t_sne

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Introduction

We are going to play with the Iris dataset:

```
X <- iris %>% dplyr::select(-Species) %>% as.matrix()
n <- nrow(X)
p <- ncol(X)
```

Euclidean distance

The way to calculate the Euclidean distance:

$$\|\mathbf{x}_i - \mathbf{x}_j\|^2 = \|\mathbf{x}_i\|^2 + \|\mathbf{x}_j\|^2 - 2\mathbf{x}_i'\mathbf{x}_j$$

where $\|\mathbf{x}_i\| = \sqrt{x_{i,1}^2 + \cdots + x_{i,p}^2}$ and $\mathbf{x}_k^2 = \mathbf{x}_k'\mathbf{x}_k = x_{k1}^2 + ... + x_{kp}^2$.

The following method apply the formula above:

```
x_diff <- function(X) {
  n <- nrow(X)
  sum_x <- apply(X^2, MARGIN=1, FUN=sum)
  sum_x_m <- t(matrix(replicate(n, sum_x), byrow=T, nrow=n))
  cross_times_minus_2 <- -2 * (X %*% t(X))
  D <- t(cross_times_minus_2 + sum_x_m) + sum_x_m
  D <- round(D, digits=4)
}
```

Perplexity

$$Perp_i = 2^{H_i}$$

Where H_i is the Shannon entropy in the point x_i of the conditional probability:

$$
\begin{aligned}
H_i &= -\sum_{j \neq i} p_{j|i} \log(p_{j|i}) \\
&= -\sum_{j \neq i} p_{j|i} \log(\frac{p_{j,i}}{p_i}) \\
&= -\sum_{j \neq i} p_{j|i} (\log(p_{j,i}) - \log(p_i)) \\
&= -\sum_{j \neq i} p_{j|i} (\log(e^{-||x_i-x_j||^2/2\sigma^2}) - \log(\sum_{k \neq i} e^{-||x_i-x_j||^2/2\sigma^2})) \\
&= -\sum_{j \neq i} p_{j|i} ((-||x_i - x_j||^2/2\sigma^2) - \log(\sum_{k \neq i} e^{-||x_i-x_j||^2/2\sigma^2})) \\
&= \sum_{j \neq i} p_{j|i} (\log(S_i) + ||x_i - x_j||^2 \frac{1}{2\sigma^2}) \\
&= \log(S_i) \sum_{j \neq i} p_{j|i} + \frac{1}{2\sigma^2} \sum_{j \neq i} p_{j|i} ||x_i - x_j||^2) \\
&= \log(S_i) + \frac{1}{2\sigma^2} \sum_{j \neq i} p_{j|i} ||x_i - x_j||^2
\end{aligned}
\tag{1}
$$

Where $S_i = \sum_{k \neq i} e^{-||x_i-x_j||^2/2\sigma^2}$ and $\sum_{j \neq i} p_{j|i} = 1$

In order to proceed with the optimization of the variance, we are going to define this term $\frac{1}{2\sigma^2}$ as the parameter β.

```r
entropy_beta <- function(D_i, beta=1) {
  P_i <- exp(-D_i * beta)
  sum_p_i <- sum(P_i)
  H_i <- log(sum_p_i) + (beta * sum(D_i * P_i) /sum_p_i)
  P_i <- P_i / sum_p_i
  return(list(entropy=H_i, probs=P_i))
}
```

The goal is to adjust the variability so that the perplexity at each point is the same. The perplexity is a way to measure the effective number of neighbors of a point. We are going to perform a binary search to get the probabilities in such a way that the conditional Gaussian has the same perplexity.

```r
index_except_i <- function(i, n) {
  index <- c(seq(1,i-1),seq(i+1,n))
  if (i == 1) {
    index <- 2:n
  } else if (i == n) {
    index <- 1:(n-1)
  }
  return(index)
}


binary_search <- function(h_diff, beta, i, beta_min, beta_max) {
  if(h_diff > 0) {
    beta_min = beta[i]
    if(beta_max == -Inf || beta_max == Inf) {
      beta[i] <- beta[i] * 2
    } else {
```

```r
      beta[i] <- (beta[i] + beta_max) / 2
    }
  } else {
    beta_max = beta[i]
    if(beta_min == -Inf || beta_min == Inf) {
      beta[i] <- beta[i] / 2
    } else {
      beta[i] <- (beta[i] + beta_min) / 2
    }
  }
  return(list(beta=beta, min=beta_min, max=beta_max))
}

binary_search_optimization <- function(D_i, i, beta, h_star, prob_star, log_perp,
                                       tolerance=1e-5) {
  beta_min <- -Inf
  beta_max <- Inf
  tries <- 0
  h_diff <- h_star - log_perp

  while(abs(h_diff) > tolerance && tries < 50) {
    beta_opt <- binary_search(h_diff, beta, i, beta_min, beta_max)
    beta <- beta_opt$beta; beta_min <- beta_opt$min; beta_max <- beta_opt$max

    res_loop <- entropy_beta(D_i, beta[i])
    h_star <- res_loop$entropy; prob_star <- res_loop$probs

    h_diff <- h_star - log_perp
    tries <- tries + 1
  }
  return(list(probs=prob_star, beta=beta))
}
```

Once we have defined these two methods, we are able to obtain the high dimensional properties:

```r
high_dimension_probs <- function(X=matrix(), tolerance=1e-5, perplexity=30) {
  n <- nrow(X)
  p <- ncol(X)

  D <- x_diff(X)

  P <- matrix(0, nrow=150, ncol=150)
  beta <- rep(1, n)
  log_perp <- log(perplexity)

  for(i in seq_len(n)) {
    column_index <- index_except_i(i, n)
    D_i <- D[i, column_index]

    res <- entropy_beta(D_i, beta[i])
    h_star <- res$entropy
    prob_star <- res$probs

    h_diff <- h_star - log_perp
```

```
    res_opt = binary_search_optimization(D_i, i, beta, h_star, prob_star,
                                          log_perp, tolerance)
    prob_star = res_opt$probs; beta = res_opt$beta

    P[i, column_index] <- prob_star
  }
  print("Values of sigma for each x_i")
  print(sqrt(1/beta))
  print(sprintf("Mean value of sigma: %01.2f", mean(sqrt(1/beta))))
  return(P)
}
```

The version of the t-SNE is the symmetric one, that has the property that $p_{ij} = p_{ji}$ and $q_{ij} = q_{ji}$ $\forall i, j$. Therefore, we define $p_{ij} = \frac{p_{i|j} + p_{j|i}}{2n}$.

```
symmetric_probs <- function(P) {
  P = (P + t(P)) / (2*nrow(P))
  return(P)
}
```

In order to initialize the lower dimension probability matrix, we are going to use the method `mvtnorm::rmvnorm` as it is described in the paper: $\mathcal{Y}^0 = \{y_1, ..., y_n\} \sim \mathcal{N}(0, 10^{-4}\mathbf{I}_n)$ which is assigned to $\mathcal{Y}^1$ and $\mathcal{Y}^2$ (the first two initial states).

Gradient Descent

$$
\begin{aligned}
C &= KL(\mathbf{P}\|\mathbf{Q}) \\
&= \sum_i \sum_j p_{ij} \log \frac{p_{ij}}{q_{ij}} \\
&= \sum_i \sum_j p_{ij} \left(\log p_{ij} - \log q_{ij} \right) \\
&= \sum_i \sum_j p_{ij} \log p_{ij} - p_{ij} \log q_{ij}
\end{aligned}
\tag{2}
$$

We define these two auxiliary variables $d_{ij} = \|\mathbf{y}_i - \mathbf{y}_j\|$ and $Z = \sum_{k \neq l} \left(1 + d_{kl}^2\right)^{-1})$

$$
\begin{aligned}
\frac{\partial C}{\partial \mathbf{y}_i} &= \sum_{j \neq i} \left[\frac{\partial C}{\partial d_{ij}} \frac{\partial d_{ij}}{\partial \mathbf{y}_i} + \frac{\partial C}{\partial d_{ji}} \frac{\partial d_{ji}}{\partial \mathbf{y}_i} \right] \\
&= \sum_{j \neq i} \left[\frac{\partial d_{ij}}{\partial \mathbf{y}_i} \left(\frac{\partial C}{\partial d_{ij}} + \frac{\partial C}{\partial d_{ji}} \right) \right]
\end{aligned}
\tag{3}
$$

Recall that $\frac{\partial}{\partial x} \sqrt{g(x)} = \frac{1}{2\sqrt{g(x)}} g'(x)$, $\frac{\partial}{\partial \mathbf{y}} \|\mathbf{y}\|^2 = 2\mathbf{y}$ and $d_{ij} = d_{ji}$

$$\begin{aligned}
\frac{\partial d_{ij}}{\partial \mathbf{y}_i} &= \frac{\partial}{\partial y_i} \|\mathbf{y}_i - \mathbf{y}_j\| \\
&= \frac{\partial}{\partial \mathbf{y}_i} (\|\mathbf{y}_i\|^2 + \|\mathbf{y}_j\|^2 - 2\mathbf{y}_i'\mathbf{y}_j)^{\frac{1}{2}} \\
&= \frac{1}{2} \frac{1}{d_{ij}} \frac{\partial}{\partial \mathbf{y}_i} (\|\mathbf{y}_i\|^2 + \|\mathbf{y}_j\|^2 - 2\mathbf{y}_i'\mathbf{y}_j) \\
&= \frac{1}{2} \frac{1}{d_{ij}} (2\mathbf{y}_i - 2\mathbf{y}_j) \\
&= \frac{(\mathbf{y}_i - \mathbf{y}_j)}{d_{ij}} \\
&= \frac{\partial d_{ji}}{\partial \mathbf{y}_i}
\end{aligned} \tag{4}$$

```r
dij.1 <- function(i, j) {
  sqrt(norm(i, type="2")^2 + norm(j, type="2")^2 - 2 * (t(i) %*% j))
}


dij.2 <- function(i, j) {
  norm(i-j, type="2")
}


y <- data.matrix(iris)[, -5]

# For rows 2 and 3
result1 <- as.numeric(round(jacobian(func=dij.2, x=y[2,], j=y[3,]), digits=7))
result2 <- as.numeric((y[2,]-y[3,])/norm(y[2,]-y[3,], type="2"))
all.equal(result2, result1) # checked
```

```
## [1] "Mean relative difference: 6e-08"
```

```r
# For row 3 and 2
result3 <- as.numeric(round(jacobian(func=dij.2, x=y[3,], i=y[2,]), digits=7))
result4 <- as.numeric((y[3,]-y[2,])/norm(y[2,]-y[3,], type="2"))
all.equal(result4, result3) # checked
```

```
## [1] "Mean relative difference: 6e-08"
```

```r
# Checked that both d(d_ij)/dy_i = d(d_ji)/dy_i
```

Recall that $d_{ij} = \|\mathbf{y}_i - \mathbf{y}_j\|$ and $Z = \sum_{k \neq l} \left(1 + d_{kl}^2\right)^{-1}$:

$$\frac{\partial C}{\partial d_{ij}} = \frac{\partial}{\partial d_{ij}} \sum_{k \neq l} \cancel{p_{kl} \log p_{kl}} - p_{kl} \log q_{kl}$$

$$= \frac{\partial}{\partial d_{ij}} \sum_{k \neq l} -p_{kl} \log q_{kl} = -\sum_{k \neq l} p_{kl} \frac{\partial (\log q_{kl})}{\partial d_{ij}}$$

$$= -\sum_{k \neq l} p_{kl} \frac{\partial (\log \frac{q_{kl} Z}{Z})}{\partial d_{ij}}$$

$$= -\sum_{k \neq l} p_{kl} \left[\frac{\partial (\log q_{kl} Z - \log Z)}{\partial d_{ij}} \right]$$

$$= -\sum_{k \neq l} p_{kl} \left[\frac{\partial (\log q_{kl} Z)}{\partial d_{ij}} - \frac{\partial \log Z}{\partial d_{ij}} \right]$$

$$= -\sum_{k \neq l} p_{kl} \left[\frac{1}{q_{kl} Z} \frac{\partial (q_{kl} Z)}{\partial d_{ij}} - \frac{1}{Z} \frac{\partial Z}{\partial d_{ij}} \right] \tag{5}$$

$$= -\sum_{k \neq l} p_{kl} \left[\frac{1}{q_{kl} Z} \frac{\partial \left(\frac{(1+d_{ij}^2)^{-1}}{\cancel{\sum_{k \neq l}(1+d_{kl}^2)^{-1}}} \cancel{\sum_{k \neq l}(1+d_{kl}^2)^{-1}} \right)}{\partial d_{ij}} - \frac{1}{Z} \frac{\partial (\sum_{k \neq l}(1+d_{kl}^2)^{-1})}{\partial d_{ij}} \right]$$

$$= 2 \frac{p_{ij}}{q_{ij} Z} (1+d_{ij}^2)^{-2} d_{ij} - 2 \sum_{k \neq l} p_{kl} \frac{(1+d_{ij}^2)^{-1}}{Z} (1+d_{ij}^2)^{-1} d_{ij}$$

$$= 2 \frac{p_{ij}}{\frac{(1+d_{ij}^2)^{-1}}{\cancel{Z}} \cancel{Z}} (1+d_{ij}^2)^{-\cancel{2}1} d_{ij} - 2 \sum_{\cancel{k \neq l}} \cancel{p_{kl}} q_{ij} (1+d_{ij}^2)^{-1} d_{ij}$$

$$= 2 p_{ij}(1+d_{ij}^2)^{-1} d_{ij} - 2 q_{ij}(1+d_{ij}^2)^{-1} d_{ij}$$

$$= 2(p_{ij} - q_{ij})(1+d_{ij}^2)^{-1} d_{ij}$$

$$\frac{\partial C}{\partial y_i} = \sum_{j \neq i} \left[\frac{\partial C}{\partial d_{ij}} \frac{\partial d_{ij}}{\partial y_i} + \frac{\partial C}{\partial d_{ji}} \frac{\partial d_{ji}}{\partial y_i} \right]$$

$$= \sum_{j \neq i} \left[\frac{\partial d_{ij}}{\partial y_i} \left(\frac{\partial C}{\partial d_{ij}} + \frac{\partial C}{\partial d_{ij}} \right) \right]$$

$$= 2 \sum_{j \neq i} \left[\frac{\partial C}{\partial d_{ij}} \right] \frac{\mathbf{y}_i - \mathbf{y}_j}{d_{ij}} \tag{6}$$

$$= 2 \sum_{j \neq i} \left[2(p_{ij} - q_{ij})(1+d_{ij}^2)^{-1} d_{ij} \right] \frac{\mathbf{y}_i - \mathbf{y}_j}{d_{ij}}$$

$$= 4 \sum_{j \neq i} (p_{ij} - q_{ij})(1+d_{ij}^2)^{-1} \cancel{d_{ij}} \frac{\mathbf{y}_i - \mathbf{y}_j}{\cancel{d_{ij}}}$$

$$= 4 \sum_{j \neq i} (p_{ij} - q_{ij})(1+\|\mathbf{y}_i - \mathbf{y}_j\|^2)^{-1} (\mathbf{y}_i - \mathbf{y}_j)$$

Cauchy distribution on the sphere

The following formula corresponds with the probability density function of the Cauchy family on the unit sphere:

$$f(\mathbf{y}; \boldsymbol{\mu}, \rho) = \frac{\Gamma\{(d+1)/2\}}{2\pi^{(d+1)/2}} \left(\frac{1 - \rho^2}{1 + \rho^2 - 2\rho\boldsymbol{\mu}'\mathbf{y}} \right)^d$$

where $\mathbf{y} \in S^d$, the location parameter $\boldsymbol{\mu} \in S^d$, the concentration parameter $\rho \in [0, 1)$ and the unit sphere in $\mathbb{R}^{d+1}$ denoted by $S^d = \{\mathbf{y} \in \mathbb{R}^{d+1}; \|\mathbf{y}\| = 1\}$. When $d = 1$ the case is the well-known Wrapped Cauchy or circular Cauchy family.

```
dsphcauchy <- function(y, mu, rho, d=2) {
  (gamma(((d+1)/2))/(2*pi^((d+1)/2))*((1-rho^2)/(1+rho^2-2*rho*(t(mu) %*% y)))^d)
}

library(Directional)
library(Rcpp)
library(rotasym)

sunspots_births$X <-
  cbind(cos(sunspots_births$phi) * cos(sunspots_births$theta),
        cos(sunspots_births$phi) * sin(sunspots_births$theta),
        sin(sunspots_births$phi))

theta_params <- spcauchy.mle(sunspots_births$X)
dsphcauchy(sunspots_births$X[1,], theta_params$mu, theta_params$rho)
```

```
##              [,1]
## [1,] 0.07968743
```

```
library(DirStats)
library(ivdoctr)
n <- nrow(sunspots_births$X)
q <- 2
x <- rbind(diag(1, nrow = q + 1), diag(-1, nrow = q + 1))
bw_rot <- bw_dir_rot(sunspots_births$X)


th <- sample(seq(0, pi, l=2000), size=1000)
ph <- sample(seq(0, 2*pi, l=2000), size=1000)
x <- to_sph(th, ph)
rgl::plot3d(0, 0, 0, xlim = c(-1, 1), ylim = c(-1, 1), zlim = c(-1, 1),
            radius = 1, type = "s", col = "lightblue", alpha = 0.25,
            lit = FALSE)
# dens <- apply(x, MARGIN=1, FUN=dsphcauchy, mu=theta_params$mu, rho=theta_params$rho)
dens <- apply(x, MARGIN=1, FUN=kde_dir, data = sunspots_births$X, h = bw_rot, L = NULL)
map2color <- function(x, pal, limits = range(x)){
  pal[findInterval(x, seq(limits[1], limits[2], length.out = length(pal) + 1),
                   all.inside=TRUE)]
}
rgl::points3d(x, col = map2color(dens, pal=heat.colors(10, alpha=0.8)))
```

Cauchy-SNE

High Dimension   For a poly-sphere $d > 2$:

$$p_{j|i} = \frac{p_{ji}}{p_i} = \frac{\frac{\Gamma\{(d+1)/2\}}{2\pi^{(d+1)/2}} \left(\frac{1-\rho^2}{1+\rho^2-2\rho\boldsymbol{\mu}'S_c(\mathbf{x}_i,\mathbf{x}_j)}\right)^d}{\frac{\Gamma\{(d+1)/2\}}{2\pi^{(d+1)/2}} \sum_{k\neq l} \left(\frac{1-\rho^2}{1+\rho^2-2\rho\boldsymbol{\mu}'S_c(\mathbf{x}_l,\mathbf{x}_k)}\right)^d} \tag{7}$$

where the cosine similarity is denoted by $S_c(\mathbf{x}_i, \mathbf{x}_j) = \cos(\theta) = \frac{\mathbf{x}_i \mathbf{x}_j}{\|\mathbf{x}_i\|\|\mathbf{x}_j\|}$.

We must adapt the configuration of each parameter to have the same perplexity fixed at the beginning, in the same way we had done with the Gaussian case.