CS205 Spring 2019 Project 2 Report: Cancer prediction using NHANES Dataset

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Abstract

Cancer as a whole is one of the most deadly diseases in the world, second only to cardiovascular disease. Every cancer is different, and there is much study into 3 predicting and treating all forms of cancer. However, there is little research on predicting cancer as a whole. In the course of this research, I analyzed and used the NHANES dataset to predict cancer based on a variety of features and using a 5 variety of strategies.

Introduction

- The goal of this project is to predict whether someone has or is at rick for any kind of cancer. I
- selected and manipulated data and programmed statistical models to make these predictions. I consulted many resources online, including research papers, technical blog posts, and coding forums.
- Environment: I set up my notebook on an AWS Sagemaker server for ease of remote access and 11
- consistent computation. 12
- Specs: 13

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- ml.t2.medium instance
- 2 virtual CPUs 15
- 4 GIB RAM 16
 - Python 3.6
- Pytorch 1.0 18

Methodology

2.1 Data Source

- The data comes from NHANES, or the National Health and Nutrition Examination Survey. NHANES 21
- "is a program of studies designed to assess the health and nutritional status of adults and children in 22
- the United States. The survey is unique in that it combines interviews and physical examinations." In 23
- this study I focused on using data from the 2015-2016 data collection, but also included data from 24
- 25 older surveys and records if they contained the same features. The NHANES py script labels 0 as has
- cancer, and 1 as no cancer (healthy).

2.2 Baseline 27

- Sample code was given for predicting arthritis more specifically predicting the target "MCQ160 -
- Doctor ever said you had arthritis." It used about 30 features.

The test accuracies using various classification methods are as follows. This served as the baseline performance to compare my new model against.

32		Random Support Logistic Re	Vector	0.756 0.758 0.767	
	accu_tst_R	FC 0.714			
		precision	recall	f1-score	support
		0.70	0.76	0.73	500
		0.74	0.67	0.70	500
	avg / tota	1 0.72	0.71	0.71	1000
33	[[381 119] [167 333]]			
34		Figure: 1	Base RFC	stats	
	accu_tst_S	VC 0.711			
		precision	recall	f1-score	support
		0 0.68	0.78	0.73	500
		1 0.75	0.64	0.69	500
	avg / tota	al 0.72	0.71	0.71	1000
35	[[392 108] [181 319]				
36		Figure: I	Base SVC	stats	
	accu_tst_L	R 0.702			
		precision	recall	f1-score	support
		0 0.67	0.78	0.72	500
		0.74	0.62	0.68	500
	avg / tota	0.71	0.70	0.70	1000
37	[[390 110] [188 312]]			
··					

Figure: Base LR stats

39 2.3 Data Preparation

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I selectively loaded data by extracting the desired features via their feature code. This allowed me to flexibly change the features I used.

2.4 Feature Selection

3 2.4.1 Exhaustive search

The first method I attempted to select features was an exhaustive search to evaluate correlation. I initially hoped to automate the entire process of extracting features from the data set, applying preprocessing, and calculating mutual information. However, automating preprocessing proved unsuccessful due to the intricate nature of selecting the correct preprocessing procedure for each type of data. Different data such as binary, categorical, continuous, and sparse each need unique preprocessing procedures. I also wish to note that a classmate who did attempt to run an exhaustive mutual information analysis stated that all the mutual information he extracted were too low and similar to use. Thus I switched to manual feature selection.

2.4.2 Manual Selection

Because of the difficulty of exhaustive search and limitation of time, I tried instead to select many features by hand via intuition. This allowed me to appropriately curate preprocessing methods for

each feature. The downside was that, due the multitude of features, it was impossible to analyze and preprocess every feature by hand, as that alone would have cost weeks. Thus I used my intuition, prior knowledge, and research, to manually select features I assumed to be the most predictive of cancer. Because most research focuses on specific types of cancer, I browsed much research of more specific cancer research, thus treating my project as a sort of ensemble research. I cross referenced the available NHANES features with existing research papers to find features that existed in prior research.

- Vaginal swab testing ¹
- Blood features ^{2 3}
 - BMI and body shape/composition ^{4 5 6}
- Age ⁷

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- Cultural background, socioeconomic status, and education ^{8 9}
- Alcohol and drugs ¹⁰ ¹¹

68 2.5 Predictive Modeling - Standard

The model itself is arguably the most important part of the predictive modeling process. There are many models so I researched some of the most popular and well-performing models used today for our type of data, especially in the application of medical analysis. I also considered factors such as complexity and efficiency. 12 13 14

- Random Forest Classifier ¹⁵ This is an ensemble learning method. It runs many different simpler decision tree models and uses bagging and boosting to compile their predictive power together to form a super-model. Ensemble methods can be very powerful because they are not confined to a single scope of the data.
- Support Vector Classifier ¹⁶ The SVM is a straightforward but still-popular model.
- Logistic Regression a statistical model that computes probabilities for each output class.
- Linear Discriminant Analysis (LDA) "attempts to express one dependent variable as a linear combination of other features or measurements." This is a commonly used model.

1 2.6 Data Size

I ran into an interesting observation after including more data (from the baseline) and increasing the train and test set sizes. I increased the training set size from 5000 to 20000, and testing set from 2000 to 10000. After doing so, the precision and recall for each classifier became heavily skewed toward the positive class and achieved much lower precision and recall for the negative class while increasing for the positive class. The classifiers are seemingly weighted too. It can be that the

¹https://jamanetwork.com/journals/jama/article-abstract/192260

²https://www.sciencedirect.com/science/article/pii/S0140673696034307

³https://www.sciencedirect.com/science/article/abs/pii/S0168827806002972

⁴https://www.sciencedirect.com/science/article/pii/S014067360860269X

⁵https://www.nature.com/articles/0802606

⁶https://academic.oup.com/ajcn/article/80/4/1012/4690349

⁷https://www.nature.com/articles/ng0299_163

⁸https://onlinelibrary.wiley.com/doi/full/10.3322/canjclin.54.2.78

⁹https://onlinelibrary.wiley.com/doi/full/10.3322/canjclin.56.3.168

¹⁰https://www.sciencedirect.com/science/article/abs/pii/S1470204506705770

¹¹https://onlinelibrary.wiley.com/doi/full/10.1002/cncr.27554

¹²https://www.sciencedirect.com/science/article/pii/S1532046401910044

¹³https://www.kaggle.com/jeffd23/10-classifier-showdown-in-scikit-learn

¹⁴https://towardsdatascience.com/which-machine-learning-model-to-use-db5fdf37f3dd

¹⁵https://www.researchgate.net/profile/Andy_Liaw/publication/228451484_Classification_and_Regression_by_RandomForest/links/53fb24cand-Regression-by-RandomForest.pdf

¹⁷https://en.wikipedia.org/wiki/Linear_discriminant_analysis

importance of rejecting False Negatives (False cancer) starts outweighing the importance of detecting True Positives(True has-cancer). It seems that a tradeoff must be made between the two. Both have their pros and cons. With less data in the model, there is more of a chance someone with cancer being detected as such, but many non-cancer patients will be falsely diagnosed. With more data, fewer patients will be falsely told they have cancer, but fewer of those who have or are at risk for cancer will be told. For the rest of this project, I will use the larger dataset that is more representative of all the data. Because this model so far is not highly accurate and should only be used as a supplement to an expert's judgement, I would personally shy away from sowing worry into a healthy patient's mind with a wrong prediction, as this itself can be dramatically detrimental to their health and even contribute to unnecessary CAT scans.

Random Forest	Classifier	0.6897572	528123149	
	precision			support
	p			
0	0.59	0.76	0.67	689
1	0.79	0.64	0.71	1000
micro avg	0.69	0.69	0.69	1689
macro avg	0.69	0.70	0.69	1689
weighted avg	0.71	0.69	0.69	1689
[[521 168]				
[356 644]]				
Support Vector				
	precision	recall	f1-score	support
0	0.59	0.79	0.68	689
1	0.81	0.62	0.70	1000
micro avg	0.69	0.69	0.69	1689
macro avg	0.70	0.71	0.69	1689
weighted avg	0.72	0.69	0.69	1689
[[544 145] [377 623]]				
Logistic Regr	ession 0.69	6269982238	30106	
		recall		support
0	0.60	0.77	0.68	689
1	0.80	0.64	0.71	1000
micro avg	0.70	0.70	0.70	1689
macro avg	0.70	0.71	0.69	1689
weighted avg	0.72	0.70	0.70	1689
[[533 156]				
[357 643]]				
Linear Discri		-		
	precision	recall	f1-score	support
0	0.60	0.79	0.68	689
1	0.81	0.79	0.72	1000
1	0.01	0.04	0.72	1000
micro avg	0.70	0.70	0.70	1689
macro avg	0.71	0.71	0.70	1689
weighted avg	0.73	0.70	0.70	1689
	0.75	2.70	2.70	2000
[[541 148]				
[359 641]]				
Figure: Cla	ssifiers wi	th 5000/2	000 train/	test solit

Figure: Classifiers with 5000/2000 train/test split

Random Forest	Classifier	0.8582510	578279267	
	precision	recall	f1-score	support
0	0.36	0.25	0.30	672
1	0.90	0.94	0.92	5000
micro avg	0.86	0.86	0.86	5672
macro avg	0.63	0.60	0.61	5672
weighted avg	0.84	0.86	0.85	5672
[[171 501] [303 4697]]				
Support Vector	r Classifie	r 0.669428	7729196051	
	precision	recall	f1-score	support
	-			
0	0.23	0.74	0.35	672
1	0.95	0.66	0.78	5000
_				
micro avg	0.67	0.67	0.67	5672
macro avg	0.59	0.70	0.56	5672
weighted avg	0.86	0.67	0.73	5672
0 0				
[[500 172] [1703 3297]]				
Logistic Regre	ession 0 67	7186177715	0017	
LOGISCIC KEBI	precision		f1-score	
	precision	recall	11-50016	support
0	0.23	0.74	0.35	672
1	0.95	0.67	0.79	5000
micro avg	0.68	0.68	0.68	5672
macro avg	0.59	0.70	0.57	5672
weighted avg	0.87	0.68	0.73	5672
[[498 174] [1657 3343]]				
Linear Discri	minant Analy	ysis 0.792	8420310296	191
	precision	recall	f1-score	support
0	0.29	0.50	0.37	672
1	0.93	0.83	0.88	5000
1	0.93	0.03	0.30	2000
micro avg	0.79	0.79	0.79	5672
macro avg	0.61	0.67	0.62	5672
weighted avg	0.85	0.79	0.82	5672
-0				
[[338 334] [841 4159]]				

Figure: Classifiers with 20000/10000 train/test split

2.7 Preprocessing and Feature Engineering

2.7.1 Principal Component Analysis

Principal component analysis (PCA) is a popular strategy for data analysis and decomposition. It reduces data complexity by reducing the data to fewer (important) dimensions. ¹⁸ I tested decomposing the data into different numbers of components. For Random Forest and LDA, the testing shows that the more components that are decomposed, the more the models skew toward predicting the major class, improving the weighted average at the cost of accuracy of the minor class. Beyond 10 components, the results do not change. It is difficult to definitively say whether this is better or worse, but should be assessed more before real use since the predictions are so skewed to one side. For SVC and LR PCA seems to simply reduce performance. test

¹⁸https://www.cs.cmu.edu/ elaw/papers/pca.pdf

Random Forest C	lassifier 0	0.6969430	780042164						
	recision		f1-score	support	Random Forest	Classifier	0.8134619	384615385	
P						precision	recall	f1-score	suppor
0	0.14	0.29	0.19	692					
1	0.88	0.75	0.81	5000	0	0.14	0.09	0.11	72
-	0.00	0175	0.01	3000	1	0.88	0.92	0.90	500
micro avg	0.70	0.70	0.70	5692					
macro avg	0.70	0.70	0.50	5692	micro avg	0.81	0.81	0.81	572
weighted avg	0.79	0.70	0.74	5692	macro avg	0.51	0.51	0.50	572
weighted avg	0.79	0.70	0.74	5092	weighted avg	0.78	0.81	0.80	572
[001 000 1]					0				
[[202 490]					[[67 653]				
[1235 3765]]	c1 .c.				[414 4586]]				
Support Vector					Support Vector	Classifie	r 0.564160	8391608391	
р	recision	recall	f1-score	support		precision		f1-score	suppor
									-appo.
0	0.18	0.64	0.28	692	0	0.13	0.42	0.20	72
1	0.92	0.59	0.72	5000	1	0.88	0.58	0.70	500
					*	0.00	0.50	0.70	300
micro avg	0.60	0.60	0.60	5692	micro avg	0.56	0.56	0.56	572
macro avg	0.55	0.62	0.50	5692	macro avg	0.50	0.50	0.45	572
weighted avg	0.83	0.60	0.67	5692	weighted avg	0.78	0.56	0.43	572
					weighted avg	0.70	0.50	0.64	5/2
[[441 251]					[[202 417]				
[2032 2968]]					[[303 417]				
Logistic Regres	sion 0.6126	141953619	9114		[2076 2924]]			2522	
р	recision	recall	f1-score	support	Logistic Regre				
						precision	recall	f1-score	suppor
0	0.18	0.61	0.28	692					
1	0.92	0.61	0.74	5000	0	0.12	0.46	0.19	72
					1	0.87	0.52	0.65	500
micro avg	0.61	0.61	0.61	5692					
macro avg	0.55	0.61	0.51	5692	micro avg	0.51	0.51	0.51	5720
weighted avg	0.83	0.61	0.68	5692	macro avg	0.49	0.49	0.42	5720
					weighted avg	0.77	0.51	0.59	5720
[[421 271]									
[1934 3066]]					[[332 388]				
Linear Discrimi	nant Analys	is 0.869	1145467322	557	[2422 2578]]	_			
	recision		f1-score	support	Linear Discrim				
P				-apport		precision	recall	f1-score	support
0	0.14	0.01	0.03	692					
1	0.14	0.99	0.93	5000	0	0.00	0.00	0.00	726
1	0.00	0.55	0.55	5000	1	0.87	1.00	0.93	5000
micro avg	0.87	0.87	0.87	5692					
	0.51	0.50	0.48	5692	micro avg	0.87	0.87	0.87	5720
macro avg					macro avg	0.44	0.50	0.47	5720
weighted avg	0.79	0.87	0.82	5692	weighted avg	0.76	0.87	0.82	572
[[10 682]					[[0 720]				
[63 4937]]					[0 5000]]				

Figure 1: PCA 1 Component Performance

Figure 2: PCA 2 Components Performance

1 2.7.2 Singular Value Decomposition

- SVD is a similar data decomposition method to PCA, and causes the models to behave similarly to
- 113 PCA with the same numbers of components.
- 114 test

115 2.8 Predictive Modeling - Neural Network

- I implemented a deep neural network using Pytorch 1.0 in python 3.6. This was my primary goal and point of analysis in this project.
- After researching and experimenting ¹⁹ ²⁰ I chose the following layers and hidden neurons for my network.

```
self.fc1 = nn.Linear(75, 120)
self.fc2 = nn.Linear(120, 400)
self.fc3 = nn.Linear(400, 800)
self.fc4 = nn.Linear(800, 300)
```

¹⁹https://www.researchgate.net/publication/258393467_Review_on_Methods_to_Fix_Number_of_Hidden_Neurons_in_Neural_Networks ²⁰https://towardsdatascience.com/beginners-ask-how-many-hidden-layers-neurons-to-use-in-artificial-networks-51466afa0d3e

Random Forest Cl	lassifier 0.8	522270	742358079						
pr	ecision r	ecall	f1-score	support	Deader Secret	c1161	0 0770000	76744406	
•					Random Forest	precision		76744186 f1-score	
0	0.11	0.02	0.04	725		precision	recall	TI-score	support
1	0.87	0.97	0.92	5000	0	0.05	0.00	0.00	676
					1	0.88	1.00	0.93	5000
micro avg	0.85	0.85	0.85	5725	*	0.00	1.00	0.55	5000
macro avg	0.49	0.50	0.48	5725	micro avg	0.88	0.88	0.88	5676
weighted avg	0.78	0.85	0.81	5725	macro avg	0.47	0.50	0.47	5676
					weighted avg	0.78	0.88	0.82	5676
[[18 707]					g g				
[139 4861]]					[[1 675]				
Support Vector C	lassifier 0.	514585	152838428		[18 4982]]				
pr	recision r	ecall	f1-score	support	Support Vector	Classifier	0.578576	4622973926	
						precision	recall	f1-score	support
0	0.13	0.49	0.20	725					
1	0.88	0.52	0.65	5000	0	0.13	0.44	0.20	676
					1	0.89	0.60	0.71	5000
micro avg	0.51	0.51	0.51	5725					
macro avg	0.50	0.51	0.43	5725	micro avg	0.58	0.58	0.58	5676
weighted avg	0.78	0.51	0.59	5725	macro avg	0.51	0.52	0.46	5676
					weighted avg	0.80	0.58	0.65	5676
[[357 368]					[[297 379]				
[2411 2589]]					[2013 2987]]				
Logistic Regress					Logistic Regre	ssion 0 471	458773784	3552	
pr	recision r	ecall	f1-score	support		precision		f1-score	support
0	0.12	0.48	0.20	725					
1	0.12	0.48	0.65	725 5000	0	0.12	0.56	0.20	676
1	0.07	0.51	0.05	5000	1	0.89	0.46	0.60	5000
micro avg	0.51	0.51	0.51	5725					
macro avg	0.50	0.50	0.42	5725	micro avg	0.47	0.47	0.47	5676
weighted avg	0.78	0.51	0.59	5725	macro avg	0.50	0.51	0.40	5676
weighted avg	0.76	0.51	0.55	3723	weighted avg	0.79	0.47	0.56	5676
[[346 379]									
[2430 2570]]					[[379 297]				
Linear Discrimin	ant Analysis	0.873	36244541484	172	[2703 2297]]				
			f1-score	support	Linear Discrim				
						precision	recall	f1-score	support
0	0.00	0.00	0.00	725	0	0.00	0.00	0.00	676
1	0.87	1.00	0.93	5000	1	0.88	0.00 1.00	0.94	676 5000
					1	0.00	1.00	0.94	5000
micro avg	0.87	0.87	0.87	5725	micro avg	0.88	0.88	0.88	5676
macro avg	0.44	0.50	0.47	5725	macro avg	0.44	0.50	0.47	5676
weighted avg	0.76	0.87	0.81	5725	weighted avg	0.78	0.88	0.83	5676
							2.30		2270
[[0 725]					[[0 676]				
[0 5000]]					[0 5000]]				

Figure 3: PCA 3 Components Performance

Figure 4: PCA 10 Components Performance

```
self.fc5 = nn.Linear(300, 2)
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```

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I used a Relu activation function between each layer and softmax (equivalent to sigmoid with this binary classification) after the final hidden layer. Most modern implementations I found utilized Relu to avoid the vanishing gradient problem. I tested multiple loss functions with multiple optimization algorithms.

I used 3 optimization algorithms - Stochastic Gradient Descent with Momentum, Adagrad, and 129 Adam. 21

- SGD Vanilla SGD is the "traditional" method, but is not very effective compared to more modern methods and is rarely used anymore. Only the learning rate is tunable so it can suffer from either slow learning or overstepping the minimum. A common augmentation is to add momentum (exponential smoothing) so that the loss can converge faster in the beginning and slow down to approach the minimum.
- Adagrad Adagrad is an algorithm that adaptively changes its learning rate automatically. I interpreted it as a smarter and more robust version of SGD with momentum. Footnote ³ explains that this makes it well suited for sparser data which could suit our dataset.

²¹https://arxiv.org/pdf/1609.04747.pdf

	lassifier	0 7000007	421727601						
	recision		f1-score	support	Random Forest C				
P	recision	recall	11-Score	Support	р	recision	recall	f1-score	support
0	0.13	0.26	0.18	686	0	0.11	0.08	0.09	701
1	0.88	0.77	0.82	5000	1	0.88	0.92	0.90	5000
micro avg	0.71	0.71	0.71	5686	micro avg	0.81	0.81	0.81	5701
macro avg	0.51	0.52	0.50	5686	macro avg	0.49	0.50	0.49	5701
weighted avg	0.79	0.71	0.74	5686	weighted avg	0.78	0.81	0.80	5701
[[179 507]					[[53 648]				
[1152 3848]]	-1				[416 4584]]				
Support Vector					Support Vector				
р	recision	recall	f1-score	support	р	recision	recall	f1-score	support
0	0.17	0.58	0.26	686	0	0.12	0.44	0.19	701
1	0.91	0.60	0.72	5000	1	0.87	0.54	0.67	5000
					_				
micro avg	0.59	0.59	0.59	5686	micro avg	0.53	0.53	0.53	5701
macro avg	0.54	0.59	0.49	5686	macro avg	0.50	0.49	0.43	5701
weighted avg	0.82	0.59	0.66	5686	weighted avg	0.78	0.53	0.61	5701
[[400 286]					[[309 392]				
[2023 2977]]					[2287 2713]]				
Logistic Regres	sion 0.596	377066479	0714		Logistic Regres	sion 0.480	617435537	625	
р	recision	recall	f1-score	support		recision		f1-score	support
					The state of the s				
0	0.16	0.57	0.26	686	0	0.12	0.51	0.20	701
0 1	0.16 0.91	0.57 0.60	0.26 0.72	686 5000	0 1	0.12 0.87	0.51 0.48	0.20 0.62	701 5000
1	0.91	0.60	0.72	5000					5000
1 micro avg	0.91 0.60	0.60	0.72 0.60	5000 5686			0.48 0.48		5000 5701
1 micro avg macro avg	0.91 0.60 0.54	0.60 0.59	0.72 0.60 0.49	5000 5686 5686	1 micro avg macro avg	0.87	0.48	0.62	5000
1 micro avg	0.91 0.60	0.60	0.72 0.60	5000 5686	1 micro avg	0.87 0.48	0.48 0.48	0.62 0.48	5000 5701
1 micro avg macro avg	0.91 0.60 0.54	0.60 0.59	0.72 0.60 0.49	5000 5686 5686	1 micro avg macro avg	0.87 0.48 0.50	0.48 0.48 0.50	0.62 0.48 0.41	5000 5701 5701
micro avg macro avg weighted avg	0.91 0.60 0.54	0.60 0.59	0.72 0.60 0.49	5000 5686 5686	micro avg macro avg weighted avg	0.87 0.48 0.50	0.48 0.48 0.50	0.62 0.48 0.41	5000 5701 5701
micro avg macro avg weighted avg	0.91 0.60 0.54 0.82	0.60 0.59 0.60	0.72 0.60 0.49 0.67	5000 5686 5686 5686	micro avg macro avg weighted avg [[361 340]	0.87 0.48 0.50 0.78	0.48 0.48 0.50 0.48	0.62 0.48 0.41 0.56	5000 5701 5701 5701
micro avg macro avg weighted avg [[393 293] [2002 2998]] Linear Discrimi	0.91 0.60 0.54 0.82	0.60 0.60 0.59 0.60	0.72 0.60 0.49 0.67	5000 5686 5686 5686	micro avg macro avg weighted avg [[361 340] [2621 2379]] Linear Discrimi	0.87 0.48 0.50 0.78	0.48 0.48 0.50 0.48	0.62 0.48 0.41 0.56	5000 5701 5701 5701
micro avg macro avg weighted avg [[393 293] [2002 2998]] Linear Discrimi	0.91 0.60 0.54 0.82 nant Analy	0.60 0.60 0.59 0.60	0.72 0.60 0.49 0.67 0664790714 f1-score	5000 5686 5686 5686 5686	micro avg macro avg weighted avg [[361 340] [2621 2379]] Linear Discrimi	0.87 0.48 0.50 0.78 ant Analy	0.48 0.48 0.50 0.48 sis 0.877	0.62 0.48 0.41 0.56	5000 5701 5701 5701 5701 712 support
micro avg macro avg weighted avg [[393 293] [2002 2998]] Linear Discrimi	0.91 0.60 0.54 0.82 nant Analy recision 0.26	0.60 0.60 0.59 0.60 vsis 0.877 recall	0.72 0.60 0.49 0.67 0664790714 f1-score 0.02	5000 5686 5686 5686 5686	micro avg macro avg weighted avg [[361 340] [2621 2379]] Linear Discrimi p	0.87 0.48 0.50 0.78 nant Analy	0.48 0.48 0.50 0.48 sis 0.877 recall	0.62 0.48 0.41 0.56	5000 5701 5701 5701 712 support
micro avg macro avg weighted avg [[393 293] [2002 2998]] Linear Discrimi	0.91 0.60 0.54 0.82 nant Analy	0.60 0.60 0.59 0.60	0.72 0.60 0.49 0.67 0664790714 f1-score	5000 5686 5686 5686 5686	micro avg macro avg weighted avg [[361 340] [2621 2379]] Linear Discrimi	0.87 0.48 0.50 0.78 ant Analy	0.48 0.48 0.50 0.48 sis 0.877	0.62 0.48 0.41 0.56	5000 5701 5701 5701 5701 712 support
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micro avg macro avg weighted avg [[393 293] [2002 2998]] Linear Discrimi p	0.91 0.60 0.54 0.82 nant Analy recision 0.26 0.88	0.60 0.60 0.59 0.60 vsis 0.877 recall 0.01 1.00	0.72 0.60 0.49 0.67 0664790714 f1-score 0.02 0.93	5000 5686 5686 5686 5000	micro avg macro avg weighted avg [[361 340] [2621 2379]] Linear Discrimi p	0.87 0.48 0.50 0.78 Manant Analy Precision 0.00 0.88	0.48 0.50 0.48 sis 0.877 recall 0.00 1.00	0.62 0.48 0.41 0.56 70391159445 f1-score 0.00 0.93	5000 5701 5701 5701 701 712 support 701 5000
micro avg macro avg weighted avg [[393	0.91 0.60 0.54 0.82 nant Analy recision 0.26 0.88	0.60 0.60 0.59 0.60 vsis 0.877 recall 0.01 1.00 0.88	0.72 0.60 0.49 0.67 0664790714 f1-score 0.02 0.93 0.88	5000 5686 5686 5686 034 support 686 5000	micro avg macro avg macro avg weighted avg [[361 340] [2621 2379]] Linear Discrimi p 0 1	0.87 0.48 0.50 0.78 .nant Analy precision 0.00 0.88	0.48 0.48 0.50 0.48 sis 0.877 recall 0.00 1.00	0.62 0.48 0.41 0.56 0391159445' f1-score 0.00 0.93	5000 5701 5701 5701 712 support 701 5000
micro avg macro avg weighted avg [[393 293] [2002 2998]] Linear Discrimi p 0 1 micro avg macro avg weighted avg	0.91 0.60 0.54 0.82 nant Analy recision 0.26 0.88 0.88 0.57	0.60 0.60 0.59 0.60 vsis 0.877 recall 0.01 1.00 0.88 0.59	0.72 0.60 0.49 0.67 0664790714 f1-score 0.02 0.93 0.88 0.48	5000 5686 5686 5686 5686 034 support 686 5000 5686 5686	micro avg macro avg weighted avg [[361 340] [2621 2379]] Linear Discrimi p 0 1 micro avg macro avg weighted avg	0.87 0.48 0.50 0.78 	0.48 0.50 0.48 sis 0.877 recall 0.00 1.00	0.62 0.48 0.41 0.56 0391159445 f1-score 0.00 0.93 0.88 0.47	5000 5701 5701 5701 5701 712 support 701 5000 5701 5701
micro avg macro avg weighted avg [[393	0.91 0.60 0.54 0.82 nant Analy recision 0.26 0.88 0.88 0.57	0.60 0.60 0.59 0.60 vsis 0.877 recall 0.01 1.00 0.88 0.59	0.72 0.60 0.49 0.67 0664790714 f1-score 0.02 0.93 0.88 0.48	5000 5686 5686 5686 5686 034 support 686 5000 5686 5686	micro avg macro avg weighted avg [[361 340] [2621 2379]] Linear Discrimi p 0 1 micro avg macro avg	0.87 0.48 0.50 0.78 	0.48 0.50 0.48 sis 0.877 recall 0.00 1.00	0.62 0.48 0.41 0.56 0391159445 f1-score 0.00 0.93 0.88 0.47	5000 5701 5701 5701 5701 712 support 701 5000 5701 5701

Figure 5: PCA 1 Component Performance

Figure 6: PCA 2 Components Performance

• Adam - Adam is a third popularly used algorithm. It stands for adaptive moment estimation. It is similar to Adagrad and adds in another element of momentum, this time by storing an exponentially decreasing weight of the history of gradients. ²²

I started with **MSE** loss. MSE is commonly used in regression problems but it is very popular so I gave it a chance.

- MSE with SGD loss performed surprisingly well, though not perfectly. It shows notable average improvement over the baseline. The loss function was interesting because it "bumped up" in the beginning instead of descending rapidly as usually happens.
- MSE with Adagrad performed the best within MSE. It converged fairly quickly at 150 epochs and obtained a high average f1-score of 0.83. However, the f1-score of the 0 class (has cancer) was low because of the unbalanced classes. The downside of MSELoss is that there is no quick way to balance the classes.
- MSE with Adam performed very poorly, predicting nearly 100% no cancer, the dominant class. Perhaps Adam has a tendency to bias toward the dominant class since it takes into account a history of previous iteration performance.

²²https://towardsdatascience.com/the-3-best-optimization-methods-in-neural-networks-40879c887873

Bandon Fan 1	c1:£:	0.000000	E4E743EE3	
Random Forest				
	precision	recall	f1-score	support
0	0.14	0.04	0.06	662
_				
1	0.88	0.97	0.92	5000
micro avg	0.86	0.86	0.86	5662
macro avg	0.51	0.50	0.49	5662
eighted avg	0.80	0.86	0.82	5662
26 636]				
165 4835]]				
pport Vector	Classifier	0.550688	8025432709	
	precision	recall	f1-score	support
0	0.11	0.41	0.18	662
1	0.88	0.57	0.69	5000
-	0.00	0.57	0.05	5000
micro avg	0.55	0.55	0.55	5662
macro avg	0.50	0.49	0.43	5662
ighted avg	0.79	0.55	0.63	5662
272 390]				
2154 2846]]				
	ession 0.516	248675379	7245	
STATE WEBL	precision		f1-score	support
0	0.12	0.48	0.19	662
1	0.88	0.52	0.66	5000
micro avg	0.52	0.52	0.52	5662
macro avg	0.50	0.50	0.42	5662
ghted avg	0.79	0.52	0.60	5662
319 343]				
2396 2604]]				
ear Discrin	ninant Analy	sis 0.883	0801836806	782
	precision	recall	f1-score	support
0	0.00	0.00	0.00	662
1	0.88	1.00	0.94	5000
-	0.00	1.00	0.54	3000
micro avg	0.88	0.88	0.88	5662
macro avg	0.44	0.50	0.47	5662
eighted avg	0.78	0.88	0.83	5662
0 6603				
0 662]				
0 5000]]				

Figure 7: PCA 3 Components Performance

Figure 8: PCA 10 Components Performance

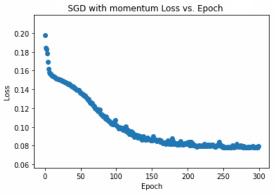
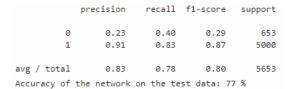


Figure: MSE with SGD Loss



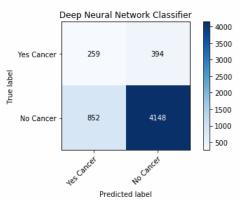


Figure: MSE with SGD Stats

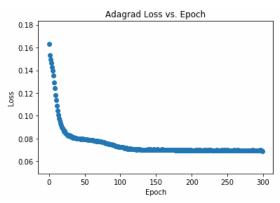


Figure: MSE with Adagrad Loss

	precision	recall	f1-score	support
0	0.26	0.18	0.21	653
1	0.90	0.93	0.91	5000
avg / total	0.82	0.85	0.83	5653
Accuracy of	the network	on the te	st data: 8	4 %

Yes Cancer - 115 538 - 4000 - 2000 - 2000 - 1000 - 1000

Figure: MSE with Adagrad Stats

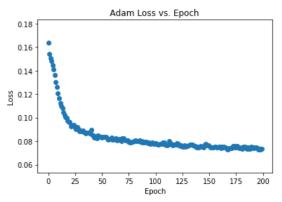


Figure: MSE with Adam Loss

support	f1-score	recall	precision	
653	0.00	0.00	0.00	0
5000	0.94	1.00	0.88	1
5653	0.83	0.88	0.78	avg / total

Accuracy of the network on the test data: 88 %

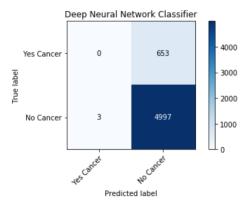


Figure: MSE with Adam Stats

I then tested **Binary Cross Entropy** as my loss function. It surprisingly performed quite similarly to MSE. I believe this could be due to the data not being well-structured enough for BCE to make as much a difference.

- The model had an interesting issue with Adam and SGD. The model would train for a certain number of epochs, but suddenly the loss would spike and stay constant. This destroyed any predictive power of the model.
- After that issue I tried limiting the number of epochs so the model would not experience the spike. This prevented the spike but the resulting model was still severely undertrained.
- The model trained terribly with SGD as well. It ran into the same "sudden loss increase issue."
- BCE with Adagrad was the best model and had the most balanced results. It still had trouble with class balancing though. I tried using the object's class weight modification parameter and manually balancing weights for classes, but could not achieve a better balance during the tests.

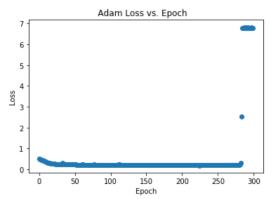
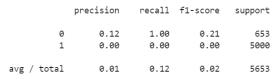


Figure: BCE with Adam Loss (bad)



Accuracy of the network on the test data: 11 %

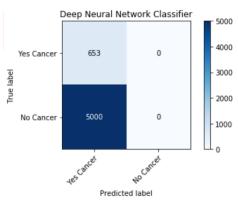


Figure: BCE with Adam Stats (bad)

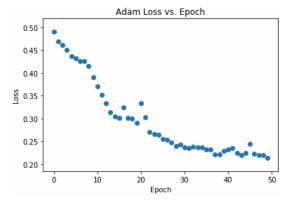


Figure: BCE with Adam Loss

```
tensor([1, 1, 1, ..., 1, 1, 1])
             precision
                          recall f1-score
         0
                  0.00
                            0.00
                                                 695
                                      0.00
                  0.88
                            1.00
                                      0.93
                                                5000
avg / total
                  0.77
                            0.88
                                      0.82
                                                5695
Accuracy of the network on the test data: 87 %
```

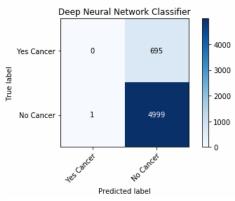


Figure: BCE with Adam Stats

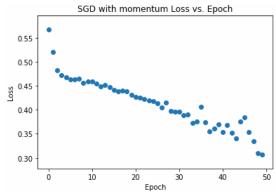
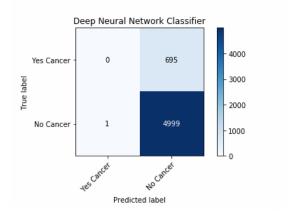


Figure: BCE with SGD Loss

support	f1-score	recall	precision	
695	0.00	0.00	0.00	0
5000	0.93	1.00	0.88	1
5695	0.82	0.88	0.77	avg / total
7 %	est data: 8	on the te	the network	Accuracy of

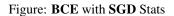


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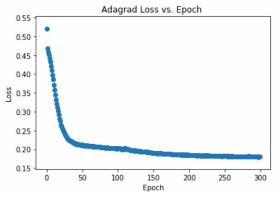


Figure: BCE with Adagrad Loss

	precision	recall f	1-score	support
0	0.26	0.21	0.23	653
1	0.90	0.92	0.91	5000
avg / total	0.82	0.84	0.83	5653
Accuracy of	the network o	on the test	data: 83 S	K

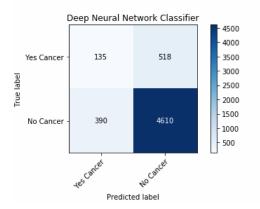
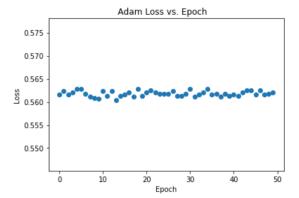
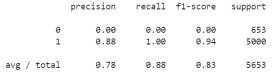


Figure: BCE with Adagrad Stats

Finally, while browsing the Pytorch forums I came across recommendations for the BCEWithLogitsLoss optimization function. This supposedly was more stable than using the original BCE. It also offered a simpler class weight parameter so I hoped it would solve my balancing issue. Unfortunately the results were subpar. It could not reduce its loss and would only predict the positive class. I may have used it wrong since it is a newer contribution to Pytorch, or it is just less stable.





Accuracy of the network on the test data: 88 %

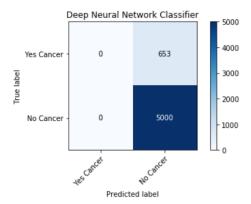


Figure: BCEWithLogitsLoss with Adam Stats

I additionally tested the neural network again with a more limited data size of 5000 train and 2000 test. This is to see if it improves the class balance issue as seen earlier in the paper, when increasing the data size also increased the gap between the classes. However, none of the stats improved and the overall performance simply decreased. This is likely due to the fact that deep neural networks can do so much more the more data they have, and is already complex enough to cause the class imbalances to appear with less data.

	precision	recall	f1-score	support
0	0.61	0.14	0.22	667
1	0.62	0.94	0.75	1000
micro avg	0.62	0.62	0.62	1667
macro avg	0.61	0.54	0.49	1667
weighted avg	0.61	0.62	0.54	1667
Confusion mat	rix, without	normaliz	ation	

[59 941]] Accuracy of the network on the test data: 61 %

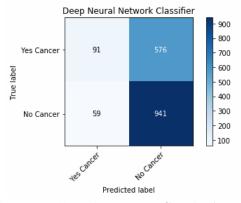


Figure: Smaller train/test sets BCE with Adagrad Stats

13 Results

- 214 The non-deep methods of Random Forest, SVC, Logistic Regression, and LDA performed decently
- but were not so accurate or robust.
- 216 In the end, the deep neural network performed the best. The best settings were Binary Cross Entropy
- with Adagrad loss. This obtained an f1-score for "has cancer" of 0.23 and an f1-score fr "healthy" of
- 218 0.91. The weighted average was 0.91.
- 219 This project shows definite predictive power of machine learning models, especially for DNNs, but I
- do not believe is is yet discriminatory enough to be used for real patients.

4 Afterthought and Further Potential

- 222 If I re-approached this project I would do more research and testing into the deep neural network. I
- believe more work can be done in terms of tuning the network architecture and weighting schemes.
- Less conventional (for this data) methods can be explored such as introducing convolution to extract
- 225 more predictive power from the data.

226 References

- 227 Please find all references in the footnotes throughout the paper.
- 228 An extra thank you to Dr. Majid Sarrafzadeh and our TA Orpaz in leading the class, presenting this
- project to us, and continued guidance throughout.