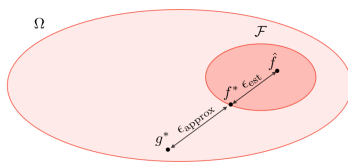

$$\mathcal{F} = \{f_w : w \in \mathbb{R}^d\}$$
$$\forall z \in]-\infty, +\infty[, \quad \sigma(z) = \frac{1}{1 + e^{-z}}$$




Remark: we have $\sigma'(z) = \sigma(z)(1 - \sigma(z))$.

The diagram shows a central blue circle labeled f_i . A green arrow points from left to right, labeled x above and $f_i(x)$ above. A red arrow points from right to left, labeled $\frac{\partial f_i(x)}{\partial x} \cdot g_i(x)$ below and $g_i(x) \triangleq \frac{\partial \text{out}}{\partial f_i(x)}$ below.

□ **Approximation and estimation error** — The approximation error ϵ_{approx} represents how far the entire hypothesis class \mathcal{F} is from the target predictor g^* , while the estimation error ϵ_{est} quantifies how good the predictor \hat{f} is with respect to the best predictor f^* of the hypothesis class \mathcal{F} .



❑ **Regularization** — Regularization aims to keep the model from overfitting to the data and thus deals with high variance issues. The following table sums up the different types of commonly used regularization techniques:

LASSO	Ridge	Elastic Net
<ul style="list-style-type: none"> Shrinks coefficients to 0 Good for variable selection 	Makes coefficients smaller	Tradeoff between variable selection, and small coefficients
 <p>$\beta_j _1 \leq t$</p>	 <p>$\beta_j _2 \leq t$</p>	 <p>$\beta_j _1 - \alpha \beta_j _2 \leq t$</p>
$-\alpha w _1$ $\lambda \in \mathbb{R}^+$	$-\alpha w _2^2$ $\lambda \in \mathbb{R}^+$	$-\alpha 1 - \alpha _2 w _2 + \alpha w _2^2$ $\lambda \in \mathbb{R}^+, \alpha \in \mathbb{R}, t \in \mathbb{R}^+$

□ **Hyperparameters** — Hyperparameters are the properties of the learning algorithm, and include architecture-related features, the regularization parameter λ , number of iterations T , step size η , etc.

□ **Sets vocabulary** — When selecting a model, we distinguish 3 different parts of the data that we have as follows:

Training set	Validation set	Testing set
<ul style="list-style-type: none"> Model is trained Usually 80% of the dataset 	<ul style="list-style-type: none"> Model is assessed Usually 20% of the dataset Also called hold-out or development set 	<ul style="list-style-type: none"> Model gives predictions Unseen data

Once the model has been chosen, it is trained on the entire dataset and tested on the unseen test set. These are represented in the figure below:



For a more detailed overview of the concepts above, check out the [Machine Learning tips and tricks cheatsheets \(teaching/cs-229/cheatsheet-machine-learning-tips-and-tricks\)](#)!

(<https://stanford.edu/~shervine/teaching/cs-221/cheatsheet-reflex-models#unsupervised-learning>)

The class of unsupervised learning methods aims at discovering the structure of the data, which may have rich latent structures.

k-means

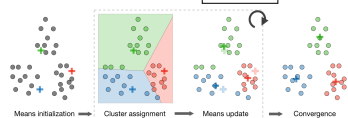
□ **Clustering** — Given a training set of n input points $\mathcal{D}_{\text{train}}$, the goal of a clustering algorithm is to assign each point $\phi(z_i)$ to a cluster $z_i \in \{1, \dots, k\}$.

□ **Objective function** — The loss function for one of the main clustering algorithms, *k*-means, is given by

$$\text{Loss}_{\text{re-embed}}(z, \mu) = \sum_{i=1}^n \|\phi(x_i) - \mu_{zi}\|^2$$

□ **Algorithm** — After randomly initializing the cluster centroids $\mu_1, \mu_2, \dots, \mu_k \in \mathbb{R}^d$, the k -means algorithm repeats the following step until convergence:

$$z_i = \arg \min_j \|\phi(x_i) - \mu_j\|^2 \quad \text{and} \quad \mu_j = \frac{\sum_{i \in \{i: j\}} \phi(x_i)}{\sum_{i \in \{i: j\}} 1}$$



Principal Component Analysis

□ **Eigenvalue, eigenvector** — Given a matrix $A \in \mathbb{R}^{d \times d}$, λ is said to be an eigenvalue of A if there exists a vector $z \in \mathbb{R}^d \setminus \{0\}$, called eigenvector, such that we have:

$$Ax = \lambda x$$

□ **Spectral theorem** — Let $A \in \mathbb{R}^{d \times d}$. If A is symmetric, then A is diagonalizable by a real orthogonal matrix $U \in \mathbb{R}^{d \times d}$. By noting $\Lambda = \text{diag}(\lambda_1, \dots, \lambda_d)$, we have:

$\exists \Lambda \text{ diagonal, } A = U\Lambda U^T$

Remark: the eigenvector associated with the largest eigenvalue is called *principal eigenvector* of matrix A

□ **Algorithm** — The Principal Component Analysis (PCA) procedure is a dimension reduction technique that projects the data on k dimensions by maximizing the variance of the data as follows:

$$\phi_j(x_i) \leftarrow \frac{\phi_j(x_i) - \mu_j}{\sigma_j} \quad \text{where} \quad \mu_j = \frac{1}{n} \sum_{i=1}^n \phi_j(x_i) \quad \text{and} \quad \sigma_j^2 = \frac{1}{n} \sum_{i=1}^n (\phi_j(x_i) - \mu_j)^2$$

- **Step 2:** Compute $\Sigma = \frac{1}{n} \sum_{i=1}^n \phi(x_i) \phi(x_i)^T \in \mathbb{R}^{d \times d}$, which is symmetric with real eigenvalues.

- **Step 3:** Compute $u_1, \dots, u_k \in \mathbb{R}^d$ the k orthogonal principal eigenvectors of Σ , i.e. the orthogonal eigenvectors of the k largest eigenvalues.

- Step 4: Project the data on $\text{span}_2(u_1, \dots, u_k)$.

This procedure maximizes the variance among all k -dimensional spaces.



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