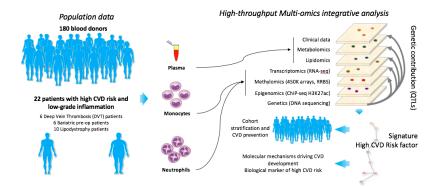
Alessandra Cabassi Dr Paul D. W. Kirk

24 April 2019





Multi-omic analysis of cardiovascular disease (CVD) risk data with Dr Denis Seyres and Dr Mattia Frontini – Department of Hæmatology



CVD risk data	Cell type	Variables	Observations
Epigenomics		25600	172
	0	26300	128
Methylomics		26214	193
	0	21442	187
Transcriptomics	۲	11370	203
	0	24224	198
Lipidomics	0	123	192
Metabolomics	0	988	200

We need:

- Scalable approximate inference method for mixture models
- That allows to combine different types of data
- And to perform feature selection

Why feature selection?

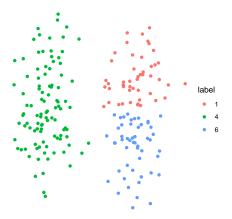
Example: Mixture of Gaussians



! Noisy features can degrade the performance of most learning algorithms Law et al. (2004)

Why feature selection?

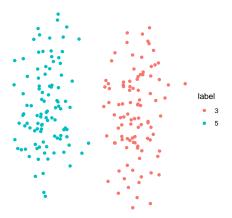
Example: Mixture of Gaussians fitted using both features



! Noisy features can degrade the performance of most learning algorithms Law et al. (2004)

Why feature selection?

Example: Mixture of Gaussians fitted using only the relevant feature



! Noisy features can degrade the performance of most learning algorithms Law et al. (2004)

Approximate inference Why?

Given a joint model for our hidden variables z and observed variables x, p(x, z), inference about the unknown is through the posterior

$$p(z|x) = \frac{p(z,x)}{p(x)}$$

For most interesting models, the denominator is not tractable, so we appeal to approximate posterior inference

Approximate inference

Which type?

Stochastic approximations \rightarrow Sampling

- + Asymptotically exact
- + Easily applicable general-purpose algorithms
- Computationally expensive
- Storage intensive

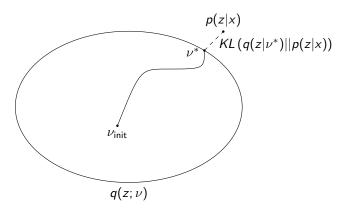
$\textbf{Deterministic approximations} \rightarrow \textbf{Structural assumptions}$

- + Computationally efficient
- + Efficient representation
- Often hard work to derive
- Not guaranteed to converge to global optimum

Brodersen (2010)

Main idea

Posit a variational family of distributions over the latent variables $q(z; \nu)$ and fit the variational parameters ν to be close (in Kullback-Leibler divergence) to the exact posterior



History

Variational inference (VI) adapts ideas from statistical physics to probabilistic inference

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History

Variational inference (VI) adapts **ideas from statistical physics** to probabilistic inference

 1980s: Peterson and Anderson (1987) used mean-field methods to fit a neural network
 Early 1990s: This idea was picked up by Jordan's lab who generalised it to many probabilistic models (a review paper is Jordan, Ghahramani, Jaakkola and Saul, 1999)

1 In parallel: Hinton and Van Camp (1993) developed mean-field for neural networks. Neal and Hinton (1993) **connected this idea to the EM algorithm**, which lead to further **variational methods for mixtures of experts** (Waterhouse et al., 1996)

Preliminary definitions

Entropy: (information theory) average rate at which information is produced by a stochastic source of data

Given a random variable x with probability density function p(x)

$$H(x) = -\int p(x) \log p(x) dx = \mathsf{IE}_p[-\log p(x)]$$

Entropy increases as the distribution becomes broader

Preliminary definitions

Kullback-Leibler divergence (relative entropy): measure of how one probability distribution is different from a second, reference probability distribution

p(z): unknown distribution q(z): approximating distribution

$$\mathsf{KL}(q||p) = -\int q(z) \log\left\{\frac{p(z)}{q(z)}\right\} dz$$
$$= -\int q(z) \log p(z) dz - \underbrace{\left(-\int q(z) \log q(z) dz\right)}_{\text{Figure 6}}$$

Entropy of q

Properties:

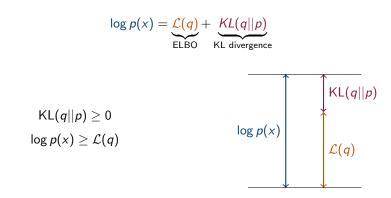
- $\mathsf{KL}(q||p) \ge 0$
- KL(q||p) = 0 iff p = q
- $KL(q||p) \neq KL(p||q)$

The evidence lower bound (ELBO)

Recall

$$p(z|x) = \frac{p(x,z)}{p(x)}$$
$$\log p(x,z) = \log \left[p(z|x) p(x) \right]$$
$$\int \log \frac{p(x,z)}{q(z)} q(z) dz = \int \log \frac{p(x)p(x|z)}{q(z)} q(z) dz$$
$$\int \log \frac{p(x,z)}{q(z)} q(z) dz = \log p(x) - \left[-\int \log \frac{p(z|x)}{q(z)} q(z) dz \right]$$
$$\underbrace{\mathcal{L}(q)}_{\text{ELBO}} = \log p(x) - \underbrace{\mathcal{KL}(q||p)}_{\text{KL divergence}}$$

The evidence lower bound (ELBO)



KL is intractable, so we optimise the ELBO instead

The evidence lower bound (ELBO)

$$\mathcal{L}(q) = \int \log p(x, z)q(z)dz - \int \log q(z)q(z)dz$$
$$= \underbrace{\mathbb{E}_q[\log p(x, z)]}_{\substack{(1)\\ \text{Exp} [\log \text{ prior } + \\ \log \text{ likelihood}]}} + \underbrace{\mathbb{E}_q[-\log q(z)]}_{\text{Entropy of q}}$$

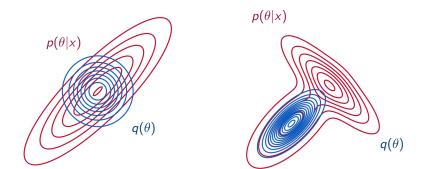
The ELBO trades off two terms:

(1) Prefers q to place its mass on the maximum a posteriori estimate(2) Encourages q to be diffuse

! The ELBO is not convex

Some properties

Bishop (2006)



 $q(\theta)$ tends to be 0 where $p(\theta|x)$ is 0.

VI may lead to a local minimum.

Mean-field approximation

We need to specify the form of q(z). The mean-field family is fully factorised:

$$q(z) = \prod_{i=1}^M q_i(z_i)$$

Optimise the ELBO. Traditionally, VI uses coordinate ascent:

$$\log q_i^*(z_i) \propto \mathsf{I\!E}_{j \neq i}[\log p(x,z)]$$

Iteratively update each parameter, holding others fixed.

Coordinate ascent (CAVI) algorithm

Input : A model $p(X, \theta)$, a dataset X **Output** : A variational density $q(\theta) = \prod_j q_j(\theta_j)$ **Initialise:** Variational factors $q_j(\theta_j)$ **do**

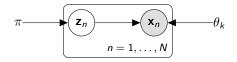
for
$$j \in \{1, ..., J\}$$
 do
 $| \text{ set } q_j(\theta_j) \propto \exp\{\mathbb{E}_{i \neq j}[\log p(X, \theta)]\}$
end

compute the ELBO $\mathcal{L}(q)$ while the ELBO has not converged; return $q(\theta)$.

Main idea

$$p(x) = \sum_{k=1}^{K} \pi_k f_x(x|\theta_k).$$

 f_x parametric density that depends on the parameter(s) θ_k π_k cluster weights



Example: Mixture of Gaussians

$$egin{aligned} & x_n \sim \prod_k \mathcal{N}(\mu_k, \Lambda_k^{-1})^{z_{nk}} \ & z_{nk} \sim \mathsf{Bernoulli}(\pi_k) \end{aligned}$$

Expectation-Maximisation (EM) algorithm

```
Input : A model p(x, z|\theta, \pi), a dataset X
Output : The parameters \theta^*, \pi^* maximising the log-likelihood
Initialise: Parameters \pi, \theta, responsibilities \mathbb{E}[z_{nk}]
do
```

Expectation step: evaluate the responsibilities $\mathbb{E}[z_{nk}]$

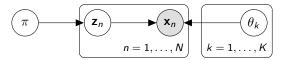
Maximisation step: update the other parameters in turn while *convergence is not reached*;

return θ^* , π^*

... in the Bayesian framework

$$p(x) = \sum_{k=1}^{K} \pi_k f_x(x|\theta_k).$$

 f_x parametric density that depends on the parameter(s) θ_k π_k cluster weights



Example: Mixture of Gaussians

$$\begin{aligned} \pi &\sim \mathsf{Dirichlet}(\alpha_0, \dots, \alpha_0) \\ \theta &= \{\mu, \Sigma\} \\ \mu_k &\sim \mathcal{N}(m_0, (\beta_0 \Lambda_k)^{-1}) \\ \Lambda_k &\sim \mathcal{W}(W_0, \nu_0) \end{aligned}$$

Approximate the true posterior with a variational distribution q

$$q(z, \theta, \pi) = q(z)q(\theta, \pi)$$

EM-type algorithm

Input : A model $p(x, z, \pi, \theta)$, a dataset X **Output** : A variational density $q(z^*, \pi^*, \theta^*) = q(z^*)q(\pi^*, \theta^*)$ **Initialise:** Parameters π, θ , responsibilities $\mathbb{E}[z_{nk}]$ **do**

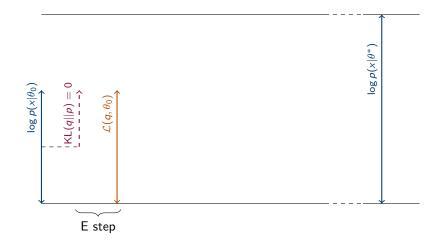
Expectation step: evaluate the responsibilities $\mathbb{E}[z_{nk}]$

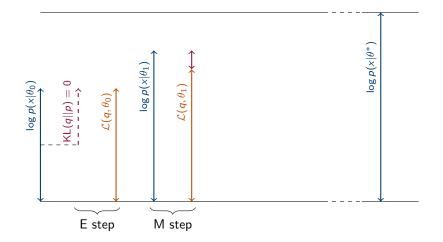
Maximisation step: update the other hyperparameters in turn while the ELBO has not converged;

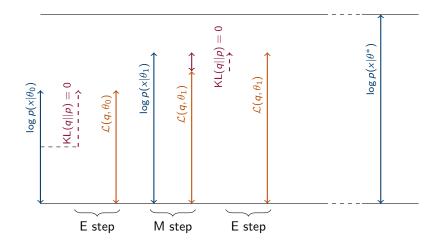
return $q(z^*, \pi^*, \theta^*)$

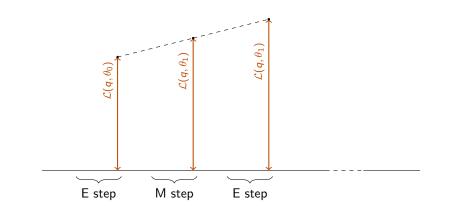
Bishop (2006)









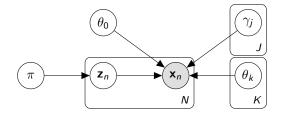


Lower bound used to check:

- Correctness of update equations
- Convergence

Feature selection

$$p(x) = \sum_{k} \pi_{k} \prod_{j} p_{x_{j}}(x_{j}|\theta_{k})^{\gamma_{j}} p_{x_{j}}(x_{j}|\theta_{0})^{1-\gamma_{j}}$$
$$\gamma_{j} \sim \text{Bernoulli}(\delta_{j})$$



Our project

 \sim

Implementation and analysis of the following mixtures using VI:

	Basic model	Feature selection	Model selection
Gaussian	\checkmark	~	\sim
Categorical			
Gaussian + Categorical			

already studied in the literature, code available online
(Bishop 2006, Ahlmann-Eltze and Yau 2018)

already studied in the literature, code *not* available online (Constantinopoulos et al. 2006)

Our project

Current status

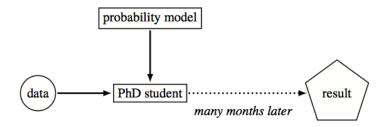


Figure 1: The "how hard could it be?"TM way of probabilistic modeling.

Kucukelbir (2015)

Our project

Current status

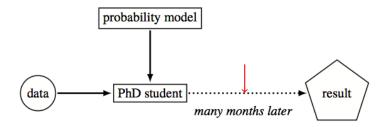


Figure 1: The "how hard could it be?"TM way of probabilistic modeling.

Kucukelbir (2015)



Future work

- Complete R package "vimix" https://acabassi.github.io/vimix/
- Apply to CVD risk data
- Explore automated tools?
 - E.g. TensorFlow Probability, PyMC3, Edward, Stan

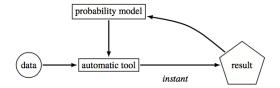


Figure 3: The probabilistic programming way of probabilistic modeling.

Thanks for listening!







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Slide 2

Mikael Häggström and A. Rad Image:Hematopoiesis (human) diagram.png by A. Rad, CC BY-SA 3.0, https://commons.wikimedia.org/w/index.php?curid=7351905