[X]^{2} = \frac{ap(ap + 1)}{(p - 1)^{2}} - \left(\frac{ap}{1 - p}\right)^{2} = \frac{ap}{(1 - p)^{2}}$$

Denouement

We have

We have
$$E[X] = \frac{ap}{1-p}$$
 $Var[X] = \frac{ap}{(1-p)^2}$
Remember $p = \frac{1}{\beta + 1}$ $1 - p = \frac{\beta}{\beta + 1}$

Then

$$E[X] = \frac{ap}{1-p} = \frac{a\left(\frac{1}{\beta+1}\right)}{\frac{\beta}{\beta+1}} = \frac{a}{\beta}$$

$$Var[X] = \frac{ap}{(1-p)^2} = \frac{a\left(\frac{1}{\beta+1}\right)}{\left(\frac{\beta}{\beta+1}\right)^2} = \frac{a}{\beta^2}(\beta+1)$$
$$= \frac{a}{\beta} + \frac{a}{\beta^2}$$
$$= \frac{a}{\beta} + \left(\frac{a}{\beta}\right)^2 \left(\frac{1}{a}\right)$$

Therefore

$$Var[X] = E[X] + \left(\frac{1}{a}\right)E[X]^2$$

We know that for poisson : $E[X_{pois}] = \lambda$ $Var[X_{pois}] = \lambda$ And for NB, $E[X_{NB}] = \mu$ $Var[X_{NB}] = \mu + c\mu^2$

Dispersion

Most efforts of most RNAseq packages...

$$E[X_{NB}] = \mu$$
$$Var[X_{NB}] = \mu + c\mu^{2}$$

Estimate dispersion using small # of samples

EdgeR : (old) assume c is identical for all genes

(new) c is different for individual genes Using hierarchical modeling to shrink to c to a consensus value

DESeq : Assume c and μ is related nonlinearly. use local regression to compute c

Why is this nice???

- Negative binomial is exponential family! So you can model complicated stuff using generalized linear model.

$$E[X_{NB}] = \mu$$
 X = design matrix
Assume $\mu = e^{X^T \beta}$ β = vector of coefficients
Estimate β

- Empirically, packages that use NB as an underlying model are shown to work better than those that don't

THE END