

Week 3 QCB 408/508 Spring 2020

$$\pi_{ij} = \frac{x_{ij}}{M_j} \quad \text{random proportion of gene } i \text{ mRNA in observation } j$$

$$\begin{aligned} E[\pi_{ij}] &= E\left[\frac{x_{ij}}{M_j}\right] \\ &= E\left[E\left[\frac{x_{ij}}{M_j} | M_j\right]\right] \\ &= E\left[E\left[\frac{a_i M_j}{M_j} | M_j\right]\right] \\ &= E[a_i] = a_i \end{aligned}$$

$$\text{Var}(\pi_{ij}) = E\left[\text{Var}(\pi_{ij} | M_j)\right] + \underbrace{\text{Var}(E\left[\pi_{ij} | M_j\right])}_{0}$$

$$\begin{matrix} \text{Var}(ct) \\ = c^2 \text{Var}(x) \end{matrix}$$

$$\begin{aligned} \text{Var}(\pi_{ij} | M_j) &= \text{Var}\left(\frac{x_{ij}}{M_j} | M_j\right) \\ &= \frac{1}{M_j^2} \text{Var}(x_{ij} | M_j) \\ &\approx \frac{1}{M_j^2} a_i M_j = \frac{a_i}{M_j} \end{aligned}$$

$$\Rightarrow \text{Var}(\pi_{ij}) \approx E\left[\frac{\pi_{ij}}{n_j}\right] = \pi_{ij} E\left[\frac{1}{n_j}\right]$$

\approx "biological variance"

Step 2.

This is assumed (here) to be completely random sampling.

Let D_j be the "read depth" of observation j , which is the total number of reads from observation j .

Note: We observe D_j , say d_j .

Reminder: Y_{ij} RNA-Seq read counts for gene i , obs. j .

$Y_{ij} | \pi_{ij}, d_j \sim \text{Binomial}(d_j, \pi_{ij})$

$\sim \text{Poisson}(d_j \pi_{ij})$

d_j large, π_{ij} small

$$E[Y_{ij}] = E[E[Y_{ij} | \pi_{ij}, d_j]]$$

$$\begin{aligned}
 &= E[\pi_{ij} d_j] \\
 &\geq d_j E[\pi_{ij}] \\
 &= d_j a_i
 \end{aligned}$$

$$\begin{aligned}
 \text{Var}(Y_{ij}) &= E[\text{Var}(Y_{ij} | \pi_{ij}, d_j)] + \text{Var}(E[Y_{ij} | \pi_{ij}, d_j]) \\
 &\approx E[\pi_{ij} d_j] + \text{Var}(\pi_{ij} d_j) \\
 &= a_i d_j + d_j^2 \text{Var}(\pi_{ij}) \\
 &= a_i d_j + d_j^2 a_i \in \left[\frac{1}{m_j} \right]
 \end{aligned}$$

$> E(Y_{ij}) \Rightarrow$ over-dispersed Poisson

$$\text{Estimate } \hat{\pi}_{ij} = \frac{Y_{ij}}{d_j}, \quad E[\hat{\pi}_{ij}] = a_i$$

$$\begin{aligned}
 \text{Var}(\hat{\pi}_{ij}) &= \frac{1}{d_j^2} \text{Var}(Y_{ij}) \\
 &= \frac{a_i}{d_j} + \text{Var}(\pi_{ij}) \\
 &= \frac{a_i}{d_j} + a_i \in \left[\frac{1}{m_j} \right]
 \end{aligned}$$

$$\text{technical variance} \approx \frac{a_i}{d_j}$$

$$\text{biological variance} \approx a_i E\left[\frac{1}{m_j}\right]$$

$$\text{Consider estimate } \hat{a}_i = \frac{\sum_{j=1}^n \hat{\pi}_{ij}}{n},$$

$$E[\hat{a}_i] = a_i.$$

$$\begin{aligned} \text{Var}(\hat{a}_i) &= \text{Var}\left(\frac{\sum_{j=1}^n \hat{\pi}_{ij}}{n}\right) \\ &= \frac{1}{n^2} \text{Var}\left(\sum_{j=1}^n \hat{\pi}_{ij}\right) \\ &= \frac{1}{n^2} \sum_{j=1}^n \text{Var}(\hat{\pi}_{ij}) \\ &= \frac{a_i}{n^2} \sum_{j=1}^n \frac{1}{d_j} + \frac{\sum_{j=1}^n \text{Var}(\hat{\pi}_{ij})}{n^2} \end{aligned}$$

Let's assume M_j are i.i.d.

Then $E\left[\frac{1}{m_j}\right]$ is the same for all j .

Coefficient of Variation, CV

$$CV = \frac{\sqrt{Var(\pi_{ij})}}{a_i} \Rightarrow \text{biological CV}$$

$$(CV)^2 = \frac{Var(\pi_{ij})}{a_i^2} = \frac{1}{a_i} E\left[\frac{1}{m_j}\right] = \phi_i$$

Let $\mu_{ij} = d_j a_i$ (population mean times
obs. read depth)

$$\Rightarrow Var(Y_{ij}) = d_j a_i + d_j^2 Var(\pi_{ij})$$

$$= d_j a_i + (d_j a_i)^2 \frac{Var(\pi_{ij})}{a_i^2}$$

$$= \mu_{ij} + \mu_{ij}^2 \phi_i$$

 estimated by
 "borrowing strength"
 across genes - with
 similar ϕ_i values

Negative Binomial

Bernoulli trials with success p

$Y =$ number of failures before r^{th} success

$Y \sim \text{Neg Bin}(r, p)$

$$\Pr(Y=y) = \binom{r+y-1}{y} p^r (1-p)^y$$

$$y=0, 1, 2, \dots$$

$$E[Y] = r \frac{(1-p)}{p} \quad \text{Var}(Y) = r \frac{(1-p)}{p^2}$$

$$\text{Let } \mu = \frac{r(1-p)}{p}$$

$$\text{Var}(Y) = \mu + \mu^2 \left(\frac{1}{r} \right) \phi$$

RNA-seq data under the above model
is therefore sometimes modeled as a
Neg Bin.

$$Y_{ij} \sim \text{NegBin}(p_{ij}, r_i)$$

$$\text{where } \mu_{ij} = \frac{r_i(1-p_{ij})}{p_{ij}} \text{ and } \phi_i = \frac{1}{r_i}.$$

Compound Gamma-Poisson Distribution

$$Y_{ij} | \lambda_{ij} \sim \text{Poisson}(\lambda_{ij})$$

$$\lambda_{ij} \sim \text{Gamma}(\alpha, \beta)$$

Then Y_{ij} is marginally a Gamma-Poisson rv.

Neg Bin \rightarrow a special case of
Gamma-Poisson.

Suppose $Y|\lambda \sim \text{Poisson}(\lambda)$

$\lambda \sim \text{Gamma}(\alpha, \beta)$

$$\text{Gamma PDF} \quad f(\lambda; \alpha, \beta) = \frac{\lambda^{\beta-1} e^{-\lambda/\alpha}}{\alpha^\beta \Gamma(\beta)} \quad \lambda > 0$$

$$E[\lambda] = \alpha\beta, \quad \text{Var}(\lambda) = \alpha\beta$$

Gamma-Poisson:

$$f(y; \alpha, \beta) = \frac{\Gamma(y+\beta) \alpha^y}{\Gamma(\beta) (1+\alpha)^{\beta+y} y!}$$

$$E[Y] = \alpha\beta \quad \text{Var}(Y) = \alpha\beta + \alpha^2\beta$$

$$\text{Let } \mu = \alpha\beta$$

$$\text{Var}(Y) = \mu + \mu^2 \cdot \frac{1}{\beta}$$

Let's map this back to the 2-step model.

$$Y_{ij} | \lambda_{ij} \sim \text{Poisson}(\lambda_{ij})$$

$$\lambda_{ij} \sim \text{Gamma}(\alpha, \beta) \xrightarrow{\substack{\text{let's determine} \\ \text{what are } \lambda_{ij}, \\ \alpha, \text{ and } \beta}}$$

λ_{ij} is $\pi_{ij} d_j$, which is random

$$E[\pi_{ij} d_j] = a_i d_j \quad \text{and}$$

$$E[\lambda_{ij}] = \alpha \beta . \quad \text{so}$$

$$\mu_{ij} = a_i d_j = \alpha \beta$$

$$\text{Var}(Y_{ij}) = \mu_{ij} + \mu_{ij}^2 \phi_i,$$

$$\text{so } \frac{1}{\beta} = \phi_i = \frac{1}{a_i} E\left[\frac{1}{M_j}\right].$$

$$\Rightarrow \beta_{ij} = a_i E\left[\frac{1}{M_j}\right]^{-1}$$

$$\Rightarrow \alpha_{ij} = a_i d_j \cdot \frac{1}{a_i} E\left[\frac{1}{M_j}\right]$$

$$= d_j \cdot E\left[\frac{1}{M_j}\right]$$

$$\Rightarrow \lambda_{ij} \sim \text{Gamma}(\alpha_{ij}, \beta_{ij}).$$

$$Y|\lambda \sim \text{Poisson}(\lambda) , \quad \lambda > 0$$

λ is r.v. with $E[\lambda]$ and $\text{Var}(\lambda)$

$$\text{Var}(Y) = E[\text{Var}(Y|\lambda)] + \text{Var}(E[Y|\lambda])$$

$$= E[\lambda] + \text{Var}(\lambda)$$

Since $\lambda > 0$, $E[\lambda]$ will "appear"

in $\text{Var}(\lambda) \Rightarrow$ mean-variance relationship

Sums of Random Variables

If X is a rv and a, b are constants,

then $E[a + bX] = a + bE[X]$ and

$$\text{Var}(a + bX) = b^2 \text{Var}(X).$$

Let X_1, X_2, \dots, X_n be n rv's. Then

$$E\left[\sum_{i=1}^n X_i\right] = \sum_{i=1}^n E[X_i]$$

$$\text{Var}\left(\sum_{i=1}^n X_i\right) = \sum_{i=1}^n \text{Var}(X_i) + \sum_{i \neq j} \text{Cov}(X_i, X_j)$$

$$= \sum_{i=1}^n \sum_{j=1}^n \text{Cov}(X_i, X_j)$$

when X_1, X_2, \dots, X_n are independent,

then $\text{Cov}(X_i, X_j) = 0$ for $i \neq j$

so $\text{Var}(\sum X_i) = \sum \text{Var}(X_i)$.

Let $\bar{X}_n = \frac{1}{n} \sum_{i=1}^n X_i$. Suppose X_1, X_2, \dots, X_n are independent.

$$\begin{aligned} E[\bar{X}_n] &= E\left[\frac{1}{n} \sum_{i=1}^n X_i\right] \\ &= \frac{1}{n} E\left[\sum_{i=1}^n X_i\right] \\ &= \frac{1}{n} \underbrace{\sum_{i=1}^n E[X_i]} \end{aligned}$$

when $E[X_1] = E[X_2] = \dots = E[X_n] = \theta$

then $E[\bar{X}_n] = \theta$.

$$\text{Var}(\bar{X}_n) = \frac{1}{n^2} \sum \text{Var}(X_i)$$

If $\text{Var}(X_1) = \text{Var}(X_2) = \dots = \text{Var}(X_n) = \sigma^2$

$$\text{Var}(\bar{X}_n) = \frac{\sigma^2}{n}$$

Convergence of RV's

Let Z_1, Z_2, \dots be a sequence of rv's.

Examples: $Z_n = \bar{X}_n$.

$$Z_n \sim \text{Binomial}(n, p)$$

① Convergence in Distribution:

$\{Z_n\}$ converges in distribution to rv W

$Z_n \xrightarrow{D} W$ as $n \rightarrow \infty$ if

$$F_{Z_n}(y) = \Pr(Z_n \leq y) \rightarrow \Pr(W \leq y) = F_W(y)$$

for all $y \in \mathbb{R}$, as $n \rightarrow \infty$.

② Convergence in Probability:

$Z_n \xrightarrow{P} W$ as $n \rightarrow \infty$ if

$$\Pr(|Z_n - W| \leq \varepsilon) \rightarrow 1 \text{ as}$$

$n \rightarrow \infty$, for $\varepsilon > 0$.

If θ is a fixed number, then
we can have $Z_n \xrightarrow{P} \theta$.

③ Almost sure convergence

$\{Z_n\}$ converges "almost surely" (a.s.)
or "with probability 1" to W

$Z_n \xrightarrow{\text{a.s.}} W$ if

$$\Pr(\{\omega : |Z_n(\omega) - W(\omega)| \xrightarrow{n \rightarrow \infty} 0\}) = 1$$

Strong Law of Large Numbers

Suppose X_1, X_2, \dots are iid r.v's with
population mean $E[X_i] = \mu$ where
 $E[|X_i|] < \infty$. Then

$$\bar{X}_n \xrightarrow{\text{a.s.}} \mu, \text{ as } n \rightarrow \infty.$$

Central Limit Theorem

Suppose X_1, X_2, \dots are iid r.v's
with $E[X_i] = \mu$ and $\text{Var}(X_i) = \sigma^2$.

Then as $n \rightarrow \infty$,

$$\sqrt{n}(\bar{X}_n - \mu) \xrightarrow{D} \text{Normal}(0, \sigma^2)$$

$$\frac{\bar{X}_n - \mu}{\sigma/\sqrt{n}} \xrightarrow{D} \text{Normal}(0, 1)$$

Note:

$$\text{Var}(\bar{X}_n - \mu) = \text{Var}(\bar{X}_n) = \frac{\sigma^2}{n}$$

$$\text{Var}\left(\frac{\bar{X}_n - \mu}{\sigma/\sqrt{n}}\right) = \frac{1}{\sigma^2/n} \text{Var}(\bar{X}_n) = 1$$

$$\frac{\bar{X}_n - \mu}{\sigma/\sqrt{n}} = \sqrt{n} \left(\frac{\bar{X}_n - \mu}{\sigma} \right) \xrightarrow{D} N(0, 1).$$

Normal r.v's - some useful facts

$$X_1, X_2, \dots, X_n \stackrel{iid}{\sim} \text{Normal}(\mu, \sigma^2)$$

$$E[\bar{X}_n] = \mu, \quad \text{Var}(\bar{X}_n) = \frac{\sigma^2}{n}$$

$$\bar{X}_n \sim \text{Normal}(\mu, \sigma^2/n)$$

because

$$X_1 + X_2 + \dots + X_n \sim \text{Normal}(n\mu, n\sigma^2)$$

$$aX_1 + b \sim \text{Normal}(a\mu + b, a^2\sigma^2)$$