Incomplete Data Analysis

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- \hookrightarrow The computation of the E and M steps simplify when it can be shown that the log likelihood of the complete data is linear in the **sufficient statistics** for θ .
- In particular, this turns out to be the case when the distribution of the complete data belongs to the exponential family.
- → But what is a sufficient statistic? And what do we mean by a distribution belonging to the exponential family?

Sufficient statistic

 \hookrightarrow A **statistic** is simply a function $T = T(Y_1, \dots, Y_n)$ of the random sample, e.g.,

$$T = \frac{1}{n} \sum_{i=1}^{n} Y_i,$$

$$T = \frac{1}{n-1} \sum_{i=1}^{n} (Y_i - \bar{Y}),$$

$$T = \max\{Y_1, \dots, Y_n\}.$$

Definition

A statistic $T = T(Y_1, \dots, Y_n)$ is said to be **sufficient** for θ if the conditional distribution of Y_1, \dots, Y_n given T = t does not depend on θ for any value of t.

Sufficient statistic

- \hookrightarrow A sufficient statistic for θ contains all the information in the sample about θ .
- \hookrightarrow Thus, given the value of T, we cannot improve our knowledge about θ by a more detailed analysis of the data Y_1, \ldots, Y_n .
- \hookrightarrow This basically, and informally, means that the statistician who knows the value of T can do as just as a good job of estimating the unknown parameter θ as the statistician who knowns the entire random sample.
- \hookrightarrow In other words, an estimate based on T=t cannot be improved by using the data Y_1,\ldots,Y_n .

Sufficient statistic

 $\,\hookrightarrow\,$ As an example, consider a sequence of independent Bernoulli trials

$$Y_i \stackrel{\text{iid}}{\sim} \text{Bernoulli}(\theta), \qquad i = 1, \dots, n.$$

- \hookrightarrow The number of successes $T = \sum_{i_1}^{n} Y_i$ is a sufficient statistic for the parameter θ .
- \hookrightarrow Additional information about the observed values Y_1, \ldots, Y_n , as e.g. the order in which the successes occurred, does not convey any information about θ .

Exponential family

Definition

One-parameter members of the exponential family have density function of the form

$$f(y; \theta) = b(y) \exp\{c(\theta)t(y) - a(\theta)\}.$$

- \hookrightarrow It can be shown (see, e.g., Rice, 2006, Chapter 8) that t(y) is a sufficient statistic for θ .
- \hookrightarrow Suppose that Y_1, \ldots, Y_n is a sample from a member of the (one-parameter) exponential family, then the joint density function is given by

$$f(\mathbf{y}; \theta) = \prod_{i=1}^{n} \{b(y_i) \exp\{c(\theta)t(y_i) - a(\theta)\}\}$$
$$= \left\{\prod_{i=1}^{n} b(y_i)\right\} \exp\left\{c(\theta) \sum_{i=1}^{n} t(y_i) - na(\theta)\right\}$$

Exponential family

Definition

A p-parameter member of the exponential family has a density function of the form

$$f(y; \theta) = b(y) \exp \left\{ \sum_{j=1}^{p} c_j(\theta) t_j(y) - a(\theta) \right\}$$

- \hookrightarrow Analogously to the single parameter case, it can be shown that $(T_1(y), \ldots, T_p(y))$ is sufficient for θ .
- \hookrightarrow Suppose that Y_1, \ldots, Y_n is a sample from a member of the (*p*-parameter) exponential family, then the joint density function is given by

$$f(\mathbf{y}; \boldsymbol{\theta}) = \prod_{i=1}^{n} \left\{ b(y) \exp \left\{ \sum_{j=1}^{p} c_{j}(\boldsymbol{\theta}) t_{j}(y) - a(\boldsymbol{\theta}) \right\} \right\}$$
$$= \left\{ \prod_{i=1}^{n} b(y_{i}) \right\} \exp \left\{ \sum_{i=1}^{n} \sum_{j=1}^{p} c_{j}(\boldsymbol{\theta}) t_{j}(y_{i}) - na(\boldsymbol{\theta}) \right\}$$

Exponential family

 \hookrightarrow Binomial distribution: $Y \sim Bin(n, \theta)$.

$$f(y) = \binom{n}{y} \theta^{y} (1 - \theta)^{n-y}$$
$$= \binom{n}{y} \exp\left\{y \log\left(\frac{\theta}{1 - \theta}\right) + n \log(1 - \theta)\right\},$$

where

$$c(\theta) = log\left(\frac{\theta}{1-\theta}\right),$$

$$t(y) = y.$$

Exponential family

 \hookrightarrow Normal distribution: $Y \sim N(\mu, \sigma^2)$.

$$\begin{split} f(y) &= \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left\{ -\frac{1}{2\sigma^2} (y - \mu)^2 \right\} \\ &= \exp\left\{ -\frac{1}{2\sigma^2} y^2 + \frac{\mu}{\sigma^2} y - \frac{\mu^2}{2\sigma^2} - \frac{1}{2} \log(2\pi\sigma^2) \right\}, \end{split}$$

with

$$c(\mu, \sigma^2) = \left(-\frac{1}{2\sigma^2}, \frac{\mu}{\sigma^2}\right),$$

$$t(\gamma) = (\gamma^2, \gamma).$$

 \hookrightarrow If $Y_i \stackrel{\text{iid}}{\sim} N(\mu, \sigma^2)$, for $i = 1, \dots, n$, then

$$t(\mathbf{y}) = \left(\sum_{i=1}^n y_i^2, \sum_{i=1}^n y_i\right).$$



- → In what follows I will be focusing in the case of a scalar parameter, but everything follows analogously to the multiparameter case.
- → Let us assume that the distribution of the complete data y belongs to the exponential family. Then the log likelihood of the complete data can be written as

$$\log f(\mathbf{y};\theta) = \log b(\mathbf{y}) + c(\theta)t(\mathbf{y}) - a(\theta).$$

$$\begin{split} Q\left(\theta\mid\theta^{(t)}\right) &= E_{Y_{\text{mis}}}\left[\log b(\mathbf{y}) + c(\theta)t(\mathbf{y}) - a(\theta)\mid\mathbf{y}_{\text{obs}},\theta^{(t)}\right] \\ &= \text{constant} + c(\theta)E_{Y_{\text{mis}}}\left[t(\mathbf{y})\mid\mathbf{y}_{\text{obs}},\theta^{(t)}\right] - a(\theta). \end{split}$$

 \hookrightarrow The M-step is

$$\frac{\mathsf{d}}{\mathsf{d}\theta} \mathcal{Q}\left(\theta \mid \theta^{(t)}\right) = 0 \Rightarrow \frac{\mathsf{d}}{\mathsf{d}\theta} c(\theta) \mathcal{E}_{\mathsf{Y}_{\mathsf{mis}}} \left[\mathit{t}(\mathbf{y}) \mid \mathbf{y}_{\mathsf{obs}}, \theta^{(t)} \right] = \frac{\mathsf{d}}{\mathsf{d}\theta} \mathit{a}(\theta).$$

→ From the log likelihood of the complete data, we know that

$$\frac{\mathsf{d}}{\mathsf{d}\theta}\log f(\mathbf{y};\theta) = \frac{\mathsf{d}}{\mathsf{d}\theta}c(\theta)t(\mathbf{y}) - \frac{\mathsf{d}}{\mathsf{d}\theta}a(\theta).$$

A known result from likelihood theory is that the (unconditional) expected value of the score function (derivative of the log likelihood) is zero. Therefore,

$$E\left[\frac{\mathsf{d}}{\mathsf{d}\theta}\log f(\boldsymbol{y};\theta)\right] = 0 = \frac{\mathsf{d}}{\mathsf{d}\theta}c(\theta)E[t(\boldsymbol{y})] - \frac{\mathsf{d}}{\mathsf{d}\theta}a(\theta) \Rightarrow \frac{\mathsf{d}}{\mathsf{d}\theta}a(\theta) = \frac{\mathsf{d}}{\mathsf{d}\theta}c(\theta)E[t(\boldsymbol{y})].$$

 \hookrightarrow Therefore, the M-step reduces to finding $\theta^{(t+1)}$ as a solution to

$$E_{Y_{\mathsf{mis}}}\left[t(\mathbf{y})\mid\mathbf{y}_{\mathsf{obs}},\theta^{(t)}
ight]=E[t(\mathbf{y})].$$



Summary of the E and M steps

- **© E-step**: Compute the expected value of the sufficient statistics for the complete data, given the observed data and using the current parameter estimate $\theta^{(t)}$. Let $\mathbf{t}^{(t)} = E_{Y_{\text{mis}}}\left[t(\mathbf{y}) \mid \mathbf{y}_{\text{obs}}, \theta^{(t)}\right]$.
- **M-step**: Set $\theta^{(t+1)}$ to the value that makes the unconditional expectation of the sufficient statistics for the complete data equal to $t^{(t)} = E_{Y_{\text{mis}}} [t(\mathbf{y}) \mid \mathbf{y}_{\text{obs}}, \theta^{(t)}]$. In other words, $\theta^{(t+1)}$ solves $E[t(\mathbf{y})] = \mathbf{t}^{(t)}$.
- Return to the E step unless a convergence criterion has been met.

- \hookrightarrow Let us revisit the bivariate normal example from week 6 where Y_1 was fully observed but only the first m values of Y_2 were available.
- \hookrightarrow The joint density of Y_1 and Y_2 can be equivalently written as

$$f(y_1, y_2) = \frac{1}{2\pi\sigma_1\sigma_2\sqrt{1-\rho^2}} \exp\left\{-\frac{1}{2(1-\rho^2)} \left(\frac{(y_1-\mu_1)^2}{\sigma_1^2} + \frac{(y_2-\mu_2)^2}{\sigma_2^2} - 2\rho\frac{(y_1-\mu_1)(y_2-\mu_2)}{\sigma_1\sigma_2}\right)\right\}.$$

- → The bivariate normal distribution belongs to the exponential family.

$$\left(\sum_{i=1}^n y_{1i}, \sum_{i=1}^n y_{1i}^2, \sum_{i=1}^n y_{2i}, \sum_{i=1}^n y_{2i}^2, \sum_{i=1}^n y_{1i}y_{2i}\right).$$



Example: bivariate normal data with one variable subject to missingness

- \hookrightarrow Remember that $\theta = (\mu_1, \sigma_1^2, \mu_2, \sigma_2^2, \sigma_{12}), \mathbf{y}_{\text{obs}} = (y_{11}, \dots, y_{1n}, y_{21}, \dots, y_{2m})$ and $\mathbf{y}_{\text{mis}} = (y_{2m+1}, \dots, y_{2n}).$

$$t_1^{(t)} = E_{Y_{\text{mis}}} \left(\sum_{i=1}^n y_{1i} \mid \mathbf{y}_{\text{obs}}, \theta^{(t)} \right) = \sum_{i=1}^n y_{1i},$$

$$t_{11}^{(t)} = \textit{E}_{\textit{Y}_{mis}}\left(\sum_{i=1}^{n} \textit{y}_{1i}^{2} \mid \mathbf{y}_{obs}, \theta^{(t)}\right) = \sum_{i=1}^{n} \textit{y}_{1i}^{2}.$$

Note that because we do not have missing values on Y_1 , the conditional expectations equal the observed values.

Example: bivariate normal data with one variable subject to missingness

→ Because Y₂ has some missing values and the expectation of those is conditional on the current parameter estimate and on what is observed (and so, in particular, on the corresponding Y₁ values) and remembering that

$$Y_{2} \mid Y_{1} = y_{1} \sim N(\beta_{0} + \beta_{1}y_{1}, \sigma_{2|1}^{2}),$$

$$\beta_{0} = \mu_{2} - \beta_{1}\mu_{1},$$

$$\beta_{1} = \frac{\sigma_{12}}{\sigma_{1}^{2}},$$

$$\sigma_{2|1}^{2} = \sigma_{2}^{2} - \frac{\sigma_{12}^{2}}{\sigma_{1}^{2}}.$$

we thus have

$$\begin{split}
& = \sum_{i=1}^{n} y_{2i} \mid y_{olos}, \theta^{(\epsilon)} \\
& = \sum_{i=1}^{n} y_{2i} + \mathbb{E}_{y_{mis}} \left(\sum_{i=m+1}^{n} y_{2i} \mid y_{olos}, \theta^{(\epsilon)} \right) \\
& = \sum_{i=1}^{n} y_{2i} + \sum_{i=m+1}^{n} \mathbb{E}_{x_{i}} \left[y_{olos}, \theta^{(\epsilon)} \right] \\
& = \sum_{i=1}^{n} y_{2i} + \sum_{i=m+1}^{n} \mathbb{E}_{x_{i}} \left[y_{olos}, \theta^{(\epsilon)} \right]
\end{split}$$

$$\frac{1}{2} \sum_{i=1}^{m} y_{i2}^{2} + \sum_{i=m+1}^{m} \frac{1}{2} \left(|\nabla_{2h}|^{2} \right)^{2} + \left(|\nabla_{2h}|^{$$

Example: bivariate normal data with one variable subject to missingness

 \hookrightarrow For the M-step we need

$$t_{1}^{(t)} = E\left(\sum_{i=1}^{n} y_{1i}\right) = n\mu_{1},$$

$$t_{11}^{(t)} = E\left(\sum_{i=1}^{n} y_{1i}^{2}\right) = n(\mu_{1}^{2} + \sigma_{1}^{2}),$$

$$t_{2}^{(t)} = E\left(\sum_{i=1}^{n} y_{2i}\right) = n\mu_{2},$$

$$t_{22}^{(t)} = E\left(\sum_{i=1}^{n} y_{2i}^{2}\right) = n(\mu_{2}^{2} + \sigma_{2}^{2}),$$

$$t_{12}^{(t)} = E\left(\sum_{i=1}^{n} y_{1i}y_{2i}\right) = n(\sigma_{12} + \mu_{1}\mu_{2}).$$

Example: bivariate normal data with one variable subject to missingness

$$\begin{split} \mu_1^{(t+1)} &= \frac{t_1^{(t)}}{n}, \\ \left(\sigma_1^{(t+1)}\right)^2 &= \frac{t_{11}^{(t)}}{n} - (\mu_1^{(t+1)})^2, \\ \mu_2^{(t+1)} &= \frac{t_2^{(t)}}{n}, \\ \left(\sigma_2^{(t+1)}\right)^2 &= \frac{t_{22}^{(t)}}{n} - (\mu_2^{(t+1)})^2, \\ \sigma_{12}^{(t+1)} &= \frac{t_{12}^{(t)}}{n} - \mu_1^{(t+1)} \mu_2^{(t+1)}. \end{split}$$

 \hookrightarrow Note that $t_1^{(t)}$, $t_{11}^{(t)}$, $\mu_1^{(t+1)}$, and $\sigma_1^{(t+1)}$ are constant across iterations because there are no missing values in Y_1 .