

SVM and SVM Ensembles in Breast Cancer Prediction

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Breast Cancer Prevention Model

- Studied Techniques: logistic regression, linear discriminate analysis, artificial neural network and so on.
- Objective: compare performance of SVM and SVM ensembles over small and large scale breast cancer datasets.

Dataset

- Small scale:

699 data samples: 458 benign (65.5%) and 241 (34.5%) malignant;

11 different features: 1. sample code number;

2-10: 9 attributes range from 1 to 10;

i.e. clump Thickness, Uniformity of Cell, Marginal Adhesion;

11. class: (benign, malignant)

- Large scale:

102294 data samples;

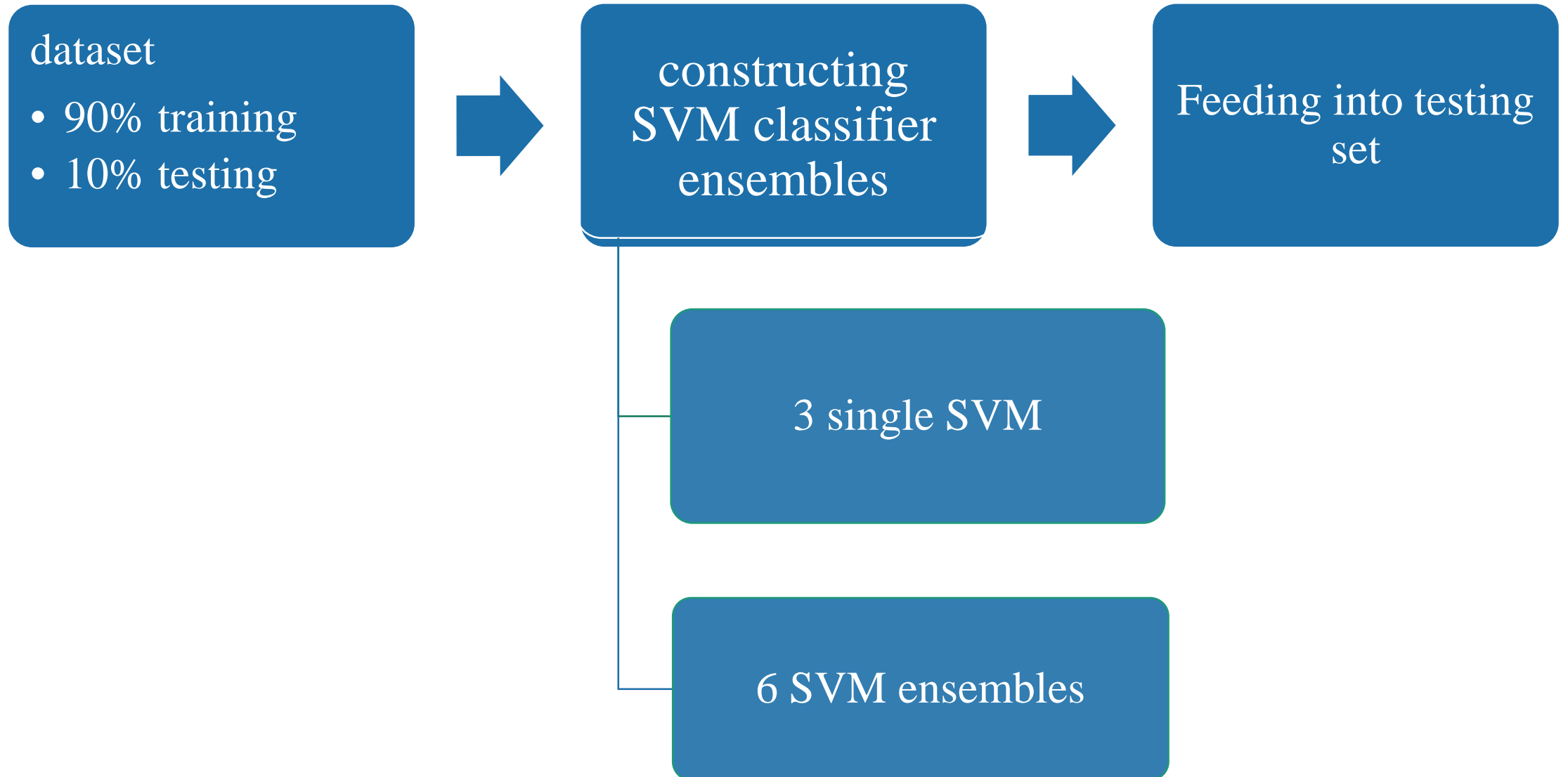
117 different features: detection of breast cancer from X-ray images of the breast

- feature selection(GA): to filter out unrepresentative features;

Small scale: 10 feature;

Large scale: 36 features

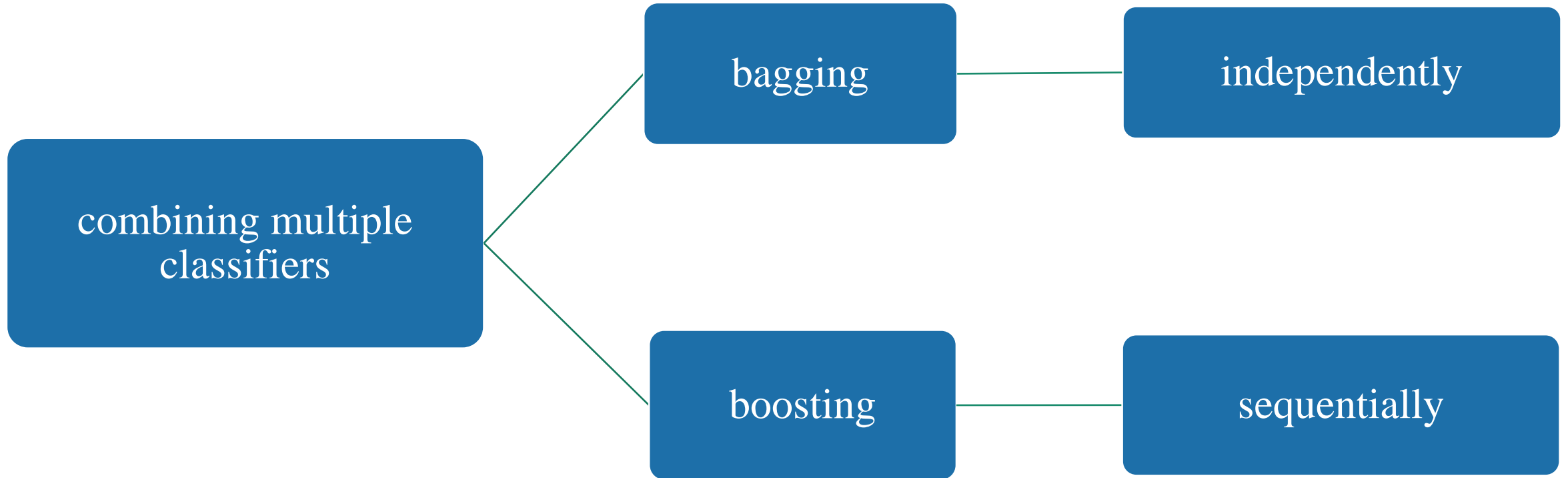
Experimental Procedure



Kernel Function

- Linear kernel function: $K(x_i, x_j) = \Phi(x_i) \cdot \Phi(x_j)$, $\Phi : \mathbb{R}^d \rightarrow \mathbb{H}^f$, $d < f$.
- RBF kernel function: $K_{Gaussian}(x_i, x_j) = e^{-\frac{\|x_i - x_j\|^2}{2\sigma}}$ (σ is Gaussian sigma)
- Polynomial kernel function: $K_{poly}(x_i, x_j) = (x_i \cdot x_j + 1)^p$ (p is the degree of polynomial)

Classifier Ensembles



Bagging

Bagging



training
set



⋮



bootstrap samples
(sampling with
replacement)



⋮



base classifiers



the ensemble

- Create Bootstrap samples of a training set using sampling with replacement
- Each bootstrap sample is used to train a different component of base classifier
- Classification is done by plurality voting

BAGGING

Training phase

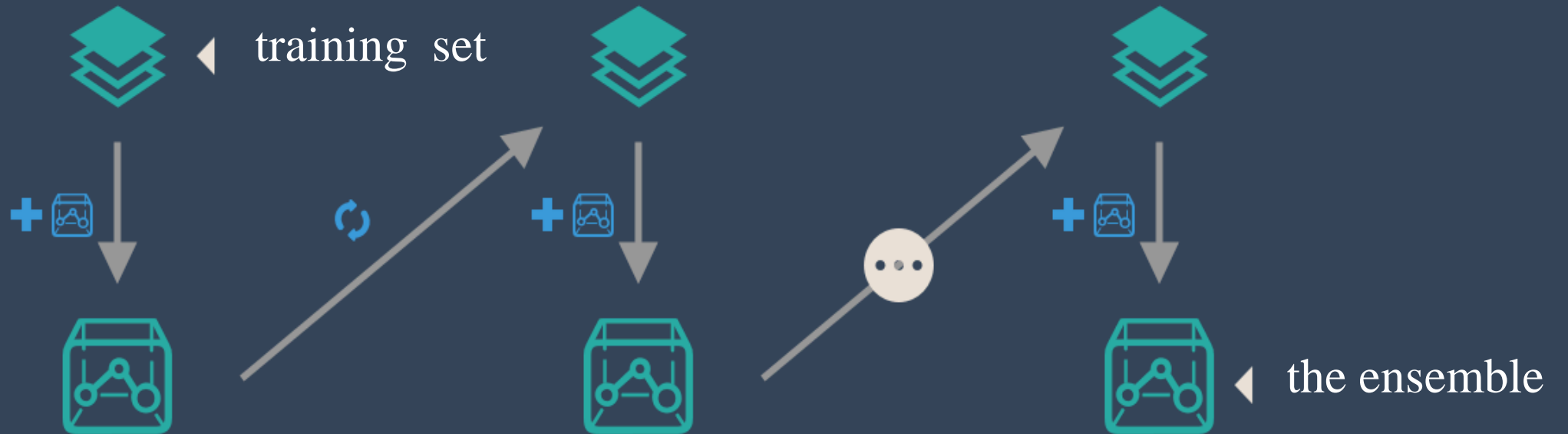
1. Initialize the parameters
 - $\mathcal{D} = \emptyset$, the ensemble.
 - L , the number of classifiers to train.
2. For $k = 1, \dots, L$
 - Take a bootstrap sample S_k from \mathbf{Z} .
 - Build a classifier D_k using S_k as the training set.
 - Add the classifier to the current ensemble, $\mathcal{D} = \mathcal{D} \cup D_k$.
3. Return \mathcal{D} .

Classification phase

4. Run D_1, \dots, D_L on the input \mathbf{x} .
5. The class with the maximum number of votes is chosen as the label for \mathbf{x} .

Boosting

Boosting



+ ensemble base classifier

update sample weights
based on the previous
classifier

- Sequential training of weak learners
 - Each base classifier is trained on data that is weighted based on the performance of the previous classifier
- Each classifier votes to obtain a final outcome

Training phase

1. Initialize the parameters
 - Set the weights $\mathbf{w}^1 = [w_1, \dots, w_N]$, $w_j^1 \in [0, 1]$, $\sum_{j=1}^N w_j^1 = 1$. (Usually $w_j^1 = \frac{1}{N}$).
 - Initialize the ensemble $\mathcal{D} = \emptyset$.
 - Pick L , the number of classifiers to train.

2. For $k = 1, \dots, L$

- Take a sample S_k from \mathbf{Z} using distribution \mathbf{w}^k .
- Build a classifier D_k using S_k as the training set.
- Calculate the weighted ensemble error at step k by

$$\epsilon_k = \sum_{j=1}^N w_j^k l_k^j,$$

($l_k^j = 1$ if D_k misclassifies \mathbf{z}_j and $l_k^j = 0$ otherwise.)

- If $\epsilon_k = 0$ or $\epsilon_k \geq 0.5$, ignore D_k , reinitialize the weights w_j^k to $\frac{1}{N}$ and continue.
- Else, calculate

$$\beta_k = \frac{\epsilon_k}{1 - \epsilon_k}, \quad \text{where } \epsilon_k \in (0, 0.5),$$

- Update the individual weights

$$w_j^{k+1} = \frac{w_j^k \beta_k^{(1-l_k^j)}}{\sum_{i=1}^N w_i^k \beta_k^{(1-l_k^i)}}, \quad j = 1, \dots, N.$$

3. Return \mathcal{D} and β_1, \dots, β_L .

Classification phase

4. Calculate the support for class ω_i by

$$\mu_i(\mathbf{x}) = \sum_{D_k(\mathbf{x})=\omega_i} \ln\left(\frac{1}{\beta_k}\right).$$

5. The class with the maximum support is chosen as the label for \mathbf{x} .

Comparison between Bagging and Boosting

- Sampling
 - Bagging: training set is chosen with replacement and train sets of each round are independent.
 - Boosting: train set does not change for each round.
- Weight
 - Bagging: evenly
 - Boosting: weight changes based on the loss function(error). Less the accuracy, larger the weight.
- classifier
 - Bagging: classifiers weight evenly
 - Boosting: every classifier has corresponding weight. Classifier which is more accurate has larger weight.
- computation
 - Bagging: classifier generated simultaneously
 - Boosting: classifier generated order by order because the parameters used in n-th model are produced by the (n-1) th model
- Variance
 - Bagging: variance decreases as number of model increases

Compare SVM and SVM ensembles

- Classification accuracy
- ROC (receiver operation characteristic)

		<u>True class</u>	
		p	n
<u>Hypothesized class</u>	Y	True Positives	False Positives
	N	False Negatives	True Negatives
Column totals:		P	N

fp rate = $\frac{FP}{N}$ tp rate = $\frac{TP}{P}$

precision = $\frac{TP}{TP+FP}$

accuracy = $\frac{TP+TN}{P+N}$

F-measure = $\frac{2}{1/precision+1/recall}$

- F-measure
- Computational times of training

Experimental Results

- The performance of single SVM classifiers vs SVM classifier ensembles

SVM ensembles > single SVM classifier

- The performance of applying genetic algorithm (GA)

Small scale: with GA > without GA

Comparison of the the classification accuracy, ROC, and F-measure of the top 3 classifiers

Classification accuracy		ROC		F-measure	
<i>Small scale dataset</i>					
1	GA+RBF SVM ensembles (boosting) (98.28%)	1	GA+linear/poly SVM ensembles (bagging) (0.98)	1	GA+RBF SVM (0.988)
2	GA+linear SVM (96.85%)	2	GA+poly SVM ensembles (boosting) (0.979)	2	GA+linear SVM ensembles (bagging/boosting) (0.966)
3	GA+linear SVM ensembles (bagging/boosting) (96.57%)	3	GA+RBF SVM ensembles (boosting) (0.977)	3	GA+RBF SVM ensembles (boosting) (0.963)
<i>Large scale dataset</i>					
1	RBF SVM ensembles (boosting) (99.52%)	1	GA+linear SVM ensembles (boosting) (0.876)	1	RBF SVM ensembles (boosting) (0.995)
2	Poly SVM ensembles (bagging) (99.51%)	2	GA+RBF SVM ensembles (boosting) (0.875)	2	Poly SVM; poly SVM ensembles (bagging); GA+poly SVM ensembles (bagging); GA+RBF SVM ensembles (boosting) (0.994)
3	Poly SVM; GA+poly SVM; RBF SVM ensembles (bagging); GA+poly SVM ensembles (bagging) (99.50%)	3	RBF SVM ensembles (boosting) (0.869)		

Reference

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