

# Machine Learning 1

## Solved Problems

### 1 Team members:

- Luis Sierra Muntané (luis.sierra@est.fib.upc.edu)
- Àlex Batlle Casellas (alex.batlle@est.fib.upc.edu)
- Aleix Torres i Camps (aleix.torres.camps@est.fib.upc.edu)

### 2 Problems

1. You may need the following procedures for several exercises below.

- (a) Write a procedure to generate random samples according to a normal distribution  $\mathcal{N}(\mu, \Sigma)$  in  $d$  dimensions.

SOLUTION:

Based on the information in Wikipedia's Multivariate Normal Distribution page, and in particular in [this part](#), we see that a  $d$ -dimensional vector  $X$  is normally distributed (it is a *normal random vector*) iff there exist a vector  $\mu$ , a matrix  $A$  of coefficients and a standard normal vector  $Z$  (meaning, it follows a standard normal distribution and the components are independent) such that  $AZ + \mu = X$ , with the covariance matrix being  $\Sigma = AA^T$ . So, we will take advantage of all this situation:

- We can calculate the matrix  $A$  by simply decomposing  $\Sigma$  (which is the matrix we are given) using the Cholesky decomposition method, which factors a symmetric positive-definite matrix into a product of a lower triangular matrix and its transpose. R has a built in function named `chol`, although this gives the upper triangular part of the decomposition.
- We just have to calculate a random vector  $Z$ , and we will do so using the function `rnorm` already included in R.
- The reason we are not doing this straight away is because of  $\Sigma$ : if it is the identity matrix, or a multiple of the identity  $\sigma^2 I_d$ , we could easily get a normally distributed  $d$ -dimensional vector by using `rnorm(d, mu, sigma)`, but as this is not necessarily the case and there could be positive covariance between the components, we have to think another way of working.

So, the solution code is this one:

```
normal = function(mu, sigma) {  
  d = length(mu)  
  L = t(chol(sigma))  
  res = rnorm(d)  
  return(L %*% res + mu)  
}
```

- (b) Write a procedure to calculate the discriminant function (of the form given in Eq. 47) for a given normal distribution and prior probability  $P(\omega_i)$ .

SOLUTION:

We will suppose we are given the probability of class  $i$ , and the  $\mu_i, \Sigma_i$  such that  $p(x|\omega_i) \sim \mathcal{N}(\mu_i, \Sigma_i)$ . Given the formula in Eq. 47,

$$g_i(x) = -\frac{1}{2}(x - \mu_i)^T \Sigma_i^{-1}(x - \mu_i) - \frac{d}{2} \ln 2\pi - \frac{1}{2} \ln \det \Sigma_i + \ln P(\omega_i),$$

the implementation is quite straight forward:

```
calc_discr = function(x, p, mu, sigma) {
  xnorm = x - mu
  inv = solve(sigma)
  d = length(mu)
  dt = det(sigma)
  return(-0.5*t(xnorm)%*%inv%*%xnorm-d/2*log(2*pi)-0.5*log(dt)+log(p))
}
```

We are using the function `solve`, which R has built-in, that solves linear systems given  $A, b$  such that  $Ax = b$ . If  $b$  is absent, the function returns the inverse of a matrix.

- (c) Write a procedure to calculate the Euclidean distance between two arbitrary points.

SOLUTION:

We will write the solution for the general  $p$ -norm distance:

```
# pre: length(x)=length(y)
euc = function(x, y, p = 2) {
  n = length(x)
  sum = 0
  for (i in 1:n) {
    sum = sum + abs(x[i]-y[i])^p
  }
  return(sum^(1/p))
}
```

Note that we could have also used the `dist` function included in R, by row-binding the coordinates of  $x$  and  $y$  into a matrix.

- (d) Write a procedure to calculate the Mahalanobis distance between the mean  $\mu$  and an arbitrary point  $x$ , given the covariance matrix.

SOLUTION:

We will be using the matrix formula for the Mahalanobis distance, which is the square root of a quadratic form, namely

$$\delta_M(x, y) = \sqrt{(x - y)^T \Sigma^{-1} (x - y)}.$$

```
mah = function(x, mu, sigma) {
  xnorm = x - mu
  return(sqrt(t(xnorm)%*%solve(sigma)%*%xnorm))
}
```