CS7641 Fall 2018 Assignment1 Supervised learning

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Datasets

Why they are interesting

Nontrival, allows tuning and comparing of different algorithms

	Task	#Instances	#Attributes				
HTRU2	Binary						
	Classification						
Breast	Binary	683 ^[1]	9 ^[2]	Real			
Cancer	Classification						
Wisconsin							

Table 1.

- [2] Removed 16 instances with missing data
- [3] ID number attribute was dropped

Scaling

Splitting into training (70%) and testing (30%)

Results and Discussion

1. Decision Tree

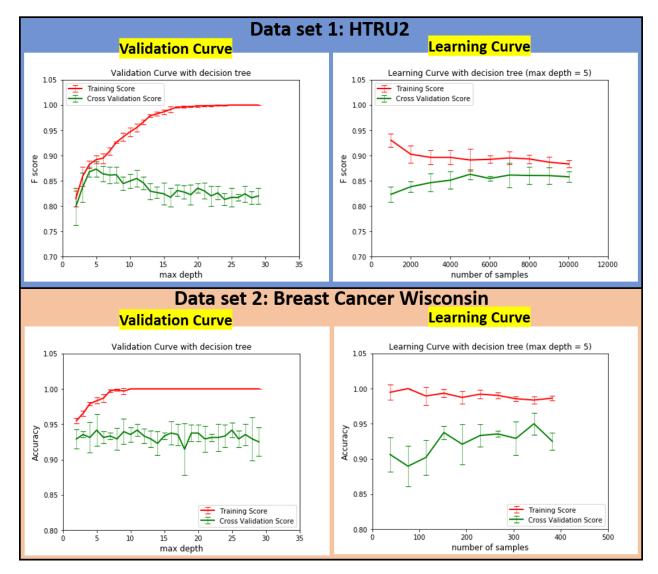


Figure 1. Decision Tree: Validation curve and learning curve (5 folds cross validation. Error bars represent standard deviation)

2. k-Nearest Neighbors

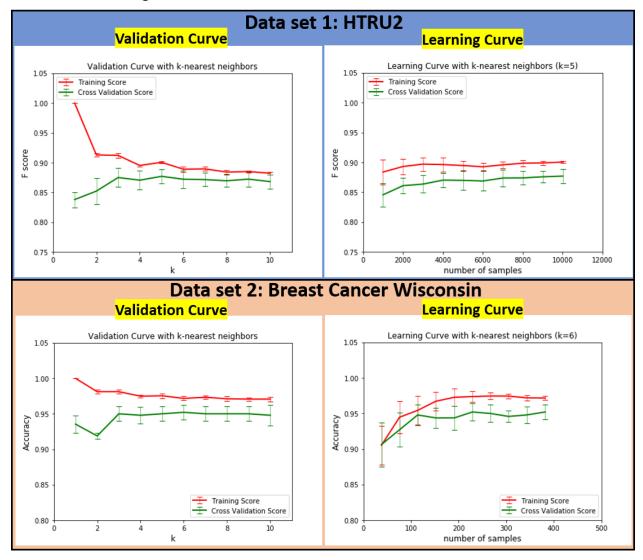


Figure 2. k-Nearest Neighbors: Validation curve and learning curve (5 folds cross validation. Error bars represent standard deviation)

The effects of different values of k
The HTRU2 dataset

3. Support Vector Machine

3.1 linear kernel

The HTRU2 dataset

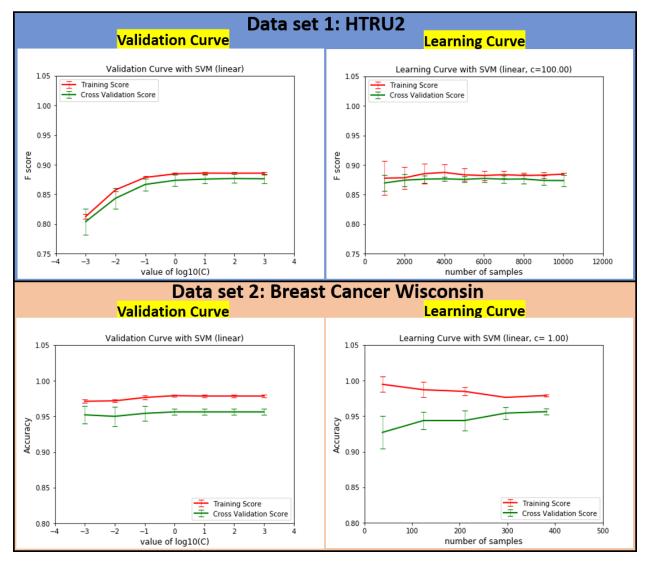


Figure 3.1. SVM with linear kernel: Validation curve and learning curve (5 folds cross validation. Error bars represent standard deviation)

3.2 RBF kernel

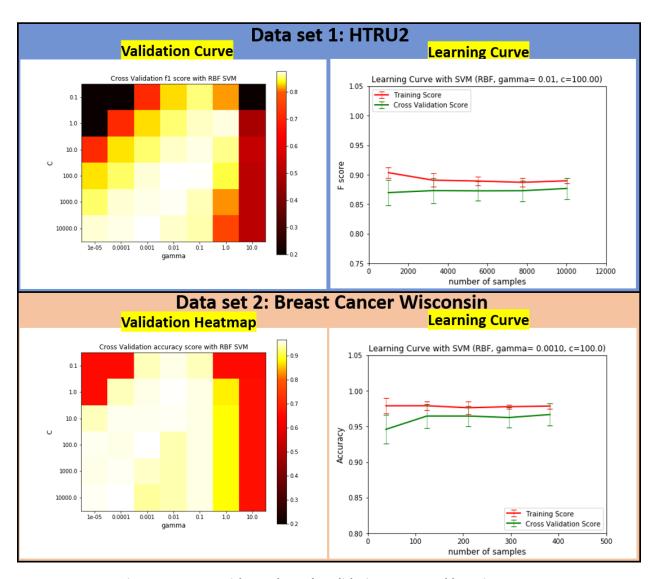


Figure 3.2. SVM with RBF kernel: Validation curve and learning curve (5 folds cross validation. Error bars represent standard deviation)

4. Boosting

Base classifier: decision tree with more aggressive pruning (max depth 3 for HTRU2 and 1 for breast cancer data set)

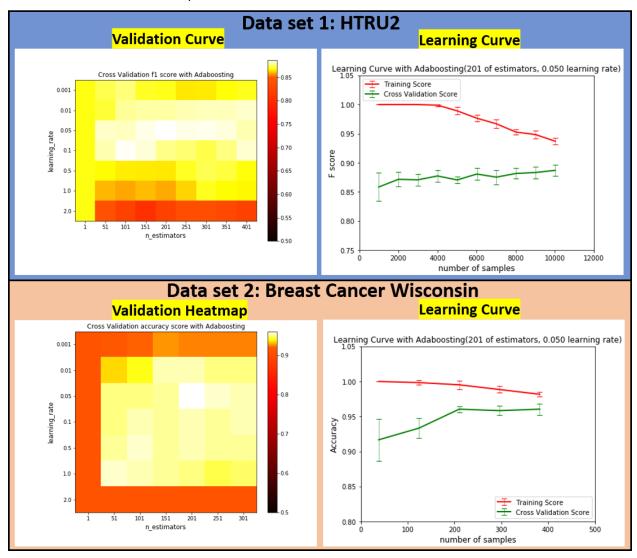


Figure 4. Adaboosting: Validation curve and learning curve

(5 folds cross validation. Error bars represent standard deviation)

5. Neural Networks

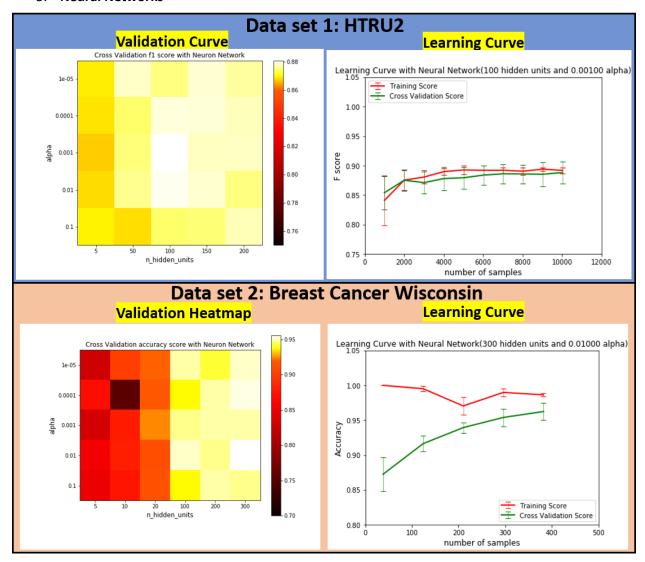


Figure 5. Neural Networks: Validation curve and learning curve (5 folds cross validation. Error bars represent standard deviation)

1 hidden layer # of hidden unites the value of alpha, activation function

6. Compare and contrast the different algorithms.

Learning	Test	Test	Test	Training	Test Time	Hyperparameters			
Algorithms	AUC Score	Accuracy	F Score	Time					
HTRU2 Data Set									
Decision Tree	0.89898	0.97821	0.85921	0.0132626	0.00346114	Max depth = 5			
k-Nearest	0.90994	0.97952	0.86967	0.0091312	0.126501	k=5			
Neighbor									
SVM	0.90952	0.98063	0.87560	3.70600	0.0311281	C=100			
(linear kernel)									
SVM	0.91270	0.98082	0.87753	0.454213	0.0916295	C=100			
(RBF kernel)						gamma=0.01			
Boosting	0.91762	0.98045	0.87691	6.54262	0.102999	Decision tree with			
						max_depth = 3			
						200 estimators			
						Learning rate 0.05			
Neural	0.91987	0.98082	0.87953	4.57695	0.00571682	alpha=0.001			
Networks						1 hidden layer			
						with 100 nodes			
		Breast (Cancer Wisc	onsin Data Se					
Decision Tree	0.94077	0.94634	0.92617	0.0006914	0.00035675	Max depth = 5			
k-Nearest	0.95128	0.95610	0.93960	0.0007125	0.00180639	k=6			
Neighbor									
SVM	0.94744	0.95122	0.93333	0.0027717	0.00044157	C=1			
(linear kernel)									
SVM	0.96744	0.96585	0.95425	0.0023225	0.00058448	C=100			
(RBF kernel)						gamma=0.001			
Boosting	0.93795	0.94634	0.92517	0.309154	0.0149498	Decision tree with			
						max_depth = 1			
						200 estimators			
						Learning rate 0.05			
Neural	0.93795	0.94634	0.92517	1.75296	0.00088109	alpha=0.01			
Networks						1 hidden layer			
						with 300 nodes			

Table 2. Comparison of different learning algorithms

Which algorithm performed best? How do you define best?

What sort of changes might you make to each of those algorithms to improve performance? How fast were they in terms of wall clock time? Iterations?

How much performance was due to the problems you chose? How about the values you chose for learning rates, stopping criteria, pruning methods, and so forth (and why doesn't your analysis show results for the different values you chose?)?

Features are computed from a digitized image of a fine needle aspirate (FNA) of a breast mass. They describe characteristics of the cell nuclei present in the image.

Reference

W.N. Street, W.H. Wolberg and O.L. Mangasarian. Nuclear feature extraction for breast tumor diagnosis. IS&T/SPIE 1993 International Symposium on Electronic Imaging: Science and Technology, volume 1905, pages 861-870, San Jose, CA, 1993. [Web Link]

O.L. Mangasarian, W.N. Street and W.H. Wolberg. Breast cancer diagnosis and prognosis via linear programming. Operations Research, 43(4), pages 570-577, July-August 1995. [Web Link]