



# BREAST TUMOR CLASSIFICATION

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# INTRODUCTION: ABOUT THE DISEASE

Cancer begins when healthy cells in the breast change and grow out of control, forming a mass or sheet of cells called a tumor.

Breast cancer spreads when the cancer grows into adjacent organs or other parts of the body or when breast cancer cells move to other parts of the body through the blood vessels and/or lymph vessels.

There are 4 stages to a Breast Cancer. Stage 0 being early-stage and stage 3 locally advanced invasive breast cancer.

Types of breast cancer:

- Ductal carcinoma
- Invasive lobular carcinoma

# SYMPTOMS OF THE DISEASE

Lump in your breast or underarm



Pain and tenderness



Skin sores



# LITERATURE SURVEY



Rangayyan RM, Ayres FJ, Leo Desautels JE. A review of computer-aided diagnosis of breast cancer: toward the detection of subtle signs. J Franklin Inst. 2007 May;344(3-4):312-48. doi: 10.1016/j.jfranklin.2006.09.003.

This paper presents an overview of digital image processing and pattern analysis techniques to address several areas in CAD of breast cancer, including: contrast enhancement, detection and analysis of calcifications, detection and analysis of masses and tumors, analysis of bilateral asymmetry, and detection of architectural distortion.



Mohammed S.A., Darrab S., Noaman S.A., Saake G. (2020) Analysis of Breast Cancer Detection Using Different Machine Learning Techniques. In: Tan Y., Shi Y., Tuba M. (eds) Data Mining and Big Data. DMBD 2020. Communications in Computer and Information Science, vol 1234. Springer, Singapore. [https://doi.org/10.1007/978-981-15-7205-0\\_10](https://doi.org/10.1007/978-981-15-7205-0_10)

In this paper, we propose an approach that improves the accuracy and enhances the performance of three different classifiers: Decision Tree (J48), Naïve Bayes (NB), and Sequential Minimal Optimization (SMO). We also validate and compare the classifiers on two benchmark datasets: Wisconsin Breast Cancer (WBC) and Breast Cancer dataset.

# DATASET USED

- The Dataset was taken from [https://archive.ics.uci.edu/ml/datasets/Breast+Cancer+Wisconsin+\(Diagnostic\)](https://archive.ics.uci.edu/ml/datasets/Breast+Cancer+Wisconsin+(Diagnostic))
- This dataset consists of 570 people with 10 different parameters. We have divided the dataset into 470 for training and the remaining 100 for testing.
- Snapshot of the Dataset:

id	diagnosis	radius_me	texture_m	perimeter	area_me	smoothne	compactn	concavity	concave p	symmetry	fractal_dir	radius_se	texture_se	perimeter	area_se	smoothne	compactn	concavity	concave p	symmetry	fractal_dir	radius_wo
842302	1	17.99	10.38	122.8	1001	0.1184	0.2776	0.3001	0.1471	0.2419	0.07871	1.095	0.9053	8.589	153.4	0.006399	0.04904	0.05373	0.01587	0.03003	0.006193	25.38
842517	1	20.57	17.77	132.9	1326	0.08474	0.07864	0.0869	0.07017	0.1812	0.05667	0.5435	0.7339	3.398	74.08	0.005225	0.01308	0.0186	0.0134	0.01389	0.003532	24.99
84300903	1	19.69	21.25	130	1203	0.1096	0.1599	0.1974	0.1279	0.2069	0.05999	0.7456	0.7869	4.585	94.03	0.00615	0.04006	0.03832	0.02058	0.0225	0.004571	23.57
84348301	1	11.42	20.38	77.58	386.1	0.1425	0.2839	0.2414	0.1052	0.2597	0.09744	0.4956	1.156	3.445	27.23	0.00911	0.07458	0.05661	0.01867	0.05963	0.009208	14.91
84358402	1	20.29	14.34	135.1	1297	0.1003	0.1328	0.198	0.1043	0.1809	0.05883	0.7572	0.7813	5.438	94.44	0.01149	0.02461	0.05688	0.01885	0.01756	0.005115	22.54
843786	1	12.45	15.7	82.57	477.1	0.1278	0.17	0.1578	0.08089	0.2087	0.07613	0.3345	0.8902	2.217	27.19	0.00751	0.03345	0.03672	0.01137	0.02165	0.005082	15.47
844359	1	18.25	19.98	119.6	1040	0.09463	0.109	0.1127	0.074	0.1794	0.05742	0.4467	0.7732	3.18	53.91	0.004314	0.01382	0.02254	0.01039	0.01369	0.002179	22.88
84458202	1	13.71	20.83	90.2	577.9	0.1189	0.1645	0.09366	0.05985	0.2196	0.07451	0.5835	1.377	3.856	50.96	0.008805	0.03029	0.02488	0.01448	0.01486	0.005412	17.06
844981	1	13	21.82	87.5	519.8	0.1273	0.1932	0.1859	0.09353	0.235	0.07389	0.3063	1.002	2.406	24.32	0.005731	0.03502	0.03553	0.01226	0.02143	0.003749	15.49
84501001	1	12.46	24.04	83.97	475.9	0.1186	0.2396	0.2273	0.08543	0.203	0.08243	0.2976	1.599	2.039	23.94	0.007149	0.07217	0.07743	0.01432	0.01789	0.01008	15.09
845636	1	16.02	23.24	102.7	797.8	0.08206	0.06669	0.03299	0.03323	0.1528	0.05697	0.3795	1.187	2.466	40.51	0.004029	0.009269	0.01101	0.007591	0.0146	0.003042	19.19
84610002	1	15.78	17.89	103.6	781	0.0971	0.1292	0.09954	0.06606	0.1842	0.06082	0.5058	0.9849	3.564	54.16	0.005771	0.04061	0.02791	0.01282	0.02008	0.004144	20.42
846226	1	19.17	24.8	132.4	1123	0.0974	0.2458	0.2065	0.1118	0.2397	0.078	0.9555	3.568	11.07	116.2	0.003139	0.08297	0.0889	0.0409	0.04484	0.01284	20.96
846381	1	15.85	23.95	103.7	782.7	0.08401	0.1002	0.09938	0.05364	0.1847	0.05338	0.4033	1.078	2.903	36.58	0.009769	0.03126	0.05051	0.01992	0.02981	0.003002	16.84
84667401	1	13.73	22.61	93.6	578.3	0.1131	0.2293	0.2128	0.08025	0.2069	0.07682	0.2121	1.169	2.061	19.21	0.006429	0.05936	0.05501	0.01628	0.01961	0.008093	15.03
84799002	1	14.54	27.54	96.73	658.8	0.1139	0.1595	0.1639	0.07364	0.2303	0.07077	0.37	1.033	2.879	32.55	0.005607	0.0424	0.04741	0.0109	0.01857	0.005466	17.46
848406	1	14.68	20.13	94.74	684.5	0.09867	0.072	0.07395	0.05259	0.1586	0.05922	0.4727	1.24	3.195	45.4	0.005718	0.01162	0.01998	0.01109	0.0141	0.002085	19.07
84862001	1	16.13	20.68	108.1	798.8	0.117	0.2022	0.1722	0.1028	0.2164	0.07356	0.5692	1.073	3.854	54.18	0.007026	0.02501	0.03188	0.01297	0.01689	0.004142	20.96
849014	1	19.81	22.15	130	1260	0.09831	0.1027	0.1479	0.09498	0.1582	0.05395	0.7582	1.017	5.865	112.4	0.006494	0.01893	0.03391	0.01521	0.01356	0.001997	27.32
8510426	0	13.54	14.36	87.46	566.3	0.09779	0.08129	0.06664	0.04781	0.1885	0.05766	0.2699	0.7886	2.058	23.56	0.008462	0.0146	0.02387	0.01315	0.0198	0.0023	15.11
8510653	0	13.08	15.71	85.63	520	0.1075	0.127	0.04568	0.0311	0.1967	0.06811	0.1852	0.7477	1.383	14.67	0.004097	0.01898	0.01698	0.00649	0.01678	0.002425	14.5



# METHODOLOGY

- In this problem statement, we need to detect the breast cancer using the parameters given like radius of tumor etc.
- For performing this, we have used various algorithms
  - KNN
  - Logistic regression
  - Linear and Polynomial Support Vector Machines
  - Decision Tree, etc.

# METHODOLOGY

Upload the CSV file  
Separate training and testing data



**Step 1**

Send the parameters to different  
functions of various models



**Step 2**

Obtain the accuracy and plot  
the graph for the same



**Step 3**



# ALGORITHMS USED

## KNN

Stores all available data and classifies a new data point based on similarity

## Logistic Regression

Can take any real-valued number and map it into a value between 0 and 1, but never exactly at those limits.

## SVM's

Can take any real-valued number and map it into a value between 0 and 1, but never exactly at those limits.

## Decision Tree

Each internal node denotes a test, each branch represents an outcome of the test, and each leaf node holds a class label.

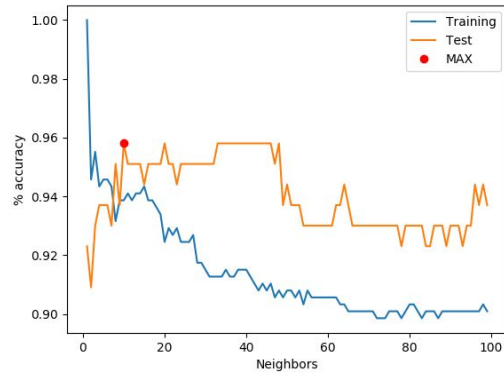
## Gradient Boosting

Ensemble of shallow trees in sequence with each tree learning and improving on the previous one.

## Naive-Bayes

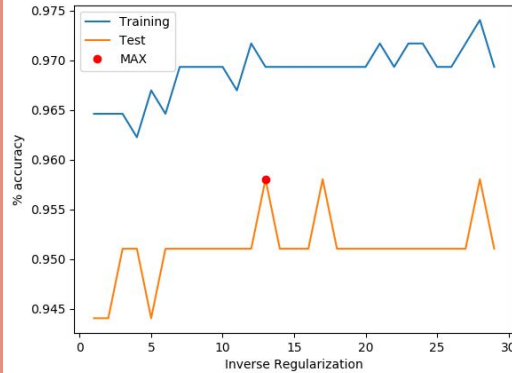
Models that assign class labels to problem instances, represented as vectors of feature values, where the class labels are drawn from some finite set.

# RESULT ANALYSIS

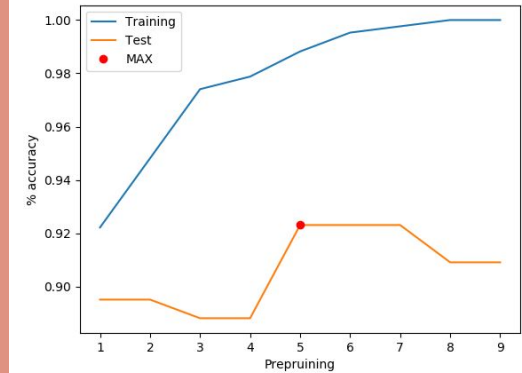


**KNN**

**Accuracy: 95.80% at  
10 neighbours**

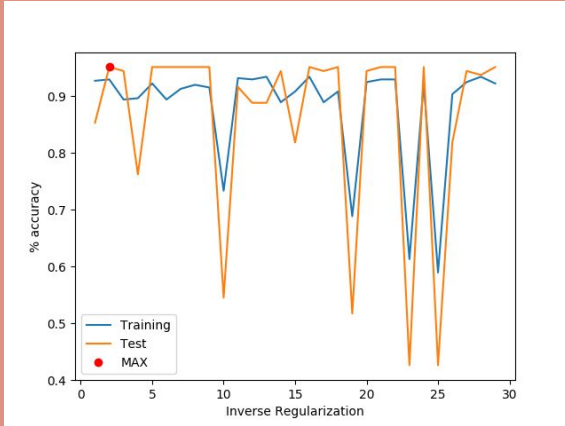


**Logistic Regression**  
**Accuracy: 95.80% at  
Inverse  
Regularization: 16**

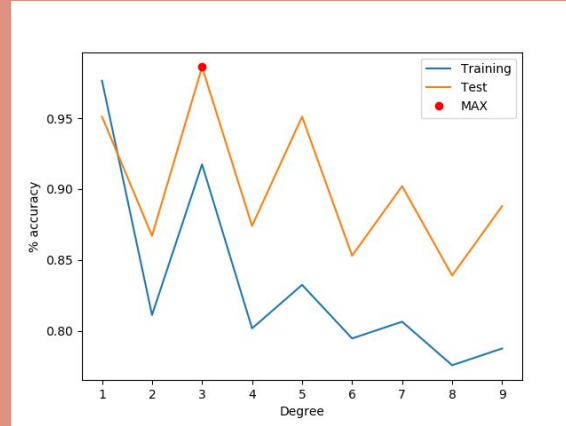


**Decision Tree**  
**Accuracy: 92.30% At  
Pre-pruning = 5**

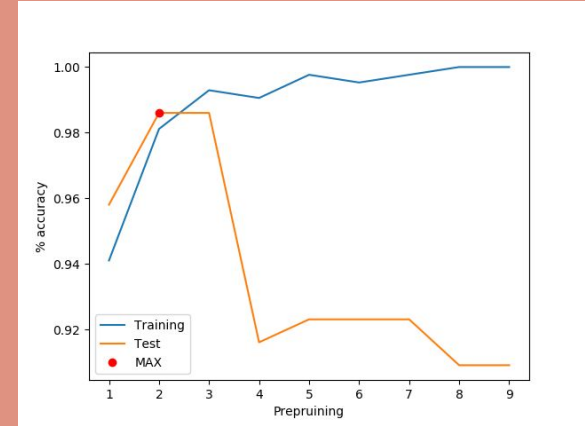
# RESULT ANALYSIS



**Linear SVM**  
Accuracy: 95.16% at  
Inverse  
Regularization: 20

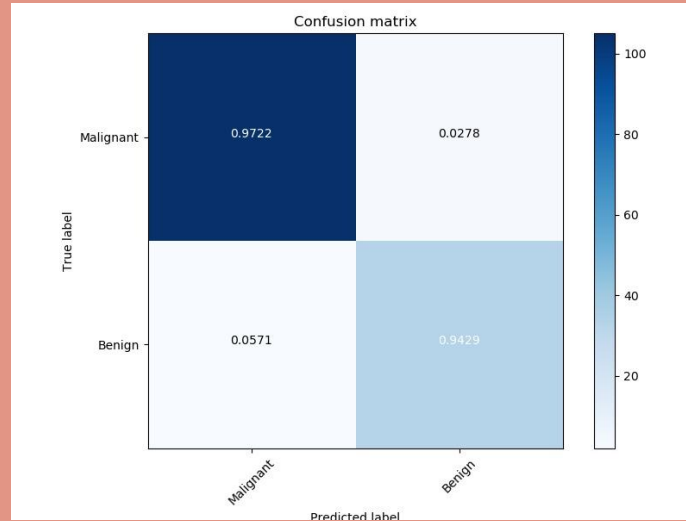


**Polynomial SVM**  
Accuracy: 98.60% at  
Degrees: 3



**Gradient Boosting**  
Accuracy: 98.60% at  
Estimator: 2

# RESULT ANALYSIS



**Naive-Bayes**  
**Accuracy: 96.50%**

From the above results obtained, we can see that Gradient Boosting and Polynomial SVM have the most accuracy of 98.60% .

# CONCLUSION



```
graph TD; A[CONCLUSION] --- B[ ]; B --- C[Implemented various algorithms. The best algorithm were Gradient Boosting and Polynomial SVM, having 98.60% accuracy]; B --- D[Creating hybrid algorithms which will improve accuracy even further]; B --- E[This helps in detecting whether the tumor is benign or malignant];
```

Implemented various algorithms. The best algorithm were Gradient Boosting and Polynomial SVM, having 98.60% accuracy

Creating hybrid algorithms which will improve accuracy even further

This helps in detecting whether the tumor is benign or malignant