Predicting Breast Cancer Using Machine Learning

Classification Project
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Source: National Breast Cancer Foundation

breast cancer in her lifetime.









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Objective

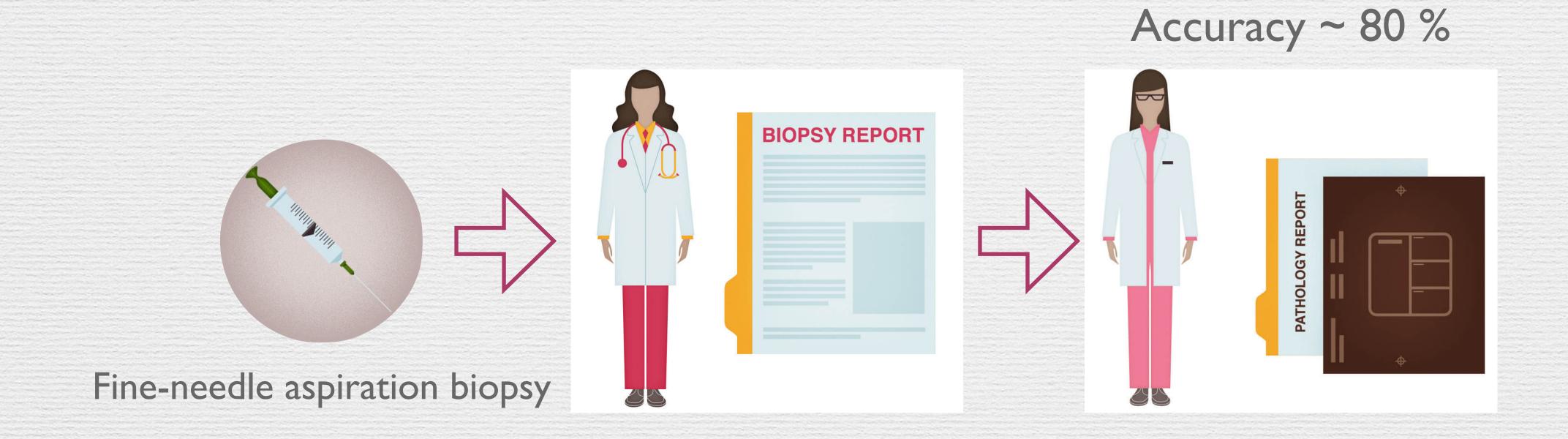




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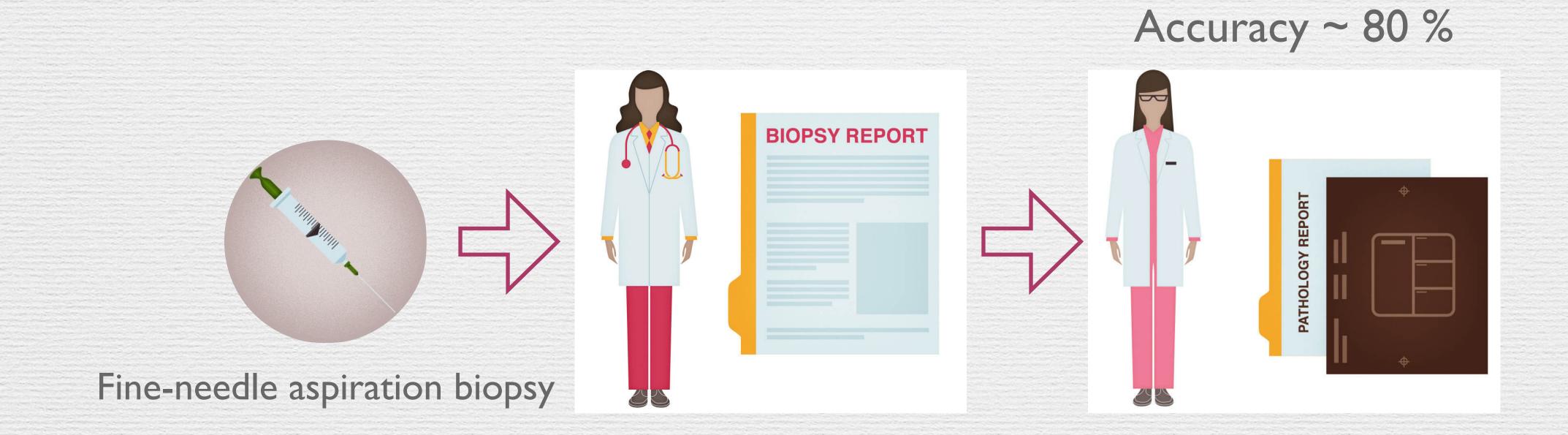
1~2 weeks



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Improve diagnosis accuracy and speed with machine learning





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Models

+ 9 classification models in SciKit-Learn



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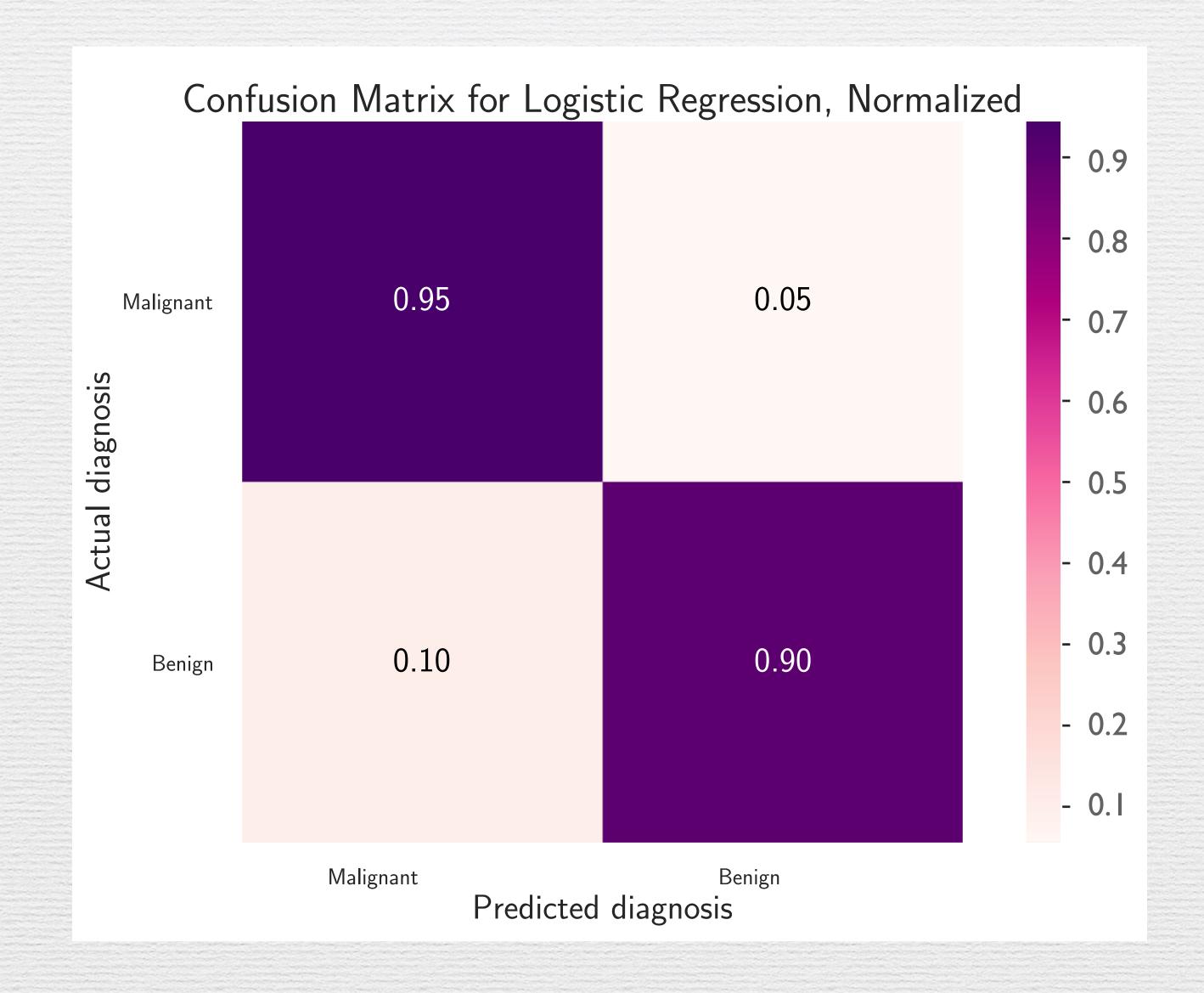
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- + 9 classification models in SciKit-Learn
 - + Top 4: Logistic Regression, Gradient Boosting, Random Forest, SVM

Results









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Future Work

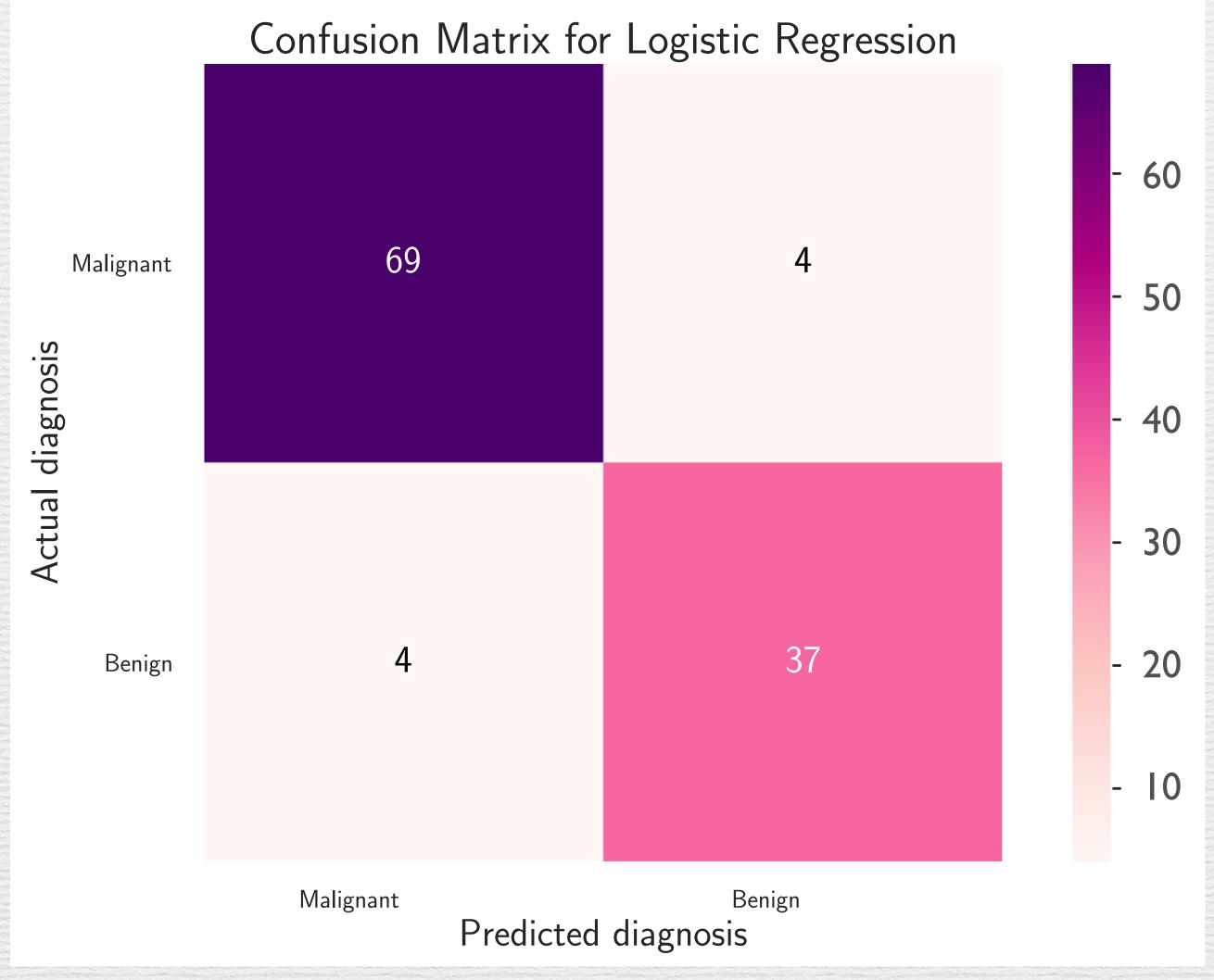


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Future Work

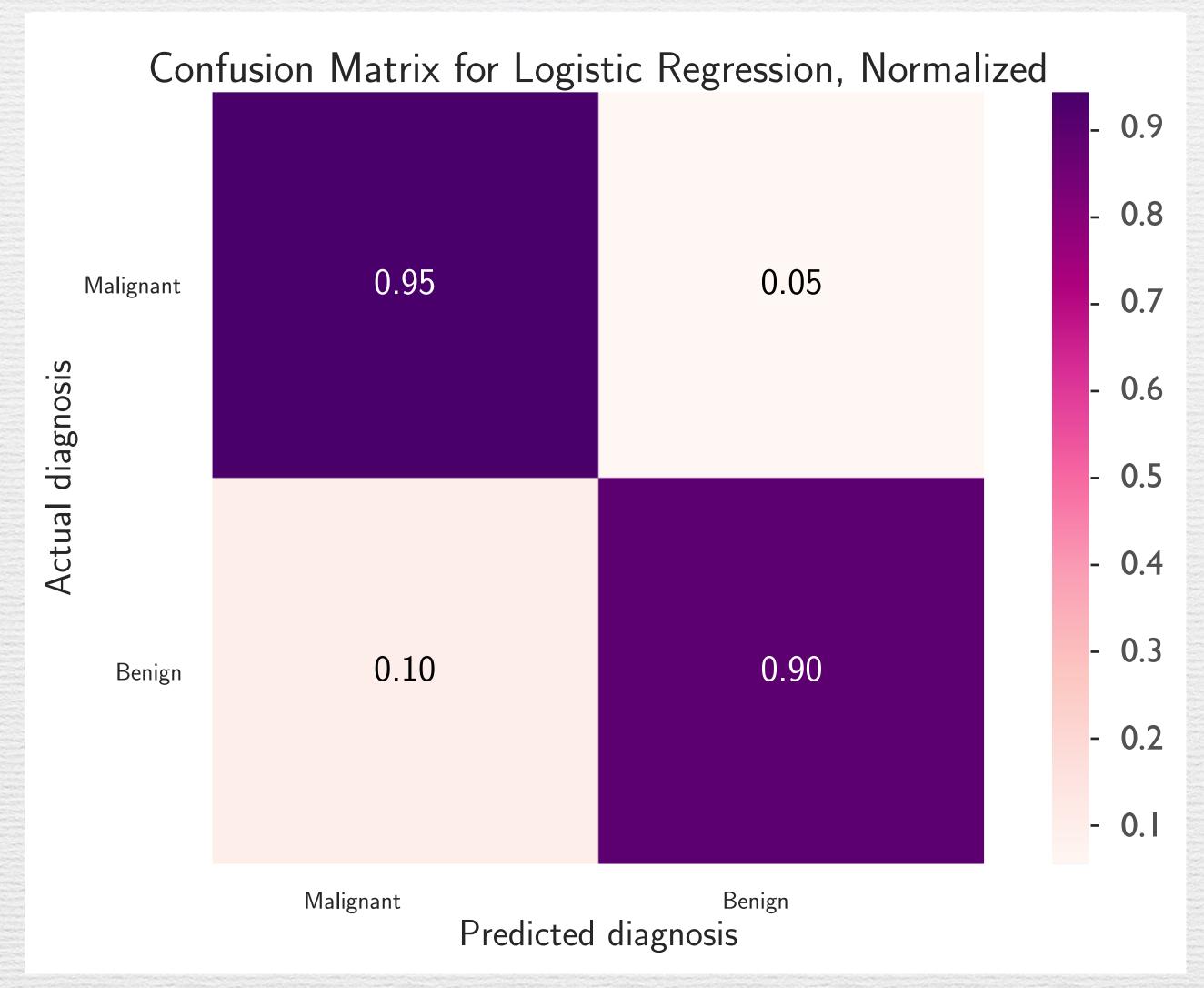
+ Image classification — "biopsy to diagnosis"





Test set, cross-validation over 5 folds



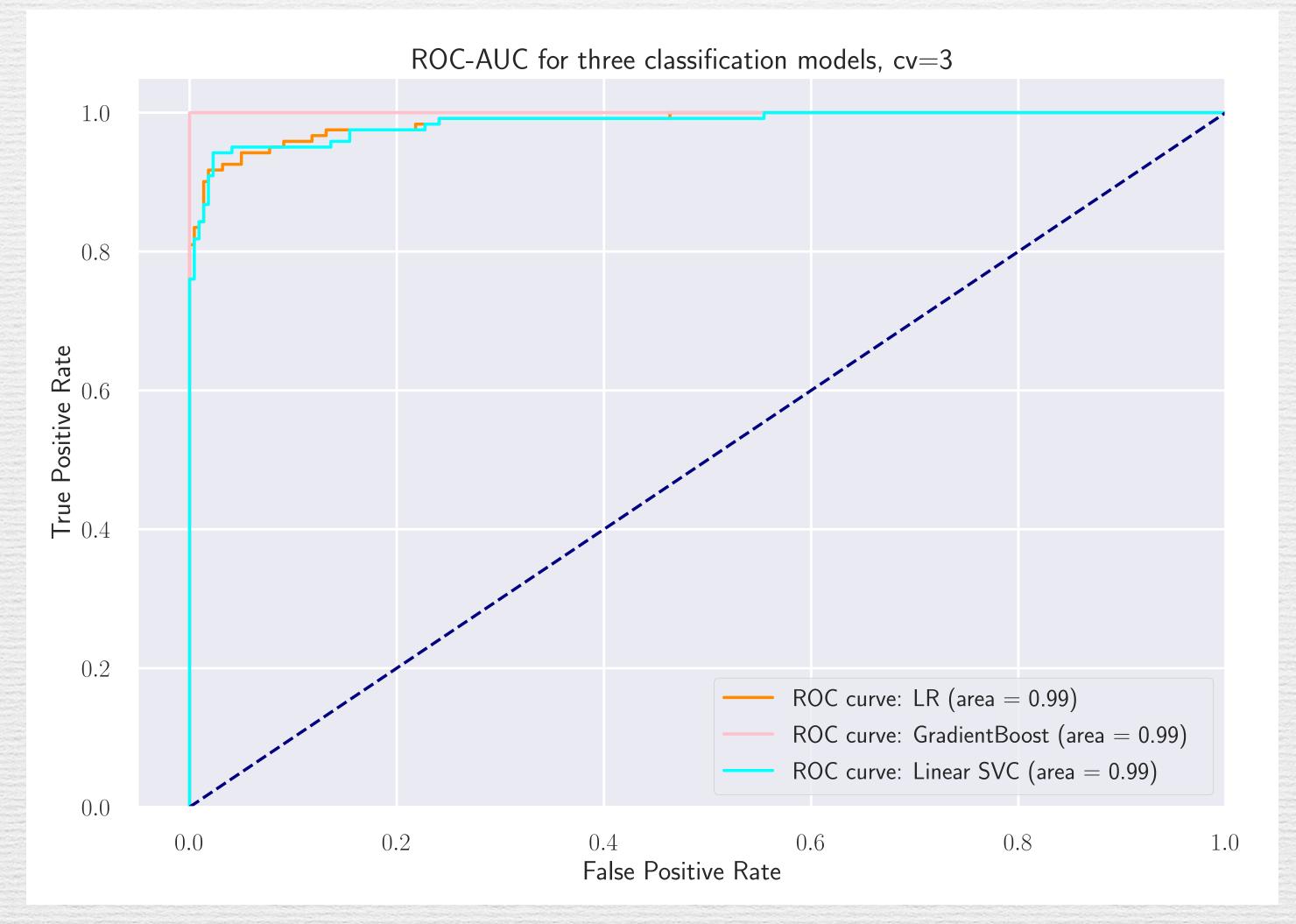


Normalization: divide prediction counts by the sum of each row



Using 10 features

LR AUC score validation set 0.9884

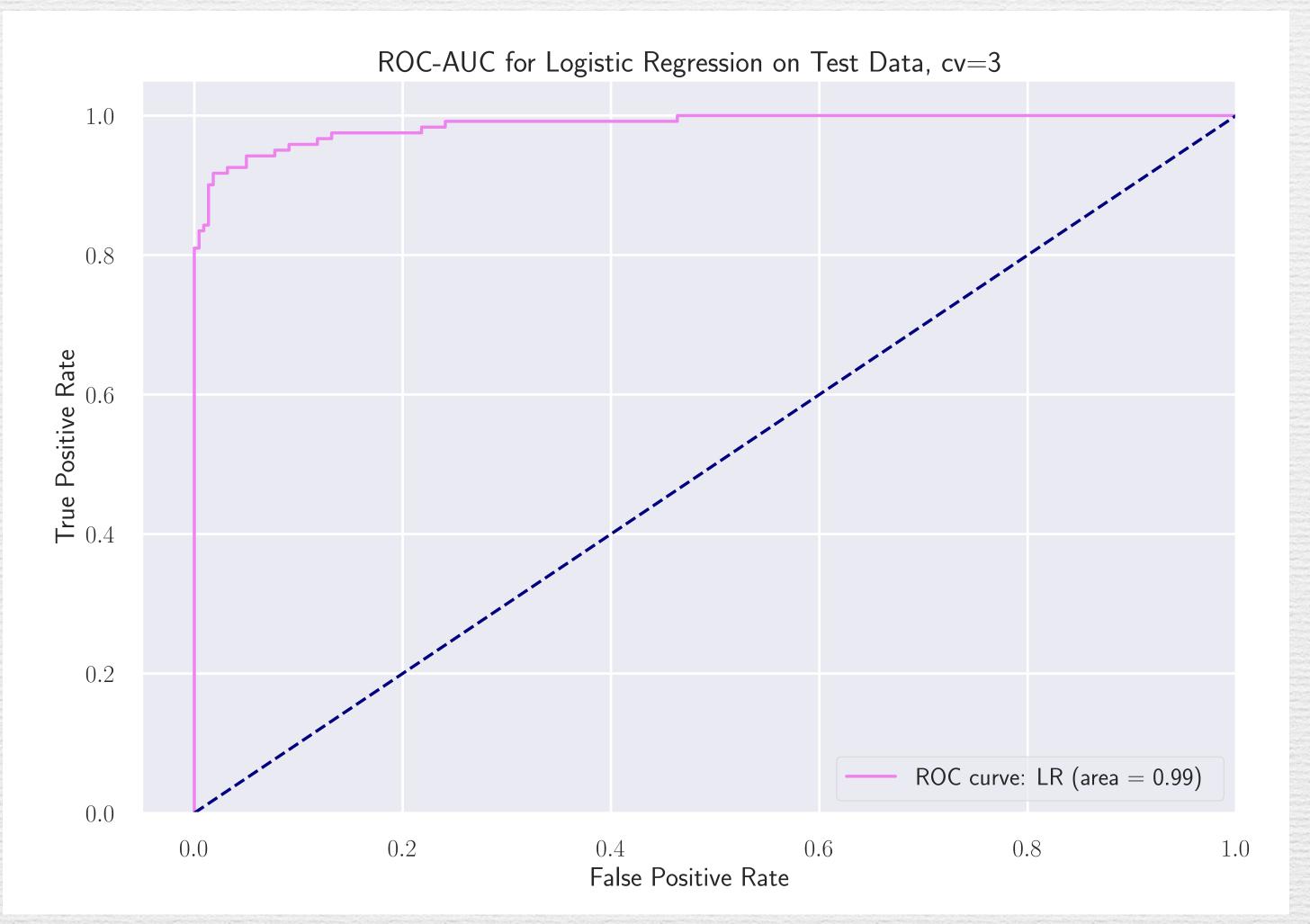


Comparison of the three models with highest roc-auc scores for validation set, with naïve model at 0.5 (dashed line)



After feature engineering: Validation set 0.9909

Test set 0.9870



Area under the curve for Logistic Regression on test data set after feature engineering, with naïve model at 0.5 (dashed line)



Model	Metric	Top features	Weights (not normalized, 4 decimals)
LogisticRegression	coefficient	area	3.8253
		concave points	2.7111
		texture	1.6457
SVM - linear SVC		area	2.4970
	coefficient	concave points	1.2092
		texture	1.0927
RandomForest	feature	concave points	0.4975
	importance	area	0.3071
		compactness	0.1300

Table showing the most impactful features for predicting diagnosis, for the models with the top roc-auc scores, after feature engineering. Area and concave points are the two most important features for our prediction.







Multicollinearity

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AUC score for LR on validation set (10 features): 0.9884 LR on test set (after feature engineering, 5 features): 0.9870

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