

# Data analytics for personalized genomics and precision medicine

## Lecture 9: Clustering and classification performance evaluation

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### Outline of the Lecture:

- ❖ Performance Evaluation
- ❖ Cross-Validation
- ❖ Multi-class Classification
- ❖ Clustering Evaluation

### Content:

#### ❖ Training

- What is training ?
  - To get  $w_h, w_w, w_0$
  - To make model fit the training data
  - To make  $\frac{1}{1+e^{-(w_h H + w_w W + w_0)}} \geq 0.5$  correct

➤  $Y^{\text{output}} = \frac{1}{1+e^{-(w_h H + w_w W + w_0)}}$

#### ➤ Loss function L

- Loss function =  $(Y^{\text{output}} - Y)^2$
- Example: Data from P1-P4

Person	Height	Weight	Gender
P1	0.625	0.875	M

- For P1,
- $L = (Y^{\text{output}} - Y)^2 = \left(1 - \frac{1}{1+e^{-(w_h H + w_w W + w_0)}}\right)^2$
- In a whole,  $L = \sum_{P_1}^{P_4} (Y^{\text{output}} - Y)^2$
- L = the smaller, the better

### ❖ Gradient descent algorithm

➤ Aim: find smallest value for L using different values of w

➤ Step 1: initialise  $w_w, W_w, w_0$

Step 2: For each data (P1,P2,P3,P4)

Calculate  $Y^{output}$

Update new weights

New  $w_i = w_i + \Delta w_i$

$\Delta w_i = 2\alpha(Y - Y^{output}) \frac{\partial Y^{output}}{\partial w_i}$ ,  $\alpha$  is a constant

Repeat until no update

### ❖ Performance evaluation -binary classification evaluation

➤ Purpose: pinpoint strong points and weak points of one method → model selection

➤ Method: confusion matrix

		Predicted class	
		Class=Yes	Class=No
Actual class	Class=Yes	a(TP)	b(FN)
	Class=No	c(FP)	d(TN)

➤ Accuracy =  $\frac{a+d}{a+b+c+d} = \frac{TP+TN}{TP+TN+FP+FN}$

➤ Higher accuracy, better the classifier is

➤ Exception: when there is imbalanced classes

➤ Precision =  $\frac{a}{a+c}$  Recall =  $\frac{a}{a+b}$  F1 score =  $\frac{2(\text{precision})(\text{recall})}{\text{precision}+\text{recall}}$

Balanced accuracy =  $0.5\left(\frac{TP}{TP+FN} + \frac{TN}{TN+FP}\right)$

### ❖ Cross-validation

➤ KNN

- Standard procedure
- After chosen distance metric and K,
  1. Normalization
  2. Compute distances
  3. Identify the K most similar data
  4. Take their class out and find the mode class
- Good K = good prediction accuracy
- Problem: no label for testing data

Solution: use part of training data as testing data  
(use each part one by one and calculate the average)

- Can use  $L_\infty$  for convenience

➤ Cross-fold validation

- Procedure to measure the performance of models
- N-fold cross-validation:

Step 1: randomly partition data into n disjoint subsets

Step 2: for  $i=1$  to n,

Validation data = i-th subset

$H \leftarrow$  classifier trained except validation data

Accuracy (i) = accuracy of h

Step3: final accuracy = mean of n recorded accuracies

❖ **Multi-class classification**

➤ Consider each class as binary classification problem

➤ Aggregate multiple values into one value:

➤ Macro-average =  $\frac{\text{sum of accuracy of each class}}{\text{number of classes}}$

➤ Micro-average =  $\frac{\text{sum of (accuracy*number of data of the class)}}{\text{number of data}}$

❖ **Clustering evaluation** (different from classification !!)

➤ Messy classification can be a good clustering

➤ Should be evaluate a pair of cells

➤ Rand index  $R = \frac{a+d}{a+b+c+d} = \frac{a+d}{\text{Number of all pair combinations}}$

➤ Number of pairs =  $\binom{n}{2} = \frac{n(n-1)}{2}$