Purity calculation

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1 Background

1.1 Kullback-Leibler divergence

In statistics, the Kullback-Leibler (KL) divergence is a measure of how one probability distribution F_1 is different from a second reference probability distribution F_2 . We may apply the KL divergence idea in the clustering problem, since the larger the KL divergence between distributions, the more pure the groups/clusters are.

For distributions F_1 and F_2 of a continuous random variable, the KL divergence is defined as:

$$D_{KL}(F_1||F_2) = \int_{-\infty}^{+\infty} f_1(x) \log(\frac{f_1(x)}{f_2(x)}) dx$$
 (1)

where f_1 and f_2 denote the probability density of F_1 and F_2 .

Besides, $log(x) \le x - 1$ is always true, then

$$\int -\log(\frac{f_2(x)}{f_1(x)})f_1(x)dx \ge \int -(\frac{f_2(x)}{f_1(x)} - 1)f_1(x)dx$$
$$= \int [f_2(x) - f_1(x)]dx = 0$$

The $D_{KL}(F_1||F_2)$ is always bigger or equal to than 0. Similarly, the $D_{KL}(F_2||F_1)$ is also always bigger or equal to than 0.

1.2 Application

In our setting, we assume the outcomes are from a linear mixed model:

$$Y = S(\beta + b + \Gamma(\alpha'x)) + \epsilon. \tag{2}$$

where,

- S is the matrix of times (intercept, linear, and quadratic term)
- $oldsymbol{\circ}$ is the vector of covariates for fixed effects of $oldsymbol{S}$
- \bullet **b** is the vector of random effects

- Γ is the vector of fixed effects of the baseline covariates.
- $\alpha'x$ is the combination of the input baseline covariates.
- α has the restriction that $||\alpha|| = 1$

Define the covariate matrix of S as z. The z contains both fixed effects and random effects.

$$z = \beta + b + \Gamma w$$

That is, we have distributions for the mixed-effect model coefficients z given $w = \alpha' x$, where

$$z|w \sim N(\beta_j + \Gamma_j w, D_j),$$

for treatment j = 1, 2. Besides, we assume the baseline biosignature x follows distribution with mean μ_x and covariance matrix Σ_x

Based on the Kullback-Leibler divergence, we define the *purity* of the data, which represents how much the differences between the treatment group distribution $f_1(x)$ and the placebo group distribution $f_2(x)$. We define the **purity function** regard to a subject with baseline biosignature \boldsymbol{x} (i.g. the **purity function**given α and the baseline biosignature \boldsymbol{x}) as:

$$g(\boldsymbol{\alpha}'\boldsymbol{x}) = D_{KL}(F_1||F_2) + D_{KL}(F_2||F_1)$$

$$= \int log(f_1(\boldsymbol{z}|\boldsymbol{\alpha}'\boldsymbol{x}))f_1(\boldsymbol{z}|\boldsymbol{\alpha}'\boldsymbol{x})dz - \int log(f_2(\boldsymbol{z}|\boldsymbol{\alpha}'\boldsymbol{x}))f_1(\boldsymbol{z}|\boldsymbol{\alpha}'\boldsymbol{x})dz$$

$$+ \int log(f_2(\boldsymbol{z}|\boldsymbol{\alpha}'\boldsymbol{x}))f_2(\boldsymbol{z}|\boldsymbol{\alpha}'\boldsymbol{x})dz - \int log(f_1(\boldsymbol{z}|\boldsymbol{\alpha}'\boldsymbol{x}))f_2(\boldsymbol{z}|\boldsymbol{\alpha}'\boldsymbol{x})dz$$
(3)

where,

$$f_1(\boldsymbol{z}|\boldsymbol{w}) = \frac{1}{\sqrt{((2\pi)^p |\boldsymbol{D}_1|)}} exp(-\frac{1}{2}(\boldsymbol{z} - \boldsymbol{\mu}_1)' \boldsymbol{D}_1^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_1))$$

$$f_2(\boldsymbol{z}|\boldsymbol{w}) = \frac{1}{\sqrt{((2\pi)^p |\boldsymbol{D}_2|)}} exp(-\frac{1}{2}(\boldsymbol{z} - \boldsymbol{\mu}_2)' \boldsymbol{D}_2^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_2))$$

$$\boldsymbol{\mu}_1 = \boldsymbol{\beta}_1 + \boldsymbol{\Gamma}_1 w, \boldsymbol{\mu}_2 = \boldsymbol{\beta}_2 + \boldsymbol{\Gamma}_2 w$$

Furthermore, we define f_w as the distribution of the combination of baseline signature, $w = \alpha' x$.

Then the purity function regards to the whole data set is defined as:

purity(
$$\boldsymbol{\alpha}$$
) = $\int g(\boldsymbol{\alpha}'\boldsymbol{x}) f_w(\boldsymbol{\alpha}'\boldsymbol{x}) d\boldsymbol{\alpha}'\boldsymbol{x}$
= $E(g(\boldsymbol{\alpha}'\boldsymbol{x}))$ (4)

Therefore, we may estimate the dataset's purity given a vector α by the mean value of g() function,

$$\hat{g}(\boldsymbol{\alpha}) = \bar{g}(\boldsymbol{\alpha}'\boldsymbol{x})$$

1.2.1 Purity Calculation

We can separate Equation(3) into four parts: $\int f_1 log f_1$, $\int f_2 log f_2$, $\int f_1 log f_2$, and $\int f_2 log f_1$.

• For $\int f_1 log f_1$ and $\int f_2 log f_2$:

$$\int f_1 log f_1 = E_1(log(f_1))$$

$$= E_1(-\frac{p}{2}log(2\pi) - \frac{1}{2}log(|\mathbf{D}_1|) - \frac{1}{2}(\mathbf{z} - \boldsymbol{\mu}_1)'\mathbf{D}_1^{-1}(\mathbf{z} - \boldsymbol{\mu}_1))$$

$$= -\frac{p}{2}log(2\pi) - \frac{1}{2}log(|\mathbf{D}_1|) - \frac{1}{2}E_1[(\mathbf{z} - \boldsymbol{\mu}_1)'\mathbf{D}_1^{-1}(\mathbf{z} - \boldsymbol{\mu}_1)]$$

And

$$E_{1}[(\boldsymbol{z} - \boldsymbol{\mu}_{1})'\boldsymbol{D}_{1}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{1})] = E_{1}[tr((\boldsymbol{z} - \boldsymbol{\mu}_{1})'\boldsymbol{D}_{1}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{1}))]$$

$$= E_{1}[tr(\boldsymbol{D}_{1}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{1})'(\boldsymbol{z} - \boldsymbol{\mu}_{1}))]$$

$$= tr(E_{1}[\boldsymbol{D}_{1}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{1})'(\boldsymbol{z} - \boldsymbol{\mu}_{1})])$$

$$= tr(\boldsymbol{D}_{1}^{-1}E_{1}[(\boldsymbol{z} - \boldsymbol{\mu}_{1})'(\boldsymbol{z} - \boldsymbol{\mu}_{1})])$$

$$= tr(\boldsymbol{D}_{1}^{-1}\boldsymbol{D}_{1}) = tr(\boldsymbol{I}_{p}) = p$$

Therefore,

$$\int f_1 log f_1 = -\frac{p}{2} log(2\pi) - \frac{1}{2} log(|\mathbf{D}_1|) - \frac{p}{2}$$
 (5)

Similarly,

$$\int f_2 log f_2 = -\frac{p}{2} log(2\pi) - \frac{1}{2} log(|\mathbf{D}_2|) - \frac{p}{2}$$
 (6)

• For $\int f_1 log f_2$ and $\int f_2 log f_1$

$$\begin{split} \int f_1 log f_2 = & E_1(log f_2) \\ = & E_1(-\frac{p}{2}log(2\pi) - \frac{1}{2}log(|\boldsymbol{D}_2|) - \frac{1}{2}(\boldsymbol{z} - \boldsymbol{\mu}_2)'\boldsymbol{D}_2^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_2)) \\ = & - \frac{p}{2}log(2\pi) - \frac{1}{2}log(|\boldsymbol{D}_2|) - \frac{1}{2}E_1[(\boldsymbol{z} - \boldsymbol{\mu}_2)'\boldsymbol{D}_2^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_2)] \end{split}$$

And

$$E_{1}[(\boldsymbol{z} - \boldsymbol{\mu}_{2})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{2})] = E_{1}[(\boldsymbol{z} - \boldsymbol{\mu}_{1} + \boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{1} + \boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})]$$

$$= E_{1}[(\boldsymbol{z} - \boldsymbol{\mu}_{1})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{1}) + (\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{1})$$

$$+ (\boldsymbol{z} - \boldsymbol{\mu}_{1})\boldsymbol{D}_{2}^{-1}(\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2}) + (\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})]$$

$$= E_{1}[(\boldsymbol{z} - \boldsymbol{\mu}_{1})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{1})] + (\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})$$

$$= E_{1}[(\boldsymbol{z} - \boldsymbol{\mu}_{1})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{1})] + 0 + 0 + (\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})$$

$$= E_{1}[tr(\boldsymbol{z} - \boldsymbol{\mu}_{1})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{1})] + (\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})$$

$$= E_{1}[tr(\boldsymbol{D}_{2}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{1})'(\boldsymbol{z} - \boldsymbol{\mu}_{1}))] + (\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})$$

$$= tr(E_{1}[\boldsymbol{D}_{2}^{-1}(\boldsymbol{z} - \boldsymbol{\mu}_{1})'(\boldsymbol{z} - \boldsymbol{\mu}_{1})]) + (\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})$$

$$= tr(\boldsymbol{D}_{2}^{-1}E_{1}[(\boldsymbol{z} - \boldsymbol{\mu}_{1})'(\boldsymbol{z} - \boldsymbol{\mu}_{1})]) + (\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})$$

$$= tr(\boldsymbol{D}_{2}^{-1}D_{1}) + (\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})'\boldsymbol{D}_{2}^{-1}(\boldsymbol{\mu}_{1} - \boldsymbol{\mu}_{2})$$

Therefore,

$$\int f_1 log f_2 = -\frac{p}{2} log(2\pi) - \frac{1}{2} log(|\mathbf{D}_2|) - \frac{1}{2} \left(tr(\mathbf{D}_2^{-1} \mathbf{D}_1) + (\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2)' \mathbf{D}_2^{-1} (\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2) \right)$$
(7)

Similarly,

$$\int f_2 log f_1 = -\frac{p}{2} log(2\pi) - \frac{1}{2} log(|\mathbf{D}_1|) - \frac{1}{2} \left(tr(\mathbf{D}_1^{-1} \mathbf{D}_2) + (\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2)' \mathbf{D}_1^{-1} (\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2) \right)$$
(8)

Therefore, the equation (1) is:

$$(3) = (5) - (7) + (6) - (8)$$

That is,

$$\int log(f_1)f_1 - \int log(f_2)f_1 + \int log(f_2)f_2 - \int log(f_1)f_2
= \left(-\frac{p}{2}log(2\pi) - \frac{1}{2}log(|\mathbf{D}_1|) - \frac{p}{2}\right)
- \left(-\frac{p}{2}log(2\pi) - \frac{1}{2}log(|\mathbf{D}_2|) - \frac{1}{2}\left(tr(\mathbf{D}_2^{-1}\mathbf{D}_1) + (\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2)'\mathbf{D}_2^{-1}(\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2)\right)\right)
+ \left(-\frac{p}{2}log(2\pi) - \frac{1}{2}log(|\mathbf{D}_2|) - \frac{p}{2}\right)
- \left(-\frac{p}{2}log(2\pi) - \frac{1}{2}log(|\mathbf{D}_1|) - \frac{1}{2}\left(tr(\mathbf{D}_1^{-1}\mathbf{D}_2) + (\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2)'\mathbf{D}_1^{-1}(\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2)\right)\right)
= -p + \frac{1}{2}tr(\mathbf{D}_2^{-1}\mathbf{D}_1) + \frac{1}{2}tr(\mathbf{D}_1^{-1}\mathbf{D}_2) + \frac{1}{2}(\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2)'(\mathbf{D}_1^{-1} + \mathbf{D}_2^{-1})(\boldsymbol{\mu}_1 - \boldsymbol{\mu}_2)$$

where $\mu_1 = \beta_1 + \Gamma_1 \alpha' x$, $\mu_2 = \beta_2 + \Gamma_2 \alpha' x$.

Besides,

$$egin{aligned} &(m{\mu}_1 - m{\mu}_2)'(m{D}_1^{-1} + m{D}_2^{-1})(m{\mu}_1 - m{\mu}_2) \ &= ig(m{eta}_1 - m{eta}_2 + (m{\Gamma}_1 - m{\Gamma}_2)m{lpha'}m{x}ig)'(m{D}_1^{-1} + m{D}_2^{-1})ig(m{eta}_1 - m{eta}_2 + (m{\Gamma}_1 - m{\Gamma}_2)m{lpha'}m{x}ig) \ &= (m{eta}_1 - m{eta}_2)'(m{D}_1^{-1} + m{D}_2^{-1})(m{eta}_1 - m{eta}_2) \ &+ 2ig[(m{eta}_1 - m{eta}_2)'(m{D}_1^{-1} + m{D}_2^{-1})(m{\Gamma}_1 - m{\Gamma}_2)m{x'}m{lpha} \ &+ m{lpha'}m{x}m{x}'m{lpha}ig((m{\Gamma}_1 - m{\Gamma}_2)ig)'(m{D}_1^{-1} + m{D}_2^{-1})ig((m{\Gamma}_1 - m{\Gamma}_2)ig) \end{aligned}$$

Therefore, the purity for a subject with baseline biosignature x is:

$$g(\boldsymbol{\alpha}'\boldsymbol{x}) = -p + \frac{1}{2}tr(\boldsymbol{D}_{2}^{-1}\boldsymbol{D}_{1}) + \frac{1}{2}tr(\boldsymbol{D}_{1}^{-1}\boldsymbol{D}_{2})$$

$$+ \frac{1}{2}\{(\boldsymbol{\beta}_{1} - \boldsymbol{\beta}_{2})'(\boldsymbol{D}_{1}^{-1} + \boldsymbol{D}_{2}^{-1})(\boldsymbol{\beta}_{1} - \boldsymbol{\beta}_{2})$$

$$+ 2[(\boldsymbol{\beta}_{1} - \boldsymbol{\beta}_{2})'(\boldsymbol{D}_{1}^{-1} + \boldsymbol{D}_{2}^{-1})(\boldsymbol{\Gamma}_{1} - \boldsymbol{\Gamma}_{2})\boldsymbol{x}'\boldsymbol{\alpha}$$

$$+ \boldsymbol{\alpha}'\boldsymbol{x}\boldsymbol{x}'\boldsymbol{\alpha}((\boldsymbol{\Gamma}_{1} - \boldsymbol{\Gamma}_{2}))'(\boldsymbol{D}_{1}^{-1} + \boldsymbol{D}_{2}^{-1})((\boldsymbol{\Gamma}_{1} - \boldsymbol{\Gamma}_{2}))\}$$

$$(9)$$

The dataset's purity, which is the expectation of the g() function is:

$$purity(\alpha) = E(g(\alpha'x))$$

$$= -p + \frac{1}{2}tr(\mathbf{D}_{2}^{-1}\mathbf{D}_{1}) + \frac{1}{2}tr(\mathbf{D}_{1}^{-1}\mathbf{D}_{2})$$

$$+ \frac{1}{2}\left\{A_{1} + 2A_{2}E(\mathbf{x}'\alpha) + A_{3}E(\alpha'\mathbf{x}\mathbf{x}'\alpha)\right\}$$

$$= A_{0} + \frac{A_{1}}{2} + A_{2}\boldsymbol{\mu}'_{x}\alpha + \frac{A_{3}}{2}[tr(\alpha'\boldsymbol{\Sigma}_{x}\alpha) + \alpha'\boldsymbol{\mu}_{x}\boldsymbol{\mu}'_{x}\alpha]$$

$$= A_{0} + \frac{A_{1}}{2} + A_{2}\boldsymbol{\mu}'_{x}\alpha + \frac{A_{3}}{2}[\alpha'\boldsymbol{\Sigma}_{x}\alpha + \alpha'\boldsymbol{\mu}_{x}\boldsymbol{\mu}'_{x}\alpha]$$

$$= A_{0} + \frac{A_{1}}{2} + A_{2}\boldsymbol{\mu}'_{x}\alpha + \frac{A_{3}}{2}[\alpha'\boldsymbol{\Sigma}_{x}\alpha + \alpha'\boldsymbol{\mu}_{x}\boldsymbol{\mu}'_{x}\alpha]$$

$$(10)$$

where

•
$$A_0 = -p + \frac{1}{2}tr(\boldsymbol{D}_2^{-1}\boldsymbol{D}_1) + \frac{1}{2}tr(\boldsymbol{D}_1^{-1}\boldsymbol{D}_2)$$

•
$$A_1 = (\boldsymbol{\beta}_1 - \boldsymbol{\beta}_2)'(\boldsymbol{D}_1^{-1} + \boldsymbol{D}_2^{-1})(\boldsymbol{\beta}_1 - \boldsymbol{\beta}_2)$$

•
$$A_2 = (\boldsymbol{\beta}_1 - \boldsymbol{\beta}_2)'(\boldsymbol{D}_1^{-1} + \boldsymbol{D}_2^{-1})(\boldsymbol{\Gamma}_1 - \boldsymbol{\Gamma}_2)$$

•
$$A_3 = (\Gamma_1 - \Gamma_2)'(D_1^{-1} + D_2^{-1})((\Gamma_1 - \Gamma_2))$$

All A_0, A_1, A_2, A_3 are scalars.

Therefore, if the distribution of x, f_1, f_2 and α are known, we can calculate the purity by:

purity(
$$\boldsymbol{\alpha}$$
) = $A_0 + \frac{A_1}{2} + A_2 \boldsymbol{\mu}_x' \boldsymbol{\alpha} + \frac{A_3}{2} [\boldsymbol{\alpha}' \boldsymbol{\Sigma}_x \boldsymbol{\alpha} + \boldsymbol{\alpha}' \boldsymbol{\mu}_x \boldsymbol{\mu}_x' \boldsymbol{\alpha}]$ (11)

If the distribution of x, f_1 , and f_2 are unknown, given an α value, we can estimated the purity by

$$\hat{\text{purity}}(\boldsymbol{\alpha}) = \hat{A}_0 + \frac{\hat{A}_1}{2} + \hat{A}_2 \hat{\boldsymbol{\mu}}_x' \boldsymbol{\alpha} + \frac{\hat{A}_3}{2} [\boldsymbol{\alpha}' \hat{\boldsymbol{\Sigma}}_x \boldsymbol{\alpha} + \boldsymbol{\alpha}' \hat{\boldsymbol{\mu}}_x \hat{\boldsymbol{\mu}}_x' \boldsymbol{\alpha}]$$
(12)

1.2.2 Optimization of α

Above Equation (11) has given us the purity function based on α . As well as the restriction of α that $||\alpha|| = 1$, we can use the method of Lagrange multiplier to find the solution of α to maximize the data purity.

Besides, we also notice that, if standardization of baseline covariates is performed, the

•
$$\mu_x = [0, ..]'_p$$

The estimation of purity in equation can be simplified as is

purity(
$$\boldsymbol{\alpha}$$
) = $A_0 + \frac{A_1}{2} + A_2 \boldsymbol{\mu}_x' \boldsymbol{\alpha} + \frac{A_3}{2} [\boldsymbol{\alpha}' \boldsymbol{\Sigma}_x \boldsymbol{\alpha} + \boldsymbol{\alpha}' \boldsymbol{\mu}_x \boldsymbol{\mu}_x' \boldsymbol{\alpha}]$
= $A_0 + \frac{A_1}{2} + \frac{A_3}{2} \boldsymbol{\alpha}' \boldsymbol{\Sigma}_x \boldsymbol{\alpha}$ (13)

Then the function to be optimized with restriction $||\boldsymbol{\alpha}|| - 1 = 0$ can be defined as $h(\boldsymbol{\alpha}; \lambda)$:

$$h(\boldsymbol{\alpha}; \lambda) = A_0 + \frac{A_1}{2} + \frac{A_3}{2} \alpha' \boldsymbol{\Sigma}_x \boldsymbol{\alpha} + \lambda (\boldsymbol{\alpha}' \boldsymbol{\alpha} - 1)$$

$$= A_0 + \frac{A_1}{2} - \lambda + \boldsymbol{\alpha}' (\frac{A_3}{2} \boldsymbol{\Sigma}_x + \lambda \boldsymbol{I}) \boldsymbol{\alpha}$$
 (14)

When A_0, A_1, A_3 are known (constant), to maximize purity $(\alpha) = \alpha' \Sigma_x \alpha$ subjects to $g(\alpha) = 0$. So by Lagrange multiplier, there is λ so that

$$\nabla \text{purity} = \lambda \nabla q$$

Note $\nabla g(\boldsymbol{\alpha}) = 2\boldsymbol{\alpha}$. On the other hand, $\nabla \text{purity}(\boldsymbol{\alpha}) = 2\boldsymbol{\Sigma}_x\boldsymbol{\alpha}$ as $\boldsymbol{\Sigma}_x$ is symmetric. Thus we have $\boldsymbol{\Sigma}_x\boldsymbol{\alpha} = \lambda\boldsymbol{\alpha}$. Therefore, $\boldsymbol{\alpha}$ is an eigenvector of $\boldsymbol{\Sigma}_x$ and λ is an eigenvalue of $\boldsymbol{\Sigma}_x$. The eigenvector of $\boldsymbol{\Sigma}_x$ with the largest eigenvalue can maximize the purity function.

1.2.3 Algorithm

Given the formulas of data purity and the solution of α , the algorithm to find the α that maximize the purity as well as the max purity can be summerzied as:

1) Set an initial $\boldsymbol{\alpha}^{(0)}$ value. And fit the LME model.

$$Y = S(\beta + b + \Gamma(\alpha'x)) + \epsilon. \tag{15}$$

2) Estimate $\hat{\Sigma}_x$ and get the $\hat{\lambda}^{(1)}$ and $\hat{\alpha}^{(1)}$, which is the largest eigenvalue of $\hat{\Sigma}_x$ and its corresponding eigenvector.

- 3) Estimate $\hat{\beta}_1^{(1)}, \hat{\beta}_2^{(1)}, \hat{\Gamma}_1^{(1)}, \hat{\Gamma}_2^{(1)}, \hat{D}_1^{(1)}, \hat{D}_2^{(1)}$.
- 4) Plug in the above estimated values in equation 13 to get the estimated purity.
- 5) Wrap the 1-4 steps into a function and optimize the function with Newton Raphson method.

1.2.4 others

If we cannot assume $\mu_x = 0$, then

we can use the method of Lagrange multiplier to find the solution of α to maximize the data purity:

$$h(\boldsymbol{\alpha}; \lambda) = A_0 + \frac{A_1}{2} + A_2 \boldsymbol{\mu}_x' \boldsymbol{\alpha} + \frac{A_3}{2} [\boldsymbol{\alpha}' \boldsymbol{\Sigma}_x \boldsymbol{\alpha} + \boldsymbol{\alpha}' \boldsymbol{\mu}_x \boldsymbol{\mu}_x' \boldsymbol{\alpha}] + \lambda (\boldsymbol{\alpha}' \boldsymbol{\alpha} - 1)$$

$$= A_0 + \frac{A_1}{2} - \lambda + A_2 \boldsymbol{\mu}_x' \boldsymbol{\alpha} + \frac{A_3}{2} \boldsymbol{\alpha}' (\boldsymbol{\Sigma}_x + \boldsymbol{\mu}_x \boldsymbol{\mu}_x' + \frac{2\lambda}{A_3} \boldsymbol{I}) \boldsymbol{\alpha}$$
(16)

Based on the facts of matrix derivatives,

$$\bullet$$
 $\frac{\partial AX}{\partial X} = A$

•
$$\frac{\partial X'AX}{\partial X} = X'(A+A')$$

The first derivative of equation (13) is

$$\frac{\partial h(\boldsymbol{\alpha}; \lambda)}{\partial \boldsymbol{\alpha}} = A_2 \boldsymbol{\mu}_x' + \frac{A_3}{2} \boldsymbol{\alpha}' [(\boldsymbol{\Sigma}_x + \boldsymbol{\mu}_x \boldsymbol{\mu}_x' + \frac{2\lambda}{A_3} \boldsymbol{I}) + (\boldsymbol{\Sigma}_x + \boldsymbol{\mu}_x \boldsymbol{\mu}_x' + \frac{2\lambda}{A_3} \boldsymbol{I})']$$

$$= A_2 \boldsymbol{\mu}_x' + A_3 \boldsymbol{\alpha}' (\boldsymbol{\Sigma}_x + \boldsymbol{\mu}_x \boldsymbol{\mu}_x' + \frac{2\lambda}{A_3} \boldsymbol{I})$$
(17)

Set Eq(14) = 0, we have

$$\hat{\boldsymbol{\alpha}} = -\frac{A_2}{A_3} (\boldsymbol{\Sigma}_x + \boldsymbol{\mu}_x \boldsymbol{\mu}_x' + \frac{2\lambda}{A_3} \boldsymbol{I})^{-1} \boldsymbol{\mu}_x$$
(18)

The second derivative of equation (13) is

$$A_3(\mathbf{\Sigma}_x + \boldsymbol{\mu}_x \boldsymbol{\mu}_x' + \frac{2\lambda}{A_3} \boldsymbol{I}) \tag{19}$$

The partial derivative of equation (13) w.r.t λ is:

$$\frac{\partial h(\boldsymbol{\alpha}; \lambda)}{\partial \lambda} = \boldsymbol{\alpha}' \boldsymbol{\alpha} - 1 \tag{20}$$

Set Eq (17) = 0 and plug in the estimated $\hat{\alpha}$ value in, we have

$$\mu'_{x}(\Sigma_{x} + \mu_{x}\mu'_{x} + \lambda_{2}I)^{-1}(\Sigma_{x} + \mu_{x}\mu'_{x} + \lambda_{2}I)^{-1}\mu_{x} - \frac{A_{3}^{2}}{A_{2}^{2}}$$

$$= \mu'_{x}(\Sigma_{x} + \mu_{x}\mu'_{x})^{-1}(\Sigma_{x} + \mu_{x}\mu'_{x})^{-1}\mu_{x} - \frac{A_{3}^{2}}{A_{2}^{2}} + 2\lambda_{2}\mu'_{x}(\Sigma_{x} + \mu_{x}\mu'_{x})^{-1}\mu_{x} + \lambda_{2}^{2}\mu'_{x}\mu_{x}$$

$$= B_{0}\lambda_{2}^{2} + 2B_{1}\lambda_{2} + B_{2}$$

$$= (\sqrt{B_{0}}\lambda_{2})^{2} + 2\frac{B_{1}}{\sqrt{B_{0}}}\sqrt{B_{0}}\lambda_{2} + \frac{B_{1}^{2}}{B_{0}} - \frac{B_{1}^{2}}{B_{0}} + B_{2}$$

$$= (\sqrt{B_{0}}\lambda_{2} + \frac{B_{1}}{\sqrt{B_{0}}})^{2} - (\frac{B_{1}^{2}}{B_{0}} - B_{2})$$

$$= 0$$

$$\rightarrow \lambda_{2} = \frac{1}{\sqrt{B_{0}}}(\sqrt{\frac{B_{1}^{2}}{\sqrt{B_{0}}} - B_{2}} - \frac{B_{1}}{\sqrt{B_{0}}}) = \sqrt{\frac{B_{1}^{2}}{B_{0}} - \frac{B_{2}}{B_{0}}} - \frac{B_{1}}{B_{0}}$$

$$(21)$$

where

•
$$\lambda_2 = \frac{2\lambda}{A_2}$$

•
$$B_0 = \boldsymbol{\mu}_x' \boldsymbol{\mu}_x$$

$$\bullet \ B_1 = \boldsymbol{\mu}_r' (\boldsymbol{\Sigma}_x + \boldsymbol{\mu}_x \boldsymbol{\mu}_r')^{-1} \boldsymbol{\mu}_x$$

•
$$B_2 = \mu_x' (\Sigma_x + \mu_x \mu_x')^{-1} (\Sigma_x + \mu_x \mu_x')^{-1} \mu_x - \frac{A_3^2}{A_2^2}$$

Plug in the λ value, we could get the estimated α :

$$\hat{oldsymbol{lpha}} = -rac{A_2}{A_3} (oldsymbol{\Sigma}_x + oldsymbol{\mu}_x oldsymbol{\mu}_x' + \lambda oldsymbol{I})^{-1} oldsymbol{\mu}_x$$

For example, if $\boldsymbol{X} \sim MVN(\boldsymbol{1}, I_{p_x})$, then

$$\bullet \ B_0 = \boldsymbol{\mu}_x' \boldsymbol{\mu}_x = p_x$$

•
$$B_1 = \mu'_x (\Sigma_x + \mu_x \mu'_x)^{-1} \mu_x = 0.5 p_x$$

•
$$B_2 = \mu_x' (\Sigma_x + \mu_x \mu_x')^{-1} (\Sigma_x + \mu_x \mu_x')^{-1} \mu_x - \frac{A_3^2}{A_2^2} = 0.25 p_x - \frac{A_3^2}{A_2^2}$$

And

$$\lambda = \sqrt{\frac{B_1^2}{B_0} - \frac{B_2}{\sqrt{B_0}}} - \frac{B_1}{B_0} = \sqrt{0.25p_x - 0.25\sqrt{p_x} + \frac{A_3^2}{A_2^2}} - 0.5$$

where p_x is the dimension of \boldsymbol{X}