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# Finite mixtures analysis by biased samples

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## Outline

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## 1. Motivating example

Imagine that the distribution of some characteristic  $\xi$  (e.g. body length) of crabs living at a sea is studied. The investigated population of crabs is divided into two sub-populations (components). The crabs belonging to the first component are more salt-loving then the ones belonging to the second component.



Component 1



Component 2

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Proportion in the local population depends on the mean salinity of the water at this site. Assume that the function describing the dependence of this proportion from the salinity is known.

We are interested in the differences in distribution of  $\xi$  for crabs belonging to different components. But the true component to which the crab belongs is not observed in the study, since it needs some expensive and time consuming tests. Therefore our inference should be based on the proportions of components at the sites. These proportions can be considered as probabilities that a crab chosen at random from the local population belongs to a given component (mixing probabilities). They can be estimated by the mean salinity data.

To catch the crabs some traps are used and it is known that the probabilities to be caught are different for crabs with different body length  $\xi$ . This causes a sampling bias in the observed distribution of  $\xi$ .



Our aim is to correct this bias and to extract the CDF of the component of interest from the mixture.

#### 2. Horwitz-Thompson approach to bias correction

Assume that there is a homogeneous infinite (very large) population of subjects O with the observed feature  $\xi(O) \in \mathbb{R}$ . Denote F(x) the CDF of  $\xi(O)$  in the entire population. A subject O can be sampled from the population with the probability depending on  $\xi(O)$  but independently of all other subjects. Let us denote this (inclusion) probability by

$$cq(t) = \mathsf{P}\{O \text{ was included to the sample } |\xi(O) = t\},$$
 (1)

where q(t) is a known function, c is an unknown constant.

The values of  $\xi(O)$  for the sampled subjects produce the sample  $Y = (\eta_1, \eta_2, \dots, \eta_n)$ .  $\eta_i$  are i.i.d. and the their CDF  $\tilde{F}(x)$  is the conditional probability of the event  $\{\xi(O) < x\}$  given that O was sampled.

Then

$$\tilde{F}(x) = \mathsf{P}\{\xi(O) < x \mid O \text{ was sampled}\} = \frac{\int_{-\infty}^{x} q(t)F(dt)}{\int_{-\infty}^{+\infty} q(t)F(dt)}.$$
 (2)

In this case the population mean  $\bar{\xi} = \mathsf{E}\,\xi = \int xF(dx)$  does not equal to the expectation of the observed values  $\mathsf{E}\,\eta_j$  due to the sampling bias. But  $\bar{\xi}$  can be estimated by the weighted sample mean with weights reciprocal to the inclusion probabilities:

$$\hat{\xi} = \frac{1}{\sum_{j=1}^{n} \frac{1}{q(\eta_j)}} \sum_{j=1}^{n} \frac{1}{q(\eta_j)} \eta_j.$$

It is the usual HT-estimate which is consistent if  $\overline{\xi}$  exists and q(t) > const > 0 for all t. The corresponding estimate for CDF F is

$$\hat{F}^{HT}(x) = \frac{1}{\sum_{j=1}^{n} \frac{1}{q(\eta_j)}} \sum_{j=1}^{n} \frac{1}{q(\eta_j)} \mathbb{I}\{\eta_j < x\}.$$
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#### 3. Mixtures with varying concentrations (MVC)

In the MVC model we assume that the subjects can belong to one of M sub-populations (components)  $\Re_1, \Re_2, \ldots, \Re_M$ . The probability to observe a subject from a given component depends on the conditions of the observations and is different for different observations. Let us denote by  $p_j^i$  the probability to observe a subject from  $\Re_i$  in the *j*-th observation. The CDF of the observed feature  $\xi(O)$  of a subject O is :

 $H_m(x) = \mathsf{P}\{\xi(O) < x \mid O \in \Re_m\}.$ 

So, the observed sample X consists of independent but not identically distributed observations  $X = (\xi_1, \ldots, \xi_n)$  with CDFs

$$F_j(x) = \mathsf{P}\{\xi_j < x\} = \sum_{m=1}^M p_j^m H_m(x).$$
(3)

Note that the component to which an observed subject belongs is unknown. One needs to infer on  $H_m$  only by the sample X and the set of mixing probabilities  $p_j^m$ , j = 1, ..., n, m = 1, ..., M which are known.

The CDF  $H_m$  may be estimated by a weighted empirical CDF

$$\hat{H}_m(x) = \frac{1}{n} \sum_{j=1}^n a_j^m \mathbb{I}\{\xi_j < x\}.$$

Here  $a_j^m$  are some weights which may depend on mixing probabilities  $p_j^i$ , but not on the observations  $\xi_j$ .

To obtain unbiased estimates one needs the following conditions to be satisfied:

$$\frac{1}{n} \sum_{j=1}^{n} a_j^m p_j^i = \mathbb{I}\{i=m\} \text{ for all } i=1,\dots,M$$

One of appropriate choices is the minimax weighting with

$$a_j^m = \sum_{i=1}^M \bar{\gamma}_{im} p_j^i,$$

where  $\bar{\Gamma} = (\bar{\gamma}_{im})_{i,m=1}^{M}$  is the matrix inverse to  $\Gamma = (\frac{1}{n} \sum_{j=1}^{n} p_j^i p_j^k)_{i,k=1}^{M}$ . To estimate the *m*-th component mean  $\bar{\xi}_m = \int x H_m(dx)$  one may use

$$\hat{\xi}_m = \frac{1}{n} \sum_{j=1}^n a_j^m \xi_j.$$

### 4. Mixture model with sampling bias

Let us assume now that the MVC model is in force together with the sampling bias. The considered subjects O belong to M different components and the CDF of their feature of interest  $\xi(O)$  is  $H_m$  for the subjects belonging to the m-th component. The proportion of the m-th component subjects in the local population from which the j-th subject was obtained is  $p_j^m$ . These probabilities are known. The CDFs  $H_m$  are unknown. The probability to sample the subject O from a local population depends on  $\xi(O)$ . This probability is defined by (1) with known q and unknown c.

The problem is to estimate the components' CDFs  $H_m$  and means  $\bar{\xi}_m = \int x H_m(dx)$ .

Analogously to (2) we obtain that the observed sample  $Y = (\eta_1, \ldots, \eta_n)$  consists of independent observations with CDFs

$$\tilde{F}_j(x) = \mathsf{P}\{\eta_j < x\} = \frac{\int_{-\infty}^x q(t)F_j(dx)}{\int_{-\infty}^{+\infty} q(t)F_j(dx)},\tag{4}$$

where  $F_j$  is defined by (3):  $F_j(x) = \sum_{m=1}^M p_j^m H_m(x)$ . From (4) we obtain

$$\tilde{F}_j(x) = \sum_{m=1}^M \frac{p_j^m \tilde{Q}_m}{\sum_{i=1}^M p_j^i \tilde{Q}_i} \tilde{H}_m(x),$$

where

$$\tilde{Q}_m = \int_{-\infty}^{\infty} q(t) H_m(dx), \ \tilde{H}_m(x) = \int_{-\infty}^{x} q(t) H_m(dx) / \tilde{Q}_m.$$

So, the sampling bias causes changes not only in the distributions of components, but also in the mixing probabilities.  $^{15}$ 

To take in account the bias in the mixing probabilities, we need to estimate  $\tilde{Q}_m$ . Notice that

$$\mathsf{E}\,\frac{1}{q(\eta_j)} = \frac{1}{\sum_{i=1}^M p_j^i \tilde{Q}_m}$$

The least squares technique suggests the estimate  $\hat{Q} = (\hat{Q}_1, \dots, \hat{Q}_M)$ for  $\tilde{Q} = (\tilde{Q}_1, \dots, \tilde{Q}_M)$  which is the minimizer of the LS functional

$$J(Q) = \sum_{j=1}^{n} \left( \frac{1}{\sum_{i=1}^{M} p_j^i Q_i} - \frac{1}{q(\eta_j)} \right)^2$$

over all  $Q = (Q_1, \ldots, Q_M)$  with  $Q_i > 0$ .

With these estimates at hands we define the weights  $\tilde{a}_j^m$  for the *m*-th component estimation as

$$\tilde{a}_{j}^{m} = \frac{1}{q(\eta_{j})} \sum_{k=1}^{M} \tilde{\gamma}_{km} \frac{p_{j}^{k}}{\sum_{i=1}^{M} p_{j}^{i} \hat{Q}_{i}},$$

where  $\hat{\Gamma}_Q = (\hat{\gamma}_{km})_{k,m=1}^M$  is the matrix inverse to

$$\tilde{\Gamma}_Q = \left(\frac{1}{n} \sum_{j=1}^n \frac{p_j^k p_j^m}{\left(\sum_{i=1}^M p_j^i \hat{Q}_i\right)^2}\right)_{k,m=1}^M$$

The resulting bias-correcting estimate for the  $H_m$  is

$$\hat{H}_{m}^{BC}(x) = \frac{1}{n} \sum_{j=1}^{n} \tilde{a}_{j}^{m} \mathbb{I}\{\eta_{j} < x\}.$$

The estimate for  $\bar{\xi}_m$  is

$$\hat{\xi}_m^{BC} = \frac{1}{n} \sum_{j=1}^n \tilde{a}_j^m \eta_j.$$

**Theorem 1** Assume that the next conditions holds.  
1) There exists 
$$c > 0$$
 such that  $q(x) > c$  for all  $x \in R$ .  
2) For all  $m = 1, ..., M$ :  $\int x^2 H_m(dx) < \infty$ .  
3) There is no linear subspace  $L \subset R^M$  with dim $L < M$  such that  $P\{(p_j^1, ..., p_j^M) \in L\} = 1$ .  
Then

$$\hat{\xi}_m^{BC} o \bar{\xi}_m$$

in probability as  $n \to \infty$ .

In this theorem we consider the concentrations  $\vec{p}_j = (p_j^1, \ldots, p_j^M)$  as generated by some stochastic mechanism. So  $\vec{p}_j$  are i.i.d. random vectors and formula (3) describes the conditional probability of  $\{\xi_j < x\}$  given  $\vec{p}_j$ :

$$F_j(x) = P\{\xi_j < x | \vec{p}_j\} = \sum_{j=1}^M p_j^i H_i(x).$$

#### 5. Simulation results

In our experiment we considered a two component mixture M = 2. The distribution of the first component was N(-1, 1), the distribution of the second one was N(1, 1).

The concentrations of the first component  $p_j^1$  were simulated as random variables, uniformly distributed on  $[0, 1], p_j^2 = 1 - p_j^1$ .

Figure 1 presents the graphs of the estimates  $\hat{H}_m^{BC}(x)$  for the components CDFs by a sample with n = 1000 observations.



Figure 1: Estimates of CDF for the first (a) and second (b) component. The true CDFs are depicted by black lines.

The biases and variances of the estimates  $\hat{\xi}_m^{BC}$  for different sample sizes n are presented in Table 1.

n	$\xi_1^{BC}$		$\xi_2^{BC}$	
	bias	Var	bias	Var
50	-0.0807	0.8011	-0.0851	0.3360
100	-0.0581	0.2257	-0.0589	0.1575
250	-0.0176	0.0849	-0.0271	0.0815
500	-0.045	0.0464	0.00210	0.0330
750	-0.0285	0.0421	-0.0162	0.0187
1000	-0.0052	0.0211	-0.0034	0.0118

Table 1: Performance of the estimates for means

#### References

1. Lohr, S. (2010) Sampling: Design and Analysis. Brooks/Cole.

2. Maiboroda, R. & Sugakova, O.(2012) Statistics of mixtures with varying concentrations with application to DNA microarray data analysis. Nonparametric statistics, **24**, iss.1, 201-215.

3. Shao, J. (2003) Mathematical statistics. Shpringer.