

# *Loci selection in model-based clustering*

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# Introduction

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- We wish to discover the unknown genetic structure of a target diploid population from a  $n$ -sample without prior information.
- It may happen that some loci are just noise or even harmful for clustering purposes.
- Which loci cluster the sample in the "best" way?
- We propose to simultaneously solve the loci selection and clustering problem by a model selection procedure for density estimation.
- An associated stand alone C++ package named MixMoGenD is available free of charge on [www.math.u-psud.fr/~toussile](http://www.math.u-psud.fr/~toussile).

# Outline

## 1 Methods

- Competing models
- Model selection via penalization

## 2 Consistency

## 3 Selection procedure

- Selection procedure in practice
- Numerical experiments using BIC

# Methods

## Framework

- Consider a random vector  $X = (X^l)_{l=1,\dots,L}$  with  $L \geq 2$ .
- With  $X^l = \{X^{l,1}, X^{l,2}\}$ , where  $X^{l,1}, X^{l,2}$  are nominal variables taking values in the set  $\{1, \dots, A_l\}$  of allele states at locus  $l$ .

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  - Assume that the clusters are characterized by:
    - (LE) Conditional complete independence of the random variables  $X^l$ ;
    - (HWE) Conditional independence of  $X^{l,1}$  and  $X^{l,2}$  at any locus  $X^l$ .
- [Pritchard et al., 2000, Chen et al., 2006, Corander et al., 2008].

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  - (HWE) Conditional independence of  $X^{l,1}$  and  $X^{l,2}$  at any locus  $X^l$ .[Pritchard et al., 2000, Chen et al., 2006, Corander et al., 2008].
- Now, assume that only some loci gathered in a subset  $S$  are relevant for clustering purposes.
- Also assume that for any  $l \notin S$ ,  $X^l$  is identically distributed across all clusters.

# Modeling

## Competing models

- $\Rightarrow$  In model-based settings,  $X \sim P_0$  of the form

$$P_{(K,S,\theta)}(x) = \left[ \sum_{k=1}^K \pi_k \prod_{l \in S} (2 - \mathbb{1}_{x^{l,1}=x^{l,2}}) \alpha_{k,l,x^{l,1}} \times \alpha_{k,l,x^{l,2}} \right] \\ \times \prod_{l \notin S} (2 - \mathbb{1}_{x^{l,1}=x^{l,2}}) \beta_{l,x^{l,1}} \beta_{l,x^{l,2}} \quad (1)$$

where  $\theta = (\pi, \alpha, \beta) \in \Theta_{(K,S)} = \dots$ .

- Model  $\mathcal{M}_{(K,S)} := \{P_{(K,S,\theta)} \mid \theta \in \Theta_{(K,S)}\}$ .
- Inferring  $(K, S) \iff$  model selection among  $\mathcal{C} = \{\mathcal{M}_{(K,S)} \mid (K, S) \in \mathbb{M}\}$  for the estimation of  $P_0$ , where  $\mathbb{M}$  is the set of all possible  $(K, S)$ .



# Methods

Model selection via penalization ([Massart, 2007])

- Selected model

$$\left(\widehat{K}_n, \widehat{S}_n\right) = \arg \min_{(K, S)} \mathbf{crit}(K, S). \quad (2)$$

- Where **crit** is a penalized maximum likelihood criterion

$$\mathbf{crit}(K, S) = \underbrace{\gamma_n \left(\widehat{P}_{(K, S)}\right)}_{\mathbb{P}_n(-\ln \widehat{P}_{(K, S)})} + \mathbf{pen}(K, S);$$
$$\mathbb{P}_n(-\ln \widehat{P}_{(K, S)}) := \frac{1}{n} \sum_{i=1}^n -\ln P_{(K, S, \widehat{\theta}_{MLE})}(X_i) \quad (3)$$

- Selected estimator  $P_{(\widehat{K}_n, \widehat{S}_n, \widehat{\theta}_{MLE})}$  and classification by MAP.

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- Selected estimator  $P_{(\widehat{K}_n, \widehat{S}_n, \widehat{\theta}_{MLE})}$  and classification by MAP.
- The most used asymptotic penalized likelihood criteria:

$$\mathbf{BIC}(K, S) = \mathbb{P}_n \left(-\ln \widehat{P}_{(K, S)}\right) + \frac{\ln n}{2n} D_{(K, S)}$$

$$\mathbf{AIC}(K, S) = \mathbb{P}_n \left(-\ln \widehat{P}_{(K, S)}\right) + \frac{1}{n} D_{(K, S)}.$$



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# Consistency of the BIC like criteria

- Although there exists a lot of articles concerning the behavior of the BIC and other penalization methods in practice, theoretical results in a mixture framework are few: the consistency of the BIC estimator is shown
  - ▶ in [Maugis et al., 2009] for a variable selection problem,
  - ▶ and in [Keribin, 2000] for the number of components, in Gaussian mixture models framework.
- But as far as we know, there is no consistency result for both a variable selection and clustering problem in a discrete distribution setting.

## Consistency of the BIC like criteria

- Consider a penalty function  $\mathbf{pen} = \mathbf{pen}(D, n)$  such that:
  - ▶ (P1): for any positive integer  $D$ ,  $\lim_{n \rightarrow \infty} \mathbf{pen}(D, n) = 0$ ;
  - ▶ (P2): for any  $\mathcal{M}_1 \subsetneq \mathcal{M}_2$ , one has

$$\lim_{n \rightarrow \infty} \left[ n \left( \mathbf{pen}(D_2, n) - \mathbf{pen}(D_1, n) \right) \right] = \infty.$$

- Let  $(\hat{K}_n, \hat{S}_n)$  be a minimizer of **crit** over a sub-collection  $\mathcal{C}_{K_{\max}}$  for a given maximum number  $K_{\max}$  of clusters.

### Theorem ([Toussile and Gassiat, 2009])

*If  $P_0 > 0$  and belongs to one of the competing models in  $\mathcal{C}_{K_{\max}}$ , then there exists an identifiable “smallest” model  $(K_0, S_0)$  such that*

$$\lim_{n \rightarrow \infty} P_0 \left[ (\hat{K}_n, \hat{S}_n) = (K_0, S_0) \right] = 1. \quad (5)$$

- Example: BIC.

# Consistency of the BIC like criteria

Definition of  $\mathcal{M}_{(K_0, S_0)}$

## Lemma

For every  $K_1$  and  $K_2$  in  $\mathbb{N}^*$ , and  $S_1$  and  $S_2$  in  $\mathcal{P}^*(L)$ ,  
 $\mathcal{M}_{(K_1, S_1)} \cap \mathcal{M}_{(K_2, S_2)} = \mathcal{M}_{(\min(K_1, K_2), S_1 \cap S_2)}$ .

The "smallest" model is defined by  $(K_0, S_0) := (K(P_0), S(P_0))$ ,  
where

$$K(P) = \min \left\{ K \mid P \in \bigcup_{S \in \mathcal{P}^*(L)} \mathcal{M}_{(K, S)} \right\}, \quad (6)$$

$$S(P) = \min \left\{ S \mid P \in \bigcup_{K \in \mathbb{N}^*} \mathcal{M}_{(K, S)} \right\}, \quad (7)$$

for every  $P$  in one of the competing models  $\mathcal{M}_{(K, S)} \in \mathcal{C}_{K_{\max}}$ .

# Consistency of the BIC like criteria

## Proof

It suffices to show that  $\lim_{n \rightarrow \infty} P_0 \left[ \gamma_n \left( \hat{P}_{(K_0, S_0)} \right) - \gamma_n \left( \hat{P}_{(K, S)} \right) > \mathbf{pen}(K, S) - \mathbf{pen}(K_0, S_0) \right] = 0$  for any  $(K, S) \neq (K_0, S_0)$ .

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①  $P_0 \in \mathcal{M}_{(K, S)}$ :

②  $P_0 \notin \mathcal{M}_{(K, S)}$ :



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①  $P_0 \in \mathcal{M}_{(K, S)}$ :

$$-m\gamma_n(P_0) \leq -m\gamma_n \left( \hat{P}_{(K_0, S_0)} \right) \leq -m\gamma_n \left( \hat{P}_{(K, S)} \right) \leq \sup_{P \in \mathcal{D}} (-m\gamma_n(P)).$$

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②  $P_0 \notin \mathcal{M}_{(K, S)}$ :

$$\begin{aligned} & \gamma_n\left(\hat{P}_{(K_0, S_0)}\right) - \gamma_n\left(\hat{P}_{(K, S)}\right) = \\ & -\inf_{\theta \in \Theta_{(K, S)}^\delta} E_{P_0} \left[ \ln P_0(X) - \ln P_{(K, S)}(X | \theta) \right] + o_{P_0}(1), \\ & \text{where } \Theta_{(K, S)}^\delta = \{ \theta \in \Theta_{(K, S)} : P_{(K, S, \theta)} \geq \delta \} \end{aligned}$$

# Consistency of the BIC like criteria

Proof

Theorem ([Toussile and Gassiat, 2009])

If  $P_0 > 0$ , there exists a real  $\delta > 0$  such that for every  $(K, S)$ , one has

$$-\gamma_n \left( \widehat{P}_{(K,S)} \right) = \sup_{\theta \in \Theta_{(K,S)}^\delta} \left\{ -\gamma_n \left( P_{(K,S,\theta)} \right) \right\} + o_{P_0}(1) \quad (8)$$

and

$$\sup_{\theta \in \Theta_{(K,S)}} E_{P_0} \left[ \ln P_{(K,S,\theta)}(X) \right] = \sup_{\theta \in \Theta_{(K,S)}^\delta} E_{P_0} \left[ \ln P_{(K,S,\theta)}(X) \right]. \quad (9)$$

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# Selection procedure in practice

- An exhaustive search of the optimum model is very painful in most situations.
- A two nested algorithm based on Backward-Stepwise proposed in [Maugis et al., 2009] could miss the optimum model in some cases, in particular in cases where the optimum subset of clustering loci is small.
- In **MixMoGenD**, we prefer a modified Backward-Stepwise algorithm with which sets  $S$  with small cardinality are always explored for any value of  $K$  [Toussile and Gassiat, 2009].
- The optimum model is then chosen between all the explored models.

## BACKWARD-STEPWISE EXPLORER(**crit**, $K$ )

```

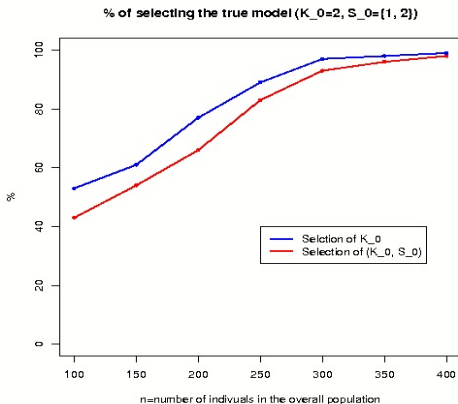
1   $S \leftarrow \{1, \dots, L\}$ ,  $c_{ex} \leftarrow 0$ ,  $c_{in} \leftarrow 0$ 
2  repeat
3      EXCLUSION( $K, S$ ) {
4           $c_{ex} \leftarrow \arg \min_{I \in S} \mathbf{crit}(K, S \setminus \{I\})$ 
5          if  $\mathbf{crit}(K, S) - \mathbf{crit}(K, S \setminus \{c_{ex}\}) \geq 0$  or  $c_{in} = 0$ 
6              then  $S \leftarrow S \setminus \{c_{ex}\}$ 
7          }
8      INCLUSION( $K, S$ ) {
9           $c_{in} \leftarrow \arg \min_{I \notin S} \mathbf{crit}(K, S \cup \{I\})$ 
10         if  $(\mathbf{crit}(K, S \cup \{c_{in}\}) - \mathbf{crit}(K, S) < 0$  and  $S \cup \{c_{in}\}$  has
11             never been the current set in an EXCLUSION step)
12             then  $S \leftarrow S \cup \{c_{in}\}$ 
13             else  $c_{in} \leftarrow 0$ 
14         }
15     until  $|S| = 1$ .

```

# Numerical experiments using BIC

## Consistency

Figure: Percentage of selecting the true model using the BIC



# Numerical experiments using BIC

- $L = 10, A_l = 10, K_0 = 5, |S_0| \in \{2, 4, 6, 8\}$ .
- 30 datasets with  $n = 1000$  for each value of  $|S_0|$ .
- $F_{ST} \in [0.0181, 0.0450]$  a range where clustering is thought to be difficult.

**Table:** Thresholds of  $F_{ST}$  for which MixMoGenD perfectly selects the true model.  $F_{ST}^S$ : with loci selection;  $F_{ST}$ : without loci selection.

$ S_0 $	8	6	4	2
$F_{ST}^S$	0.0342	0.0307	0.0316	0.0248
$F_{ST} >$	0.0425	0.0410	0.0413	0.0350

- The improvement on the estimation of  $K$  and the prediction capacity is obviously due to the variable selection procedure.



# Numerical experiments using BIC

Data	$F_{ST}$	$\hat{K}_n$	% WA	$\hat{K}_n^s$	% WA <sup>s</sup>	Data	$F_{ST}$	$\hat{K}_n$	% WA	$\hat{K}_n^s$	% WA <sup>s</sup>
1	0.0306	3	-	3	-	16	0.0381	5	10.90	5	10.30
2	0.0318	3	-	3	-	17	0.0382	5	09.30	5	08.80
3	0.0328	3	-	3	-	18	0.0390	4	-	5	09.10
4	0.0331	3	-	3	-	19	0.0400	5	08.80	5	08.00
5	0.0335	3	-	4	-	20	0.0404	4	-	5	09.50
6	0.0337	3	-	3	-	21	0.0425	5	06.30	5	05.40
7	0.0340	4	-	4	-	22	0.0427	5	07.10	5	07.50
8	0.0342	3	-	5	11.80	23	0.0427	5	05.90	5	05.90
9	0.0348	3	-	5	12.40	24	0.0435	5	06.70	5	06.50
10	0.0362	3	-	5	09.10	25	0.0436	5	07.10	5	06.60
11	0.0373	4	-	5	08.90	26	0.0440	5	05.50	5	05.70
12	0.0373	5	08.50	5	07.60	27	0.0442	5	07.20	5	06.80
13	0.0377	5	11.40	5	10.40	28	0.0449	5	07.20	5	06.70
14	0.0377	5	10.50	5	10.20	29	0.0449	5	06.10	5	06.30
15	0.0377	5	10.30	5	10.20	30	0.0450	5	06.10	5	05.60

**Table:** 30 samples each with  $n = 1\,000$ ,  $K_0 = 5$ ,  $L = 10$ ,  $|S_0| = 8$  and  $F_{ST} \in [0.0306, 0.0450]$ . % WA and % WA<sup>s</sup> = percentage of wrongly assigned individuals without and with loci selection respectively;  $\hat{K}_n$  and  $\hat{K}_n^s$  = the estimates of the number of populations without and with loci selection respectively.  $\hat{S}_n = S_0$ .

## Conclusion and perspectives

- Theoretical result on the consistency of the **BIC** type criteria is also valid for the variable selection problem in clustering with multinomial mixture models.
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  - As expected, the variable selection procedure significantly improves the inference on the number of clusters and the prediction capacity.
- 
- Robustness of the selection procedure with respect to HWE and LE assumptions.
  - Is it the same set  $S$  of loci that discriminates all populations?
  - **BIC**, as well as **AIC**, relies on a strong asymptotic assumption, and can thus be inappropriate for small sample sizes.



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