

Stochastic Optimal Transport with applications to flow cytometry data

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Flow cytometry data

Quantifying cellular markers in a biological sample (e.g. blood draw) cell-by-cell.

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- The biological markers are stained.
- The light emitted by a marker indicates whether the marker is present or not on the cell.

Modelling

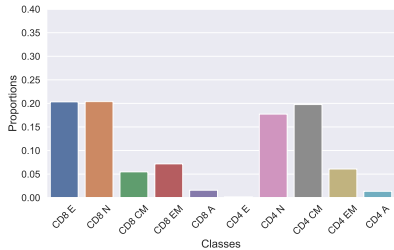
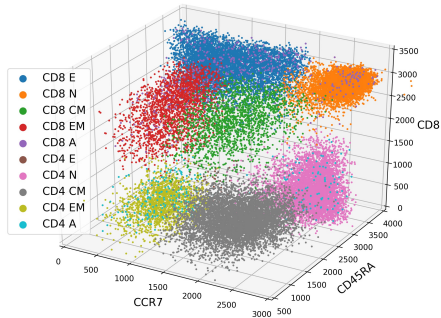
- The observation $X_i \in \mathbb{R}^d$ corresponds to the measures on the i^{th} cell.
- For $m \in \{1, \dots, d\}$, the coefficient $X_i^{(m)}$ corresponds to the light emitted by the biological marker m .
- In a data set X_1, \dots, X_I , the number of observation range from 10 000 to 200 000.

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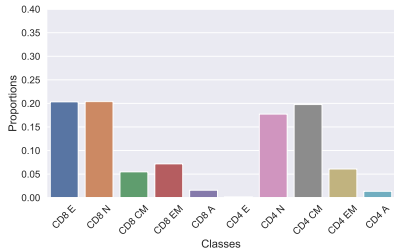
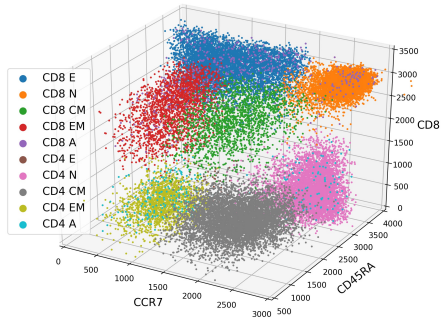
CCR7	CD4	CD45RA	CD3	HLADR	CD38	CD8
717.3	1146.5	3094.8	2526.3	1333.1	1510.2	3203.7

Table 1: Cytometry measurement for one cell. $d = 7$.



Objective

Quantify relative abundance of cell types within the sample.



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Quantify relative abundance of cell types within the sample.

Application

clinical practice: Monitor human disease and response to therapy.

Data analysis

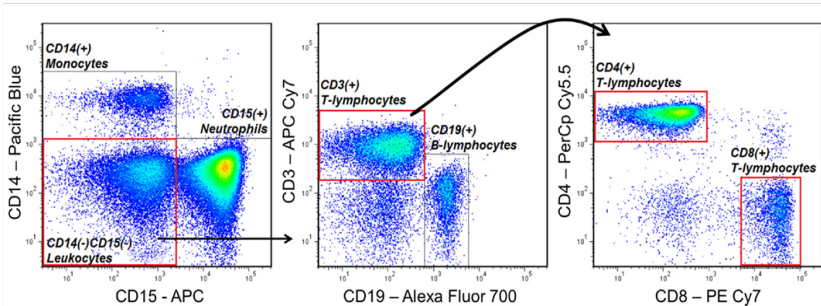


Figure 2: Front. Immunol., 27 July 2015.

Manual gating

Drawbacks: Time consuming, expensive and poorly reproducible.

Unsupervised methods

- Kmeans
- Hierarchical clustering
- Mixture models

Automated methods

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Supervised methods

- Deep learning
- Quadratic discriminant analysis
- Random Forest

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Unsupervised methods

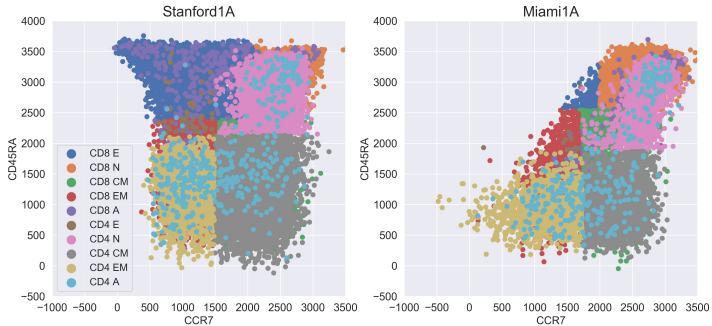
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Manual gating is still the benchmark methods.

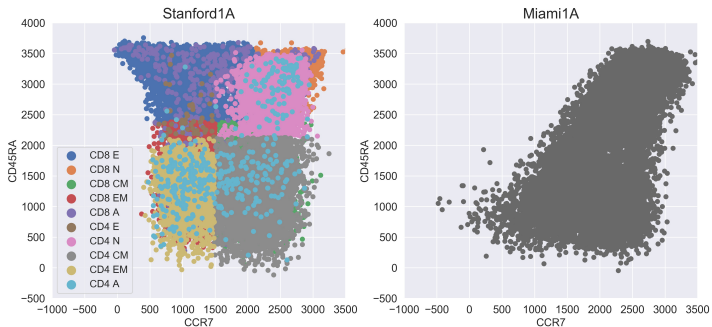
Challenges of the automated analysis



Technical variability

Two samples analysed from the same patient. Cytometry measurements were performed in two different laboratories.

Challenges of the automated analysis



- One gated data set.
- Derive the class proportions in an unclassified data set.

Wasserstein Distance

Let α and β two probability measures on \mathbb{R}^d with finite second moment.

Let $\Pi(\alpha, \beta)$ be the set of probability measures on $\mathbb{R}^d \times \mathbb{R}^d$ with marginals α and β .

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Definition

The Wasserstein distance between α and β is defined as

$$W_2^2(\alpha, \beta) = \min_{\pi \in \Pi(\alpha, \beta)} \int_{\mathbb{R}^d \times \mathbb{R}^d} c(x, y) d\pi(x, y). \quad (1)$$

where $c(x, y) = \|x - y\|_2^2$.

Wasserstein Distance

Discrete Setting

If $\alpha = \sum_i^I a_i \delta_{x_i}$ and $\beta = \sum_{j=1}^J b_j \delta_{y_j}$ are two discrete probability distributions on \mathbb{R}^d the Wasserstein distance reads:

$$W_2^2(\alpha, \beta) = \min_{P \in U(a,b)} \sum_{i,j} C_{i,j} P_{i,j} \quad (2)$$

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Computational Cost

Suppose α and β are two measures with equal size N .

- Requires to store a $N \times N$ matrix.
- Linear programming problem.
- $O(N^3 \log(N))$ operations required.

Entropic regularization (M.Cuturi 2013)

Regularized Wasserstein Distance

For α and β two probability measures the regularized Wasserstein distance is defined as:

$$W^\varepsilon(\alpha, \beta) = \min_{\pi \in \Pi(\alpha, \beta)} \int_{\mathbb{R}^d \times \mathbb{R}^d} c(x, y) d\pi(x, y) + \varepsilon H(\pi). \quad (3)$$

where $\varepsilon > 0$ and

$$H(\pi) = \int_{\mathbb{R}^d \times \mathbb{R}^d} \log \left(\left(\frac{d\pi}{d\alpha \otimes \beta}(x, y) \right) - 1 \right) d\pi(x, y)$$

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Dual problem

$$W^\varepsilon(\alpha, \beta) = \sup_{u, v \in \mathcal{C}(\mathbb{R}^d)} \int u(x) d\alpha(x) + \int v(y) d\beta(y) - \varepsilon \int e^{\frac{u(x) + v(y) - c(x, y)}{\varepsilon}} d\alpha \otimes \beta(x, y) \quad (4)$$

Dual problem in a discrete setting

$$W^\varepsilon(\alpha, \beta) = \max_{u \in \mathbb{R}^I, v \in \mathbb{R}^J} \langle u, a \rangle + \langle v, b \rangle - \varepsilon \langle e^{\frac{u \oplus v - C}{\varepsilon}}, a \otimes b \rangle. \quad (5)$$

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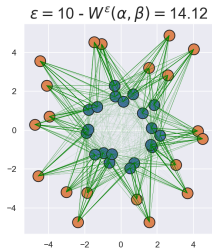
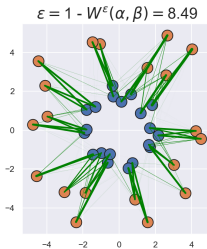
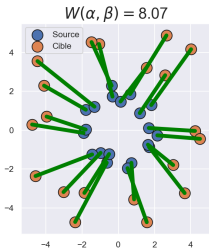
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Block coordinate ascent strategy \rightarrow Sinkhorn algorithm.

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Block coordinate ascent strategy \rightarrow Sinkhorn algorithm. In the case where $I = J = N$, computation of a solution in $O(N^2 \log(N))$ operations.



From Dual to Semi-Dual

Maximizing over a single potential $u \in \mathbb{R}^I$

For $u \in \mathbb{R}^I$, the unique maximizer of the function

$$F(v) = \langle u, a \rangle + \langle v, b \rangle - \varepsilon \langle e^{\frac{u \oplus v - c}{\varepsilon}}, a \otimes b \rangle$$

is the smoothed c -transform of u :

$$u_{c,\varepsilon}(y_j) = -\varepsilon \log \left(\sum_{i=1}^I e^{\frac{u(x_i) - c(x_i, y_j)}{\varepsilon}} a_i \right)$$

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$$\text{And } e^{\frac{u_{c,\varepsilon}(y_j)}{\varepsilon}} = \left(\langle e^{\frac{u - C(\cdot, y_j)}{\varepsilon}}, a \rangle \right)^{-1}$$

Thus

Semi Dual Formulation

$$W^\varepsilon(\alpha, \beta) = \max_{u \in \mathbb{R}^I} \langle u, a \rangle + \langle u_{c, \varepsilon}, b \rangle - \varepsilon$$

In the case where $\alpha = \sum_{i=1}^I a_i \delta_{x_i}$ and $\beta = \sum_{j=1}^J b_j \delta_{y_j}$ are discrete.

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$$W^\varepsilon(\alpha, \beta) = \max_{u \in \mathbb{R}^I} \sum_{i=1}^I u_i a_i + \int_{\mathbb{R}^d} u_{c,\varepsilon} d\beta - \varepsilon$$

And

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→ Expectation formulation of the regularized Wasserstein distance.

Stochastic optimal transport

Let β be any probability measure and $\alpha = \sum_{i=1}^I a_i \delta_{x_i}$.

Proposition (Genevay, Cuturi, Peyré and Bach (2016))

let $\varepsilon \geq 0$,

$$W_\varepsilon(\alpha, \beta) = \max_{u \in \mathbb{R}^I} \mathbb{E}[g_\varepsilon(Y, u)] \quad (6)$$

- Y is a random variable with distribution β .
- $g_\varepsilon(y, u) = \sum_{i=1}^I u_i a_i + u_{c,\varepsilon}(y) - \varepsilon$ is easy to compute for all $y \in \mathbb{R}^d$, and all $u \in \mathbb{R}^I$

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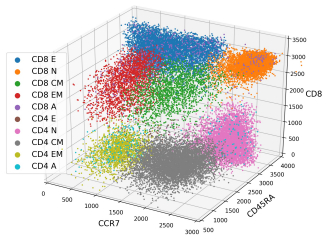
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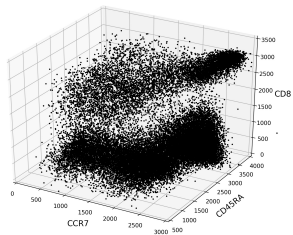
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- Stochastic optimization techniques can be applied.
- No need to store the full cost matrix.

Application to flow cytometry data



(a) Stanford Patient 1



(b) Stanford Patient 3

Framework

- The source distribution α is a mixture model.
- The target distribution β is a mixture model.

Domain adaptation (R.Flammary et al. (2019))

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Re-weight the source distribution in order to reduce the Wasserstein distance $W(\alpha, \beta)$ between the source and target distribution.

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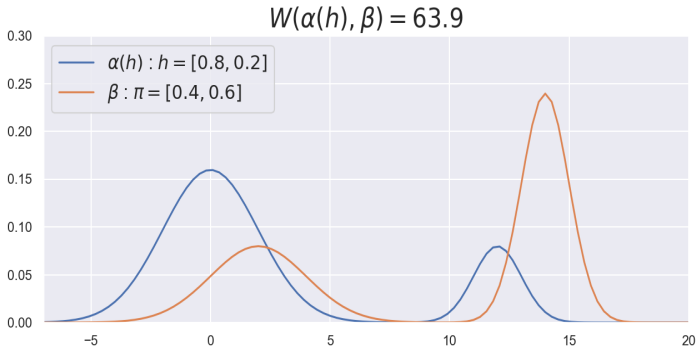
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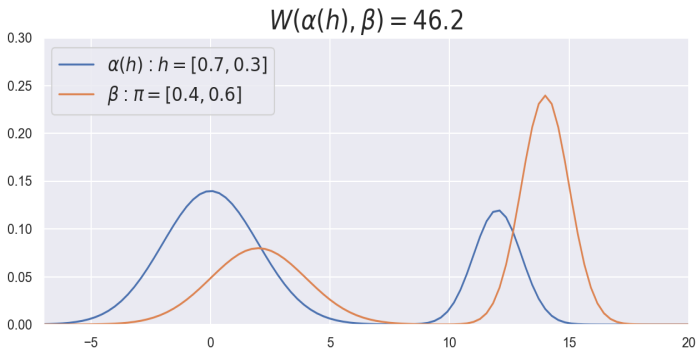
Goal

Estimation of the weights of the mixture π in the target distribution β .

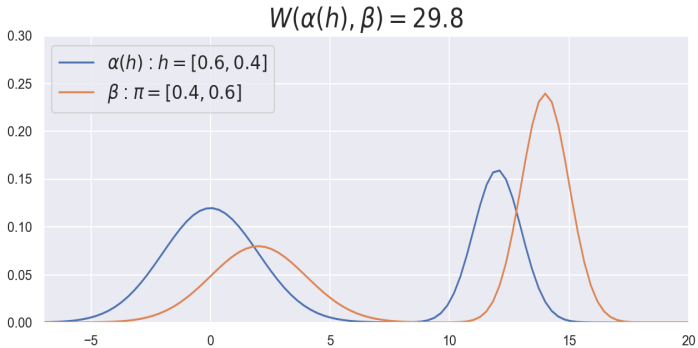
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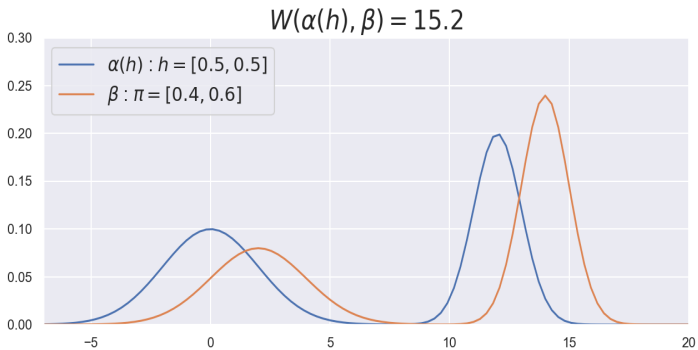
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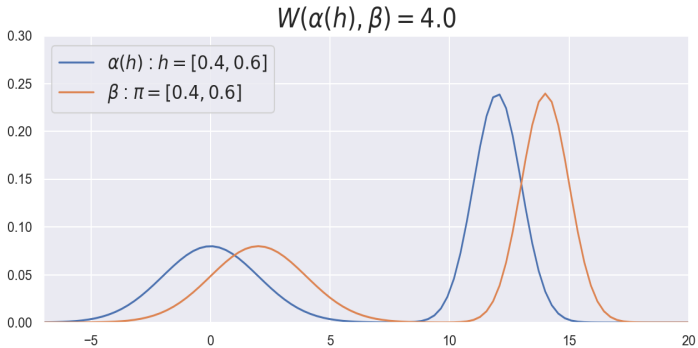
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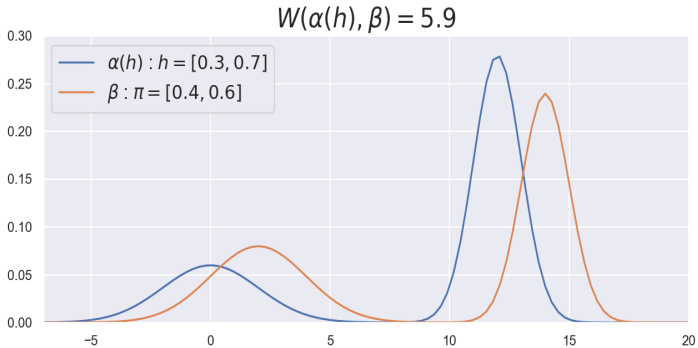
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Re-weighting of the source data

For $h = (h_1, \dots, h_K) \in \Sigma_K$, the measure α re weighted by h is:

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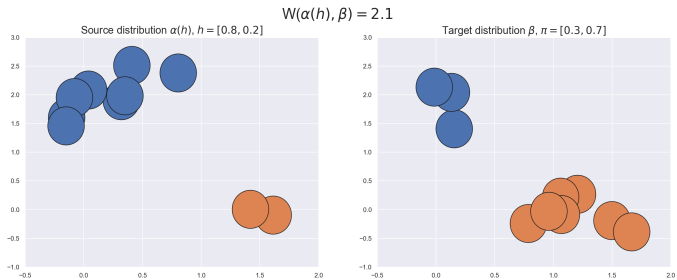
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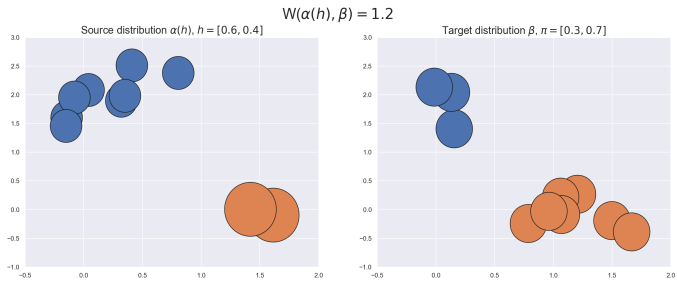
Estimation of the class proportions in the target data set:

$$\hat{\pi} \in \arg \min_{h \in \Sigma_K} W^\varepsilon(\alpha(h), \beta) \quad (8)$$

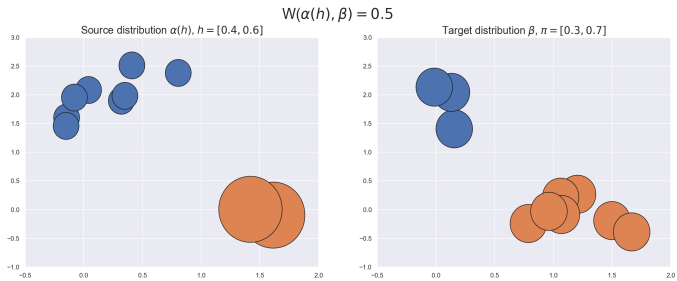
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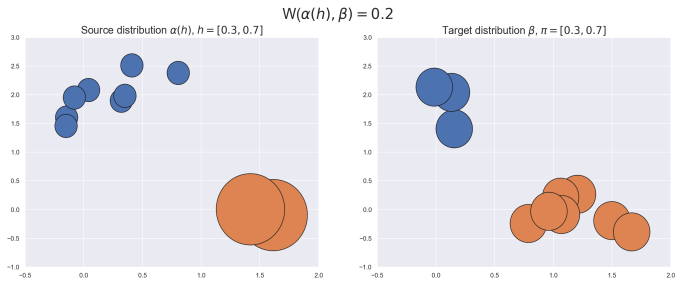
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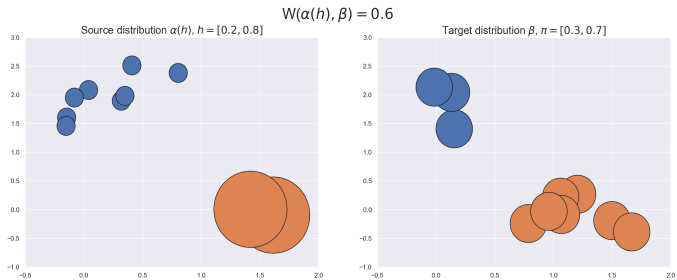
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Descent Ascent procedure

$$\min_{h \in \Sigma_K} W^\varepsilon(\alpha(h), \beta)$$

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New minimization problem:

$$\min_{z \in \mathbb{R}^K} W^\varepsilon(\alpha(\sigma(z)), \beta) = \min_{z \in \mathbb{R}^K} \max_{u \in \mathbb{R}^I} \mathbb{E}[g_\varepsilon(Y, u, \sigma(z))] \quad (10)$$

Objective function:

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for $z \in \mathbb{R}^K$, we need to solve $\max_{u \in \mathbb{R}^I} \mathbb{E}[g_\varepsilon(Y, u, \sigma(z))]$.

→ Stochastic optimal transport computation.

Approximation of $\nabla_z F(z)$

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Double loop algorithm.

Addition of a regularizing term on h (M.Ballu et al. (2020))

Regularization:
$$\varphi(h) = \sum_{k=1}^K h_k \log(h_k). \quad (11)$$

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for $u \in \mathbb{R}^I$, we can compute an explicit solution $h(u) \in \Sigma_K$ of the problem $\min_{h \in \Sigma_K} \mathbb{E}[g_\varepsilon(Y, u, h)] + \lambda \varphi(h)$.

$$k \in \{1, \dots, K\}, \quad (h(u))_k = \frac{\exp\left(-\frac{(\Gamma^T u)_k}{\lambda}\right)}{\sum_{l=1}^K \exp\left(-\frac{(\Gamma^T u)_l}{\lambda}\right)}$$

Using the expression of $h(u)$, problem (12) boils down to

$$\max_{u \in \mathbb{R}^I} \mathbb{E}_{Y \sim \beta} [f_{\varepsilon, \lambda}(Y, u)] \quad (13)$$

where Y is a random variable with distribution β .

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Single loop algorithm.

Simulation study

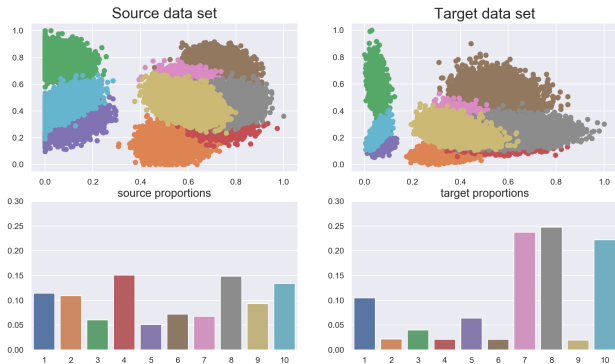


Figure 4: 2D projection of simulated data where $X_i \in \mathbb{R}^{10}$.

Spatial shift between the two data sets

Simulation study

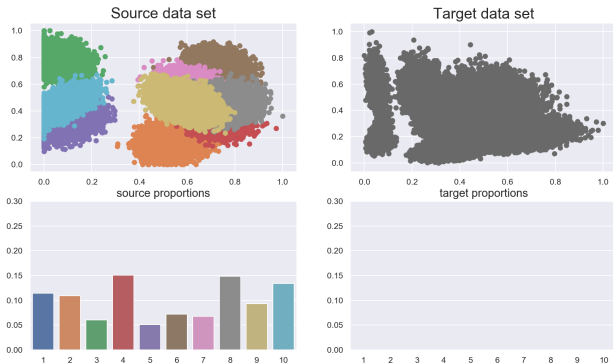
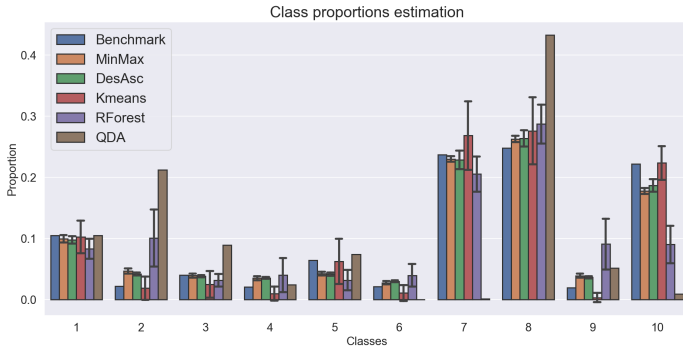


Figure 5: 2D projection of simulated data where $X_i \in \mathbb{R}^{10}$.

- Unsupervised method : Kmeans.
- Supervised methods : QDA and Random Forest.



Results on the flow cytometry data

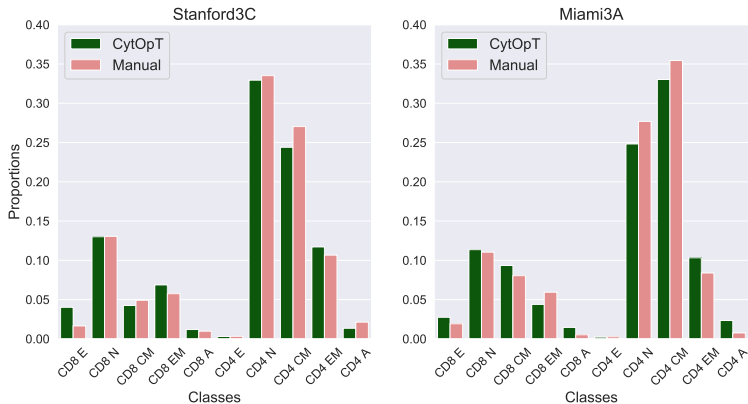


Figure 6: Comparison of the estimated proportions $\hat{\pi}$ by CytOpT with the manual gating benchmark π . Source data set: Stanford1A.

Results on the flow cytometry data

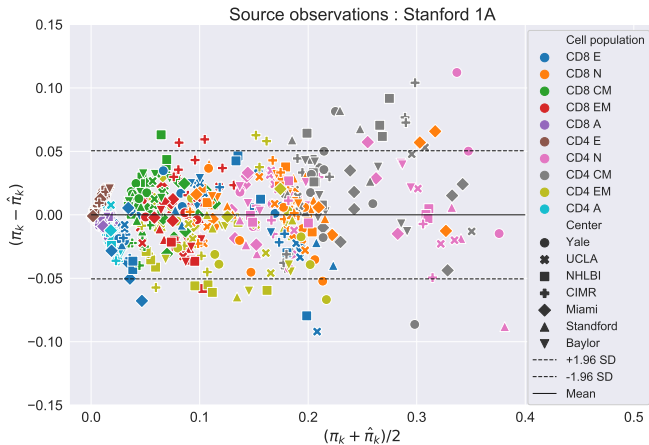


Figure 7: Comparison of the proportions $\hat{\pi}$ estimated with Cyt0pT and the manual benchmark π on the HIPC database.

$\alpha \in \mathcal{M}_+^1(\mathbb{R}^d)$ a probability measure that can be decomposed as a mixture of K probability measure $\alpha_1, \dots, \alpha_K$:

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Let $\beta \in \mathcal{M}_+^1(\mathbb{R}^d)$ an other probability measure. we define

$$\theta_* \in \arg \min_{\theta \in \Sigma_K} W(\alpha_\theta, \beta)$$

Empirical versions of α and β

- $\hat{\alpha} = \frac{1}{T} \sum_{i=1}^I \delta_{X_i} = \sum_{k=1}^K \frac{n_k}{T} \sum_{X_i \in C_k} \delta_{X_i} = \sum_{k=1}^K \frac{n_k}{T} \hat{\alpha}_k.$
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Question 1: How to choose ε, λ in order to minimize the quantity $\mathbb{E} [\|\theta_{\varepsilon, \lambda} - \theta_*\|^2]$?

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Question 2: Study of the convergence of the optimization algorithms to solve (14).



B.Bercu and J.Bigot (2020)

Asymptotic distribution and convergence rates of stochastic algorithms for entropic optimal transportation between probability measures

Annals of Statistics



I.Redko, N.Courty, R.Flamary and D.Tuia (2019)

Optimal Transport for Multi-source Domain Adaptation under Target Shift

International Conference on Artificial Intelligence and Statistics



G. Peyré et M.Cuturi (2018)

Computational Optimal Transport

$$f_{\varepsilon, \lambda}(y_j, u) = \varepsilon \left(\log(b_j) - \log \left(\sum_{i=1}^I \exp \left(\frac{u_i - c(x_i, y_j)}{\varepsilon} \right) \right) \right) - \lambda \log \left(\sum_{l=1}^K \exp \left(-\frac{(\Gamma^T u)_l}{\lambda} \right) \right) - \varepsilon \quad (15)$$