## Méthodes d'analyse biostatistique projet 1

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**TEXT** 

## 0.1 Exercise 1

Fix  $\alpha > 0$ . For a  $\theta \in (0, \alpha)$ , let  $\{X_i\}_{i=1}^n$  be a sequence of real independent identically distributed random variable defined on some probability space  $(\Omega, F, P)$  with common probability density function with respect to the Lebesgue measure:

$$f_{\theta}(x) = \begin{cases} \frac{2x}{\alpha\theta} & x \in [0, \theta] \\ \frac{2(\alpha - x)}{\alpha(\alpha - \theta)} & x \in [\theta, \alpha] \\ 0 & \text{otherwise} \end{cases}$$

Prove that the maximum likelihood estimation of  $\theta$  must be one of the given observation but not necessarily any particular observation. In case  $\alpha=5$  and n=3, compute the maximum likelihood estimate of  $\theta$  when the observations are (1,2,4) or (2,3,4).

**Solution** Let  $x = (x_1, \dots, x_n) \in \mathbb{R}^n$  be given. Write  $g_{\theta}(x)$  for the joint probability density function of  $X_i$ 's and  $x_{(i)}$  for the *i*th smallest coordinate in x. If  $x_i < 0$  or  $x_i > \alpha$  for some i, then  $g_{\theta}(x) = 0$  for any  $\theta$  so that any estimate would be a maximum likelihood estimate. We exclude this pathology and prove that:

**Théorème 1.** If  $0 \leqslant x_{(1)} \leqslant \cdots \leqslant x_{(n)} \leqslant \alpha$ , then  $\theta_0 = x_{(i)}$  for some i.

*Proof.* It never happens that  $\theta_0 < x_{(1)}$  because  $g_{\theta_1}(x) > g_{\theta_0}(x)$  whenever  $\theta_1 \in (\theta_0, x_{(1)})$ . Similarly, it never happens that  $\theta_0 > x_{(n)}$  because  $g_{\theta_1}(x) > g_{\theta_0}(x)$  whenever  $\theta_1 \in (x_{(n)}, \theta_0)$ . We assume from now on that  $\theta_0 \in [x_{(i)}, x_{(i+1)}]$  for some  $i \in \{1, \dots, n-1\}$ . Suppose, for the sake of contradiction, that  $x_{(i)} < \theta_0 < x_{(i+1)}$ . We have:

$$g_{\theta_{\theta_0}}(x) = \left(\frac{2}{\alpha}\right)^n \frac{x_{(1)}}{\theta_0} \cdots \frac{x_{(i)}}{\theta_0} \frac{\alpha - x_{(i+1)}}{\alpha - \theta_0} \cdots \frac{\alpha - x_{(n)}}{\theta_0}$$

The numerator does not depend on  $\theta_0$ . This motivates us to define function  $h:[x_{(i)},x_{(i+1)}]\to\mathbb{R}$  by:

$$h(\theta) = \frac{1}{\theta^i} \frac{1}{(\alpha - \theta)^{n-i}}$$

Then the second derivative is:

$$h''(\theta) = i(i+1)\theta^{-i-2}(\alpha - \theta)^{i-n} + (n-i)(n-i+1)\theta^{-i}(\alpha - \theta)^{i-n-2} > 0$$

Therefore, h is strictly convex and the maximum can only be at the boundary points.

We now demonstrate that the choice of i is not unique in the above theorem. The simplest case will be  $x_i = x_j$  for any i and any j. For a nontrivial example, let  $\alpha = 5$ , n = 3. If x = (2, 3, 4), then the maximum likelihood estimate is one of  $\{2, 3, 4\}$ . An estimate of 3 or 4 yields maximum likelihood  $\frac{8}{375}$  while an estimate of 2 yields likelihood  $\frac{16}{1125}$ . Therefore, the maximum likelihood estimate can be 3 or 4 and is not unique.

Finally, the additional example x=(1,2,4), estimate  $\theta=1,2,4$  gives likelihood  $\frac{3}{250},\frac{4}{375},\frac{1}{125}$  repsectively. We conclude that  $\theta=1$  is the maximum likelihood estimate in this case.///

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