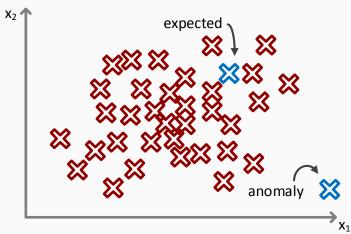
anomaly detection 🔁 basics

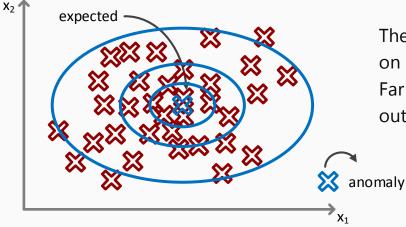
density estimation

problem motivation



when given a dataset of features $\{x^{(1)}, x^{(2)}, ..., x^{(m)}\}$, the points plotted (red) represent the data correlations. When a new dataset is added to the model (x_{test}) , points that fall within the majority or expected, but any far reaching points are considered to be an anomaly

formally, a model p(x) will determine anomaly detection: $p(x_{test}) < \varepsilon$ anomaly $(x_{test}) \ge \varepsilon$ expected

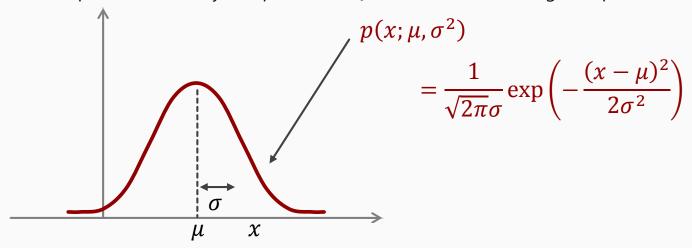


The model p(x) will set thresholds based on logic acceptable to the given data. Far reaching points will be labeled as an outlier or anomaly.

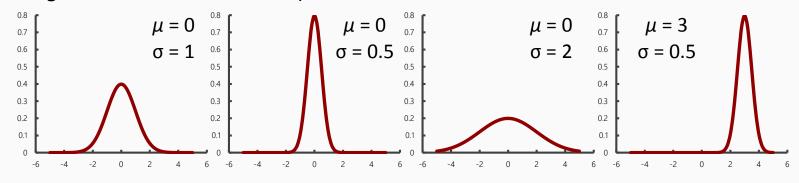
gaussian (normal) distribution

 $x \in \mathbb{R}$. if x is a distributed gaussian with mean μ , variance σ^2 , denoted as $x \sim \mathcal{N}(\mu, \sigma^2)$

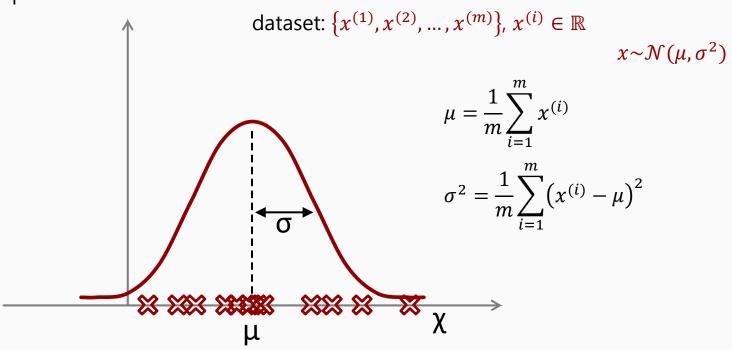
x represents a row values random variable (x is a row number). \sim tilde represents the variable "distributed as". The script N represents the "normal" gaussian distribution and is parameterized by two parameters μ mu (mean) and σ^2 sigma squared (variance)



gaussian distribution examples



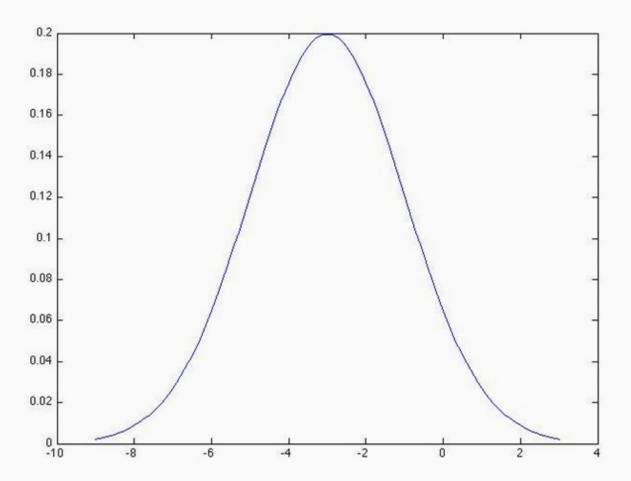
parameter estimation



The formula for the Gaussian density is:

$$p(x) = rac{1}{\sqrt{2\pi}\sigma} \exp\left(-rac{(x-\mu)^2}{2\sigma^2}
ight)$$

Which of the following is the formula for the density to the right?



$$p(x)=rac{1}{\sqrt{2\pi} imes2}\exp\!\left(-rac{(x-3)^2}{2 imes4}
ight)$$

$$p(x)=rac{1}{\sqrt{2\pi} imes 4}\exp\left(-rac{(x-3)^2}{2 imes 2}
ight)$$

$$^{ ext{\tiny{(a)}}} \ p(x) = rac{1}{\sqrt{2\pi} imes 2} \expigg(-rac{(x+3)^2}{2 imes 4}igg)$$

Correct Response

$$p(x)=rac{1}{\sqrt{2\pi} imes 4}\exp\left(-rac{(x+3)^2}{2 imes 2}
ight)$$

anomaly detection algorithm

$$\begin{array}{ll} \text{density estimation} & x \sim \mathcal{N}(\mu_1, \sigma_1^2) \\ \text{dataset: } \{x^{(1)}, x^{(2)}, \dots, x^{(m)}\}, & x \sim \mathcal{N}(\mu_2, \sigma_2^2) \\ \text{each example is } x^{(i)} \in \mathbb{R}^n & x \sim \mathcal{N}(\mu_3, \sigma_3^2) \end{array}$$

$$p(x) = p(x_1; \mu_1, \sigma_1^2) p(x_2; \mu_2, \sigma_2^2) p(x_3; \mu_3, \sigma_3^2) \dots p(x_n; \mu_n, \sigma_n^2)$$

$$= \prod_{j=1}^{n} p(x_j; \mu_j, \sigma_j^2)$$

to describe the product operator in compassion to the summation operator:

$$\sum_{i=1}^{n} 1 + 2 + 3 + \dots + n \qquad \text{versus} \qquad \prod_{i=1}^{n} 1 \times 2 \times 3 \times \dots \times n$$

estimating the above distribution p(x) is referred to as the density estimation problem

given training set $\{x^{(1)}, x^{(2)}, ..., x^{(m)}\}$, each would be estimated as (note $\mu_j \in \mathbb{R}, \sigma_j^2 \in \mathbb{R}$):

$$\mu_j = \frac{1}{m} \sum_{i=1}^m x_j^{(i)}$$
 and $\sigma_j^2 = \frac{1}{m} \sum_{i=1}^m (x_j^{(i)} - \mu_j)^2$

steps in the anomaly detection algorithm

choose features x_i that might be indicative of anomalous examples: $\{x^{(1)}, x^{(2)}, ..., x^{(m)}\}$ fit parameters $\mu_1, ..., \mu_n, \sigma_1^2, ..., \sigma_n^2$ vectorized version:

$$\mu_{j} = \frac{1}{m} \sum_{i=1}^{m} x_{j}^{(i)} \qquad p(x_{j}; \mu_{j}, \sigma_{j}^{2}) \qquad u_{1}, \mu_{2}, \dots \mu_{n}$$

$$\sigma_{j}^{2} = \frac{1}{m} \sum_{i=1}^{m} (x_{j}^{(i)} - \mu_{j})^{2} \qquad \mu = \begin{bmatrix} \mu_{1} \\ \mu_{2} \\ \vdots \\ \mu_{n} \end{bmatrix} = \frac{1}{m} \sum_{i=1}^{m} x^{(i)}$$

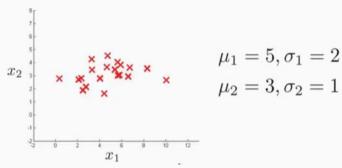
given new example x, compute p(x):

$$p(x) = \prod_{j=1}^{n} p(x_j; \mu_j, \sigma_j^2) = \prod_{j=1}^{n} \frac{1}{\sqrt{2\pi}\sigma^2} \exp\left(-\frac{(x_j - \mu_j)^2}{2\sigma_j^2}\right)$$

anomaly if $p(x) < \varepsilon$

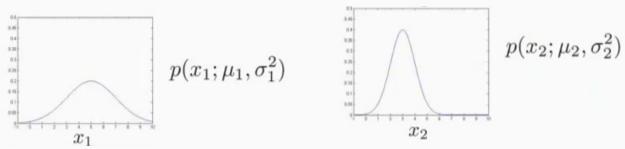
anomaly detection example

The dataset below has an average measured feature x_1 of ~5 with a standard deviation of ~2 and the feature x_2 has an average value of ~3 with a standard deviation ~1

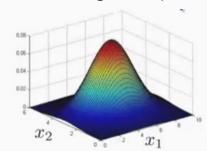


Note the use of σ (standard deviation) as opposed to σ^2 (variance) used previously

The features above plotted as in terms of $p(x_i; \mu_i, \sigma_i^2)$ distribution would appear as:



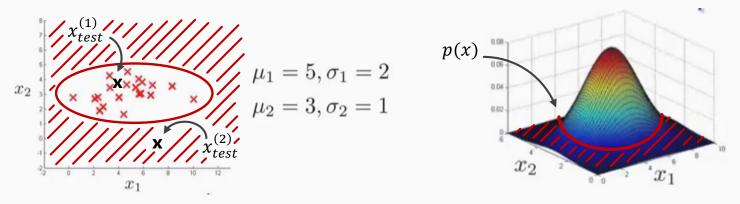
p(x) for both can be taken and plotted in a surface plot as seen below. The height of the surface given a particular values is p(x):



the height of the plot is literally equal to:

$$p(x) = p(x_1; \mu_1, \sigma_1^2) \times p(x_2; \mu_2, \sigma_2^2)$$

if additional data points are added to the dataset:



to determine if either is an anomaly: a value is set for epsilon ε , (in this case, $\varepsilon = 0.02$):

$$p\left(x_{test}^{(1)}\right) = 0.0426 \ge \varepsilon$$
 (non-anomaly) and $p\left(x_{test}^{(2)}\right) = 0.0021 < \varepsilon$ (anomaly)