

An Approximate Fisher Scoring Algorithm for Finite Mixtures of Multinomials

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(www.umbc.edu/hpcf)

Background

- Morel and Neerchal (1991, 1993, 1998, 2005) proposed a multinomial model for overdispersion — Random Clumped Multinomial — and studied estimation under it.
- They obtained a large cluster approximation to the Fisher Information Matrix (FIM), and used it to formulate an Approximate Fisher Scoring Algorithm (AFSA).
- Liu (2005, PhD Thesis) extended the idea to general mixtures of multinomials, and found some interesting connections between AFSA and Expectation Maximization (EM).
- This work extends Liu (2005), further investigating the quality of the FIM approximation and the connection between AFSA and EM.

Mixture of Multinomials Example

Example: Housing satisfaction survey

Non-metropolitan area				Metropolitan area			
Neighborhood	US	S	VS	Neighborhood	US	S	VS
1	3	2	0	19	0	4	1
2	3	2	0	20	0	5	1
3	0	5	0	21	0	3	2
\vdots				\vdots			
17	4	1	0	35	4	1	0
18	5	0	0				

With labels, a reasonable likelihood is product of two multinomials

$$L(\theta) = \left[\prod_{i=1}^{18} f(\mathbf{x}_i \mid \mathbf{p}_1, m) \right] \left[\prod_{i=19}^{35} f(\mathbf{x}_i \mid \mathbf{p}_2, m) \right], \quad m = 5.$$

J. R. Wilson, Chi-Square Tests for Overdispersion with Multiparameter Estimates.
Journal of the Royal Statistical Society (Series C), 38(3):441–453, 1989.

Mixture of Multinomials Example

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Neighborhood	US	S	VS	Neighborhood	US	S	VS
1	3	2	0	19	0	4	1
2	3	2	0	20	0	5	1
3	0	5	0	21	0	3	2
\vdots				\vdots			
17	4	1	0	35	4	1	0
18	5	0	0				

Without labels, a reasonable likelihood is mixture of two multinomials

$$L(\theta) = \prod_{i=1}^{35} \left\{ \pi f(\mathbf{x}_i \mid \mathbf{p}_1, m) + (1 - \pi) f(\mathbf{x}_i \mid \mathbf{p}_2, m) \right\}, \quad m = 5.$$

J. R. Wilson, Chi-Square Tests for Overdispersion with Multiparameter Estimates.
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Mixture of Multinomials

- Suppose we have s multinomial populations

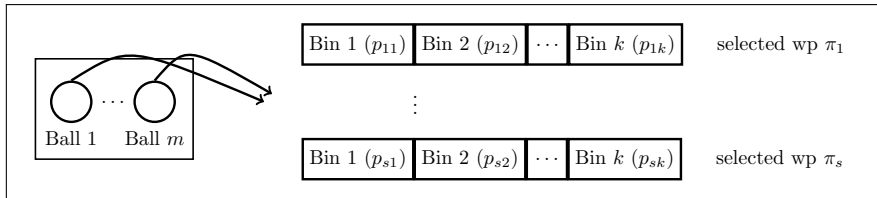
$$f(\mathbf{x} \mid \mathbf{p}_\ell, m) = \frac{m!}{x_1! \dots x_k!} p_{\ell 1}^{x_1} \dots p_{\ell k}^{x_k} \cdot I(\mathbf{x} \in \Omega), \quad \ell = 1, \dots, s$$

which occur in the total population with probabilities π_1, \dots, π_s .

- If we draw \mathbf{T} from the mixed population,

$$\mathbf{T} \sim f(\mathbf{x} \mid \boldsymbol{\theta}) = \sum_{\ell=1}^s \pi_\ell f(\mathbf{x} \mid \mathbf{p}_\ell, m), \quad \boldsymbol{\theta} = (\mathbf{p}_1, \dots, \mathbf{p}_s, \boldsymbol{\pi})$$

We'll write $\mathbf{T} \sim \text{MultMix}_k(\boldsymbol{\theta}, m)$.



Estimation Problem

- Suppose our sample is:

$$\mathbf{X}_i \stackrel{\text{ind}}{\sim} \text{MultMix}_k(\boldsymbol{\theta}, m_i), \quad i = 1, \dots, n.$$

Then likelihood is

$$L(\boldsymbol{\theta}) = \prod_{i=1}^n \left\{ \sum_{\ell=1}^s \pi_{\ell} \left[\frac{m_i!}{x_{i1}! \dots x_{ik}!} p_{\ell 1}^{x_{i1}} \dots p_{\ell k}^{x_{ik}} \cdot I(\mathbf{x}_i \in \Omega) \right] \right\}.$$

- To find MLE $\hat{\boldsymbol{\theta}} = (\hat{\mathbf{p}}_1, \dots, \hat{\mathbf{p}}_s, \hat{\boldsymbol{\pi}})$, which maximizes the log-likelihood
- No nice closed form, so use iterative methods
 - ▶ Expectation Maximization (EM)
 - ▶ Newton-Raphson, **Fisher Scoring**, Quasi-Newton methods

$$\boldsymbol{\theta}^{(g+1)} = \boldsymbol{\theta}^{(g)} - \alpha \mathbf{H}^{-1} S(\boldsymbol{\theta}^{(g)}), \quad g = 1, 2, \dots$$

Fisher Scoring Algorithm

- The iterations become

$$\boldsymbol{\theta}^{(g+1)} = \boldsymbol{\theta}^{(g)} + \mathcal{I}^{-1}(\boldsymbol{\theta}^{(g)})S(\boldsymbol{\theta}^{(g)}), \quad g = 1, 2, \dots,$$

but $\mathcal{I}(\boldsymbol{\theta})$ may not be easy to compute.

- Naive summation works when sample space Ω is small

$$\mathcal{I}(\boldsymbol{\theta}) := \sum_{\mathbf{x} \in \Omega} \left\{ -\frac{\partial^2}{\partial \boldsymbol{\theta} \partial \boldsymbol{\theta}^T} \log f(\mathbf{x} \mid \boldsymbol{\theta}) \right\} f(\mathbf{x} \mid \boldsymbol{\theta}).$$

- Monte Carlo approximation
- For large m , an approximate FIM (shown for $\mathbf{X}_1 \sim \text{MultMix}_k(\boldsymbol{\theta}, m)$) is

$$\tilde{\mathcal{I}}(\boldsymbol{\theta}) := \text{Blockdiag}(\pi_1 \mathbf{F}_1, \dots, \pi_s \mathbf{F}_s, \mathbf{F}_\pi),$$

$$\mathbf{F}_\ell = m \left[\text{Diag}(p_{\ell 1}^{-1}, \dots, p_{\ell, k-1}^{-1}) + p_{\ell k}^{-1} \mathbf{1} \mathbf{1}^T \right]$$

$$\mathbf{F}_\pi = \text{Diag}(\pi_\ell^{-1}, \dots, \pi_{s-1}^{-1}) + \pi_s^{-1} \mathbf{1} \mathbf{1}^T$$

Approximate FIM Properties I

- **Result:** $\tilde{\mathcal{I}}(\boldsymbol{\theta}) - \mathcal{I}(\boldsymbol{\theta}) \rightarrow \mathbf{0}$ as $m \rightarrow \infty$.
- **Result:** $\tilde{\mathcal{I}}(\boldsymbol{\theta})$ is “complete data” FIM of (\mathbf{X}, Z) . Assume that

$$Z = \begin{cases} 1 & \text{wp } \pi_1 \\ \vdots & \\ s & \text{wp } \pi_s, \end{cases} \quad \text{and} \quad (\mathbf{X} \mid Z = \ell) \sim \text{Mult}_k(\mathbf{p}_\ell, m).$$

Then we have $\tilde{\mathcal{I}}(\boldsymbol{\theta}) \equiv \mathbb{E} \left\{ -\frac{\partial^2}{\partial \boldsymbol{\theta} \partial \boldsymbol{\theta}^T} \log f(\mathbf{x}, z \mid \boldsymbol{\theta}) \right\}$

- Note that EM is based on maximizing

$$Q(\boldsymbol{\theta}, \boldsymbol{\theta}') = \mathbb{E}_{\boldsymbol{\theta}'} \left[\log f(\mathbf{x}, z \mid \boldsymbol{\theta}) \mid \mathbf{x} \right].$$

Approximate FIM Properties II

- $\tilde{\mathcal{I}}(\theta)$ is a block diagonal matrix of Multinomial FIMs.
 - ▶ Simple forms for inverse, trace, and determinant
- **Result:** $\tilde{\mathcal{I}}(\theta) - \mathcal{I}(\theta)$ is positive semi-definite.
 - ▶ Fairly general result for complete data FIMs, not just for multinomial mixtures
- **Result:** Standard errors derived from $\tilde{\mathcal{I}}^{-1}(\theta)$ are systematically too optimistic (small)
- **Result:** The inverses also converge

$$\mathcal{I}^{-1}(\theta) - \tilde{\mathcal{I}}^{-1}(\theta) \rightarrow \mathbf{0} \quad \text{as } m \rightarrow \infty.$$

Approximate FIM Properties III

Some recommendations for approximation FIM

- Large cluster size (m) needed for

$$\tilde{\mathcal{I}}(\boldsymbol{\theta}) \approx \mathcal{I}(\boldsymbol{\theta}) \quad \text{and} \quad \tilde{\mathcal{I}}^{-1}(\boldsymbol{\theta}) \approx \mathcal{I}^{-1}(\boldsymbol{\theta});$$

inverses appear to converge faster.

- Approximate FIM and inverse are not recommended for general inference.
- But useful as a tool for estimation, as we will see.

Approximate Fisher Scoring Algorithm

- Using the approximate FIM in place of the true FIM gives AFSA

$$\boldsymbol{\theta}^{(g+1)} = \boldsymbol{\theta}^{(g)} + \tilde{\mathcal{I}}^{-1}(\boldsymbol{\theta}^{(g)})S(\boldsymbol{\theta}^{(g)}), \quad g = 1, 2, \dots$$

until $|\log L(\boldsymbol{\theta}^{(g+1)}) - \log L(\boldsymbol{\theta}^{(g)})| < \varepsilon$.

- Result:** Under $\mathbf{X}_1, \dots, \mathbf{X}_n \stackrel{\text{iid}}{\sim} \text{MultMix}_k(\boldsymbol{\theta}, m)$, EM and AFSA iterations are “equivalent”, given the same starting place $\boldsymbol{\theta}^{(g)}$

$$\tilde{\pi}_{\ell}^{(g+1)} = \hat{\pi}_{\ell}^{(g+1)}, \quad \tilde{p}_{\ell j}^{(g+1)} = \left(\frac{\hat{\pi}_{\ell}^{(g+1)}}{\pi_{\ell}^{(g)}} \right) \hat{p}_{\ell j}^{(g+1)} + \left(1 - \frac{\hat{\pi}_{\ell}^{(g+1)}}{\pi_{\ell}^{(g)}} \right) p_{\ell j}^{(g)}.$$

- Don't get exact equality under the “independent but not iid” case.

Equivalence of AFSA and EM

- A more general connection is known between EM and iterations of the form

$$\boldsymbol{\theta}^{(g+1)} = \boldsymbol{\theta}^{(g)} + \mathcal{I}_c^{-1}(\boldsymbol{\theta}^{(g)}) S(\boldsymbol{\theta}^{(g)}), \quad g = 1, 2, \dots$$

- Titterton (1984) shows the two are approximately equivalent (under some regularity conditions)
- The equivalence is exact when the complete data likelihood is a regular exponential family

$$L(\boldsymbol{\mu}) = \exp \left\{ b(\mathbf{x}) + \boldsymbol{\eta}^T \mathbf{t} + a(\boldsymbol{\eta}) \right\},$$

$\boldsymbol{\mu} = E(\mathbf{t}(\mathbf{X}))$: the parameter of interest.

- For MultMix problem, equivalence is approximate, not exact.
 - ▶ AFSA's originally justified from $\tilde{\mathcal{I}}(\boldsymbol{\theta})$ and Blischke (1964).
 - ▶ But this result justifies AFSA for finite mixtures other than multinomial.

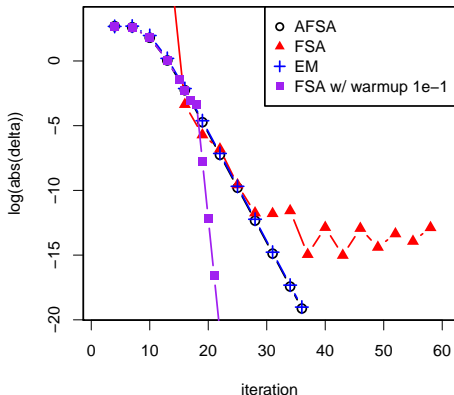
Comparison between algorithms

Consider the mixture of two trinomials

$$\mathbf{x}_i \stackrel{\text{iid}}{\sim} \text{MultMix}_3(\boldsymbol{\theta}, m = 20), \quad i = 1, \dots, n = 500$$

$$\begin{pmatrix} \mathbf{p}_1^T \\ \mathbf{p}_2^T \end{pmatrix} = \begin{pmatrix} 1/3 & 1/3 & 1/3 \\ 0.1 & 0.3 & 0.6 \end{pmatrix}, \quad \begin{pmatrix} \pi \\ 1 - \pi \end{pmatrix} = \begin{pmatrix} 0.75 \\ 0.25 \end{pmatrix}.$$

Convergence of competing algorithms



method	tol	iter
AFSA	4.94×10^{-09}	36
EM	5.50×10^{-09}	36

Monte Carlo Comparison of EM and AFSA

Consider a scenario with varying cluster sizes

$$\mathbf{Y}_i \stackrel{\text{ind}}{\sim} \text{MultMix}_k(\boldsymbol{\theta}, m_i), \quad i = 1, \dots, n = 500, \quad \boldsymbol{\pi} = (0.75, 0.25)$$

$$W_1, \dots, W_n \stackrel{\text{iid}}{\sim} \text{Gamma}(\alpha, \beta), \quad m_i = \lceil W_i \rceil.$$

Ran 1000 reps of nine scenarios and looked at the quantity

$$\bar{D} = \frac{1}{1000} \sum_{r=1}^{1000} D_r \quad \text{where} \quad D_r = \bigvee_{j=1}^q \left| \frac{\tilde{\theta}_j^{(r)} - \hat{\theta}_j^{(r)}}{\tilde{\theta}_j^{(r)}} \right|.$$

(kth probability not shown)		m_i equal	$\alpha = 100$	$\alpha = 25$
\mathbf{p}_1	\mathbf{p}_2	$m_i = 20$	$\text{Var}(m_i) \approx 4.083$	$\text{Var}(m_i) \approx 16.083$
0.1	0.5	2.178×10^{-6}	2.019×10^{-6}	2.080×10^{-6}
0.3	0.5	4.073×10^{-5}	3.501×10^{-5}	3.890×10^{-5}
0.35	0.5	8.683×10^{-4}	2.625×10^{-4}	2.738×10^{-4}
0.4	0.5	9.954×10^{-3}	6.206×10^{-2}	6.563×10^{-2}
(0.1, 0.3)	(1/3, 1/3)	1.342×10^{-3}	1.009×10^{-3}	1.878×10^{-3}
(0.1, 0.5)	(1/3, 1/3)	1.408×10^{-6}	1.338×10^{-6}	1.334×10^{-6}
(0.3, 0.5)	(1/3, 1/3)	3.884×10^{-6}	3.943×10^{-6}	3.885×10^{-6}
(0.1, 0.1, 0.3)	(0.25, 0.25, 0.25)	8.389×10^{-7}	8.251×10^{-7}	8.440×10^{-7}
(0.1, 0.2, 0.3)	(0.25, 0.25, 0.25)	1.523×10^{-6}	1.472×10^{-6}	1.408×10^{-6}

Conclusions

AFSA obtained as a Newton-type algorithm using approximate FIM.

- Nearly equivalent to EM iterations — similar solutions are obtained at similar rates of convergence.
- (EM advantage) M-step can be formulated so it won't wander outside parameter space.
- (AFSA advantage) May be easier to formulate when missing data structure is complicated.
E.g. Random-Clumped Multinomial (Morel & Neerchal 1993).

Result of Titterton (1984) suggests AFSA approach is reasonable for finite mixtures in general.

- What about the FIM approximation?

Both EM and AFSA suffer from a slow convergence rate.

- Hybrid is recommended for fast convergence and robustness.
- ...if true FIM is feasible to compute.

Conclusions

For details and references, see our technical report:

www.umbc.edu/hpcf \Rightarrow Research \Rightarrow Publications

A. M. Raim, M. Liu, N. K. Neerchal & J. G. Morel. An Approximate Fisher Scoring Algorithm for Finite Mixtures of Multinomials. Technical Report HPCF-2012-14, UMBC High Performance Computing Facility, University of Maryland, Baltimore County, 2012.

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How good is the FIM approximation?

Consider a mixture $\text{MultMix}_2(\theta, m)$ of three binomials, with parameters

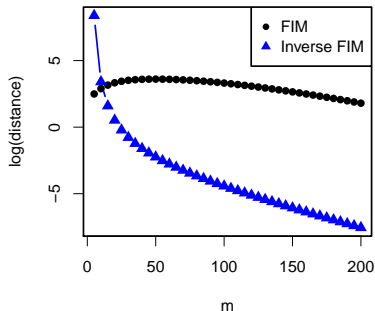
$$(p_1 \ p_2 \ p_3) = (1/7 \ 1/3 \ 2/3), \quad \pi = (1/6 \ 2/6 \ 3/6),$$

and two matrix distances

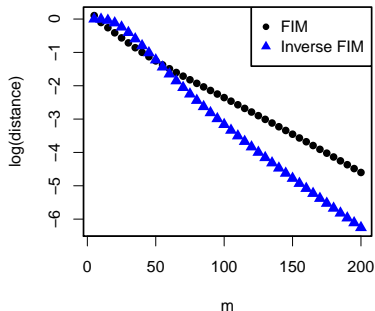
$$d(\mathbf{A}, \mathbf{B}) = \|\mathbf{A} - \mathbf{B}\|_F$$

$$d(\mathbf{A}, \mathbf{B}) = \frac{\|\mathbf{A} - \mathbf{B}\|_F}{\|\mathbf{B}\|_F}$$

**Log of Frobenius Distance
b/w Exact and Approx Matrices**



**Log of Scaled Frobenius Distance
b/w Exact and Approx Matrices**



Large m is needed for a good approximation. Inverses are converging faster.