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1 Problem 1

- a. The probability that all of the M dimensions of $x - y$ are between $-\epsilon$ and ϵ is $\boxed{\rho = (2\epsilon)^M}$.
For each dimension i of χ , the probability that $|x_i - y_i| \leq \epsilon$ is equivalent to

$$\begin{aligned} P(|x_i - y_i| \leq \epsilon) &= \\ P(-\epsilon \leq x_i - y_i \leq \epsilon) &= \\ P(-\epsilon - x_i \leq -y_i \leq \epsilon - x_i) &= \\ P(\epsilon + x_i \geq y_i \geq x_i - \epsilon) &= \\ P(x_i - \epsilon \leq y_i \leq \epsilon + x_i) & \end{aligned}$$

This distribution function is equivalent to $\int_{\epsilon+x_i}^{x_i-\epsilon} f(x)dx$, where $f(x)$ is the PDF of y_i , which we know to have a uniform distribution, so $f(x) = \frac{1}{b-a} = 1$. Thus, we get:

$$\begin{aligned} \int_{\epsilon+x_i}^{x_i-\epsilon} 1dx &= \\ \epsilon + x_i - (x_i - \epsilon) &= 2\epsilon \end{aligned}$$

Because we want to know the probability that all of the M dimensions of $x - y$ are between $-\epsilon$ and ϵ , we simply take $\prod_{i=1}^M P(|x_i - y_i| \leq \epsilon) = (2\epsilon)^M$.

- b. The probability of $\max_m |x_m - y_m| \leq \epsilon$ is at most ρ because as shown in (a), ρ does not depend on x_i and thus holds for all x_i . In addition, logically, if x is the center point, the average distance from it to any other point y is at most $\frac{1}{2}$ for any one dimension. As x moves farther and farther away from the center, the average distance increases so that it becomes at most 1 in any one dimension. So, if x is not in the center $\max_m |x_m - y_m|$ grows and is less likely to be less than ϵ , decreasing that probability so that it is less than ρ .
- c. We will show that $\|x - y\| \geq \max_m |x_m - y_m|$.

$$\begin{aligned} \|x - y\| &= \sqrt{\sum_{m=1}^M (x_m - y_m)^2} \\ \sqrt{\sum_{m=1}^M (x_m - y_m)^2} &\geq \max_m |x_m - y_m| \\ \sum_{m=1}^M (x_m - y_m)^2 &\geq (\max_m |x_m - y_m|)^2 \end{aligned}$$

This is true because the left side of the inequality includes the right side in its sum. $\|x - y\|$ is the total Euclidean distance between two points whereas $\max_m |x_m - y_m|$ is only the distance between one dimension of two points. The left side must be larger.

If x is any point in χ , and y is a point in χ drawn randomly from a uniform distribution on χ , then the probability that $\|x - y\| \leq \epsilon$ is also at most ρ because $\|x - y\|$ is greater than or equal to $\max_m |x_m - y_m|$, making it less likely to be less than ϵ and thus giving it a probability lower than ρ of being less than ϵ .

- d. Lowerbound on number N of points needed to guarantee
- e. We can conclude that the effectiveness of the hierarchical agglomerative clustering algorithm in high dimensional spaces

2 Problem 2

a Given a prior distribution $Pr(\theta)$ and likelihood $Pr(D|\theta)$, the predictive distribution $Pr(x|D)$ for a new datum,

(a) ML: $Pr(x|D) = Pr(x|\theta) = \arg \max_{\theta} (\ln(Pr(D|\theta)))$

(b) MAP: $Pr(x|D) = Pr(x|\theta) = \arg \max_{\theta} (\ln(P(D|\theta)P(\theta)))$

(c) FB: $Pr(x|D) = \int \theta P(\theta|D) d\theta$

b MAP can be considered "more Bayesian" than ML because it takes into account the distribution of θ instead of assuming same weight or uniformity.

c One advantage the MAP method enjoys over the ML method

d The Beta distribution is the conjugate prior of the Bernoulli.

e Under the ML approach

3 Problem 3

a The K -means clustering objective is to minimize the sum of squared distances between prototype and data.

b PCA relates to K -means

4 Problem 4