

The slide features three blue spheres of varying sizes. A small sphere is in the upper left, a medium sphere is in the upper right, and a large sphere is in the lower right. Thin blue lines extend from the top-left and top-right corners towards the spheres, and another line extends from the top-right corner towards the large sphere in the bottom right.

Multi-Cancer Classification

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Your data preparation process

1. Resizing images.
2. Convert from RGB to gray scale .

Description of the models and techniques :

1. Brain Scan Breast Scan model : We use Bag Of Words , test and Train by SVM
2. Brain model : We use HOG To Extarct Features , test and train it by SVM Then save and load Model using pickle .
3. Breast model : : We use HOG To Extarct Features , test and train it by SVM Then save and load Model using pickle .

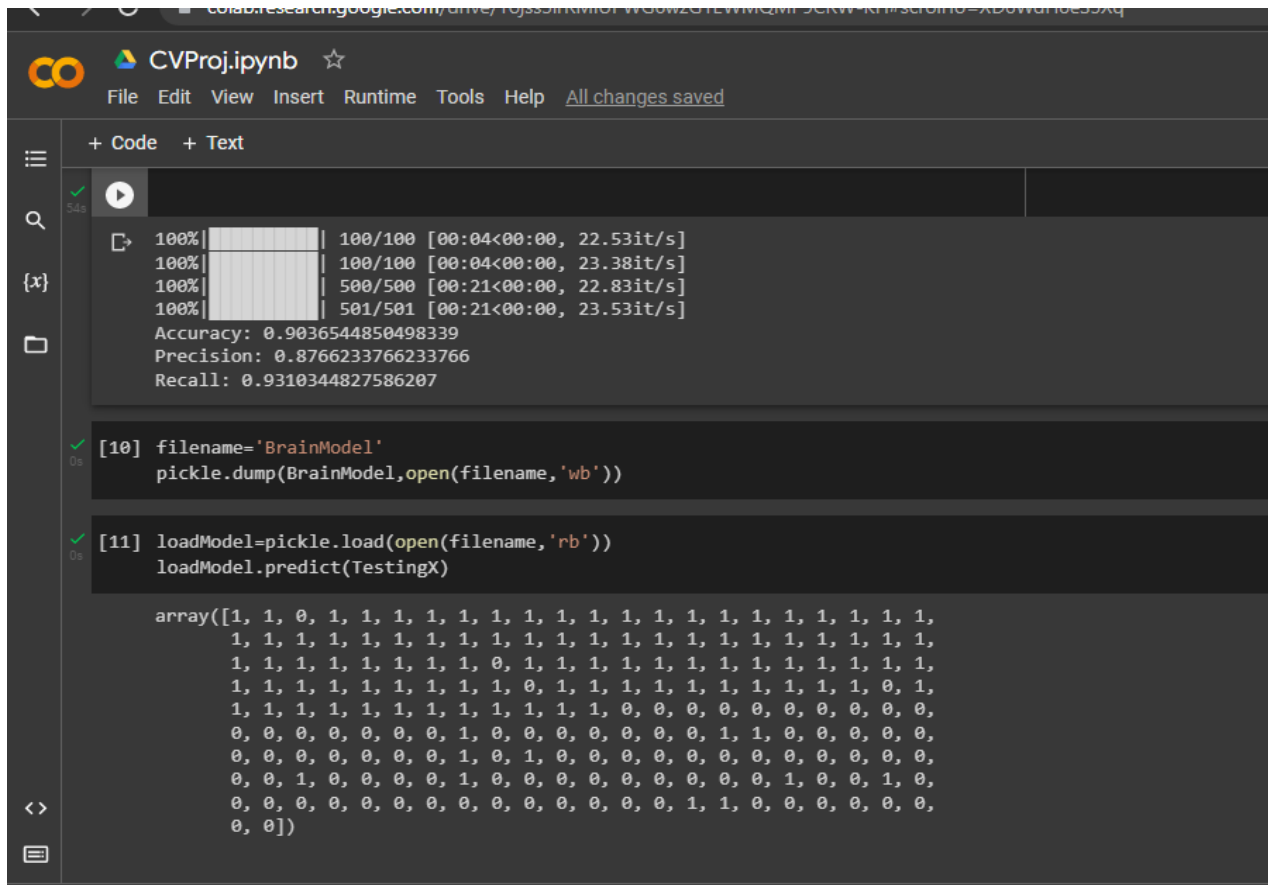
Training and Testing times for each model :

Models	Training	Testing
Brain Scan Breast Scan		
Brain	1 min : 40s	8s
Breast	1 min : 58s	5s

Image Classification training and testing accuracy:

Models	Training accuracy	Testing accuracy
Brain Scan Breast Scan		
Brain	100%	90.3%
Breast	100%	72.2%

The test sets classification with visualization



The screenshot shows a Jupyter Notebook titled "CVProj.ipynb" with a dark theme. The interface includes a top bar with "File", "Edit", "View", "Insert", "Runtime", "Tools", and "Help" menus, and a status bar indicating "All changes saved". The left sidebar contains icons for file management, search, and code execution. The main area displays a code cell with the following content:

```
+ Code + Text

100%|██████████| 100/100 [00:04<00:00, 22.53it/s]
100%|██████████| 100/100 [00:04<00:00, 23.38it/s]
100%|██████████| 500/500 [00:21<00:00, 22.83it/s]
100%|██████████| 501/501 [00:21<00:00, 23.53it/s]
Accuracy: 0.9036544850498339
Precision: 0.8766233766233766
Recall: 0.9310344827586207

[10] filename='BrainModel'
pickle.dump(BrainModel,open(filename,'wb'))

[11] loadModel=pickle.load(open(filename,'rb'))
loadModel.predict(TestingX)

array([1, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
       1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
       1, 1, 1, 1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
       1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 1, 1, 1, 1, 1, 1, 1, 1, 1,
       1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0,
       0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0, 0,
       0, 0, 0, 0, 0, 0, 0, 1, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
       0, 0, 1, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0,
       1, 0,
       0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0, 0,
       0, 0, 0, 0])
```

The output shows progress bars for four data points, followed by accuracy, precision, and recall metrics. The code cell contains three lines of Python code: saving a model, loading the model, and making predictions. The final output is a large array of 0s and 1s.

```
print("Accuracy:", metrics.accuracy_score(y_test, y_pred))
```

```
100%|██████████| 39/39 [00:03<00:00, 12.11it/s]
100%|██████████| 30/30 [00:01<00:00, 16.87it/s]
100%|██████████| 30/30 [00:01<00:00, 16.54it/s]
100%|██████████| 398/398 [00:22<00:00, 17.41it/s]
100%|██████████| 180/180 [00:10<00:00, 17.58it/s]
100%|██████████| 103/103 [00:05<00:00, 17.39it/s]
Accuracy: 0.7226277372262774
```

```
[19] filename='BreastModel'
pickle.dump(BreastModel, open(filename, 'wb'))
```

```
loadModel=pickle.load(open(filename, 'rb'))
loadModel.predict(X_test)
```

```
array([3, 5, 5, 5, 5, 5, 5, 5, 3, 5, 5, 5, 5, 5, 5, 5, 4, 5, 5, 5,
        5, 5, 5, 5, 4, 4, 5, 5, 5, 5, 5, 5, 5, 5, 4, 5, 5, 5, 3, 4,
        5, 5, 5, 5, 5, 5, 5, 3, 5, 5, 5, 5, 4, 5, 5, 5, 5, 4, 5, 5,
        5, 5, 5, 5, 4, 5, 5, 5, 5, 3, 4, 4, 5, 5, 5, 3, 5, 3, 5, 5,
        4, 3, 5, 5, 5, 5, 5, 4, 4, 4, 5, 5, 5, 4, 5, 4, 5, 3, 5, 5,
        5, 5, 4, 5, 4, 3, 5, 5, 4, 4, 5, 5, 4, 5, 5, 4, 4, 5, 4, 5,
        5, 3, 5, 4, 5])
```

Disk 84.47 GB available

0s completed at 8:34 PM

Brain Model Accuracy and Testing Time:

```
BrainModel=svm.SVC(kernel='linear')
BrainModel.fit(X_train, y_train)
y_pred = BrainModel.predict(X_test)
N_pred=BrainModel.predict(X_train)
```

```
print("Testing Accuracy:", metrics.accuracy_score(y_test, y_pred))

print("Precision:", metrics.precision_score(y_test, y_pred))

print("Recall:", metrics.recall_score(y_test, y_pred))

print(" Training Accuracy:", metrics.accuracy_score(y_train, N_pred))
```

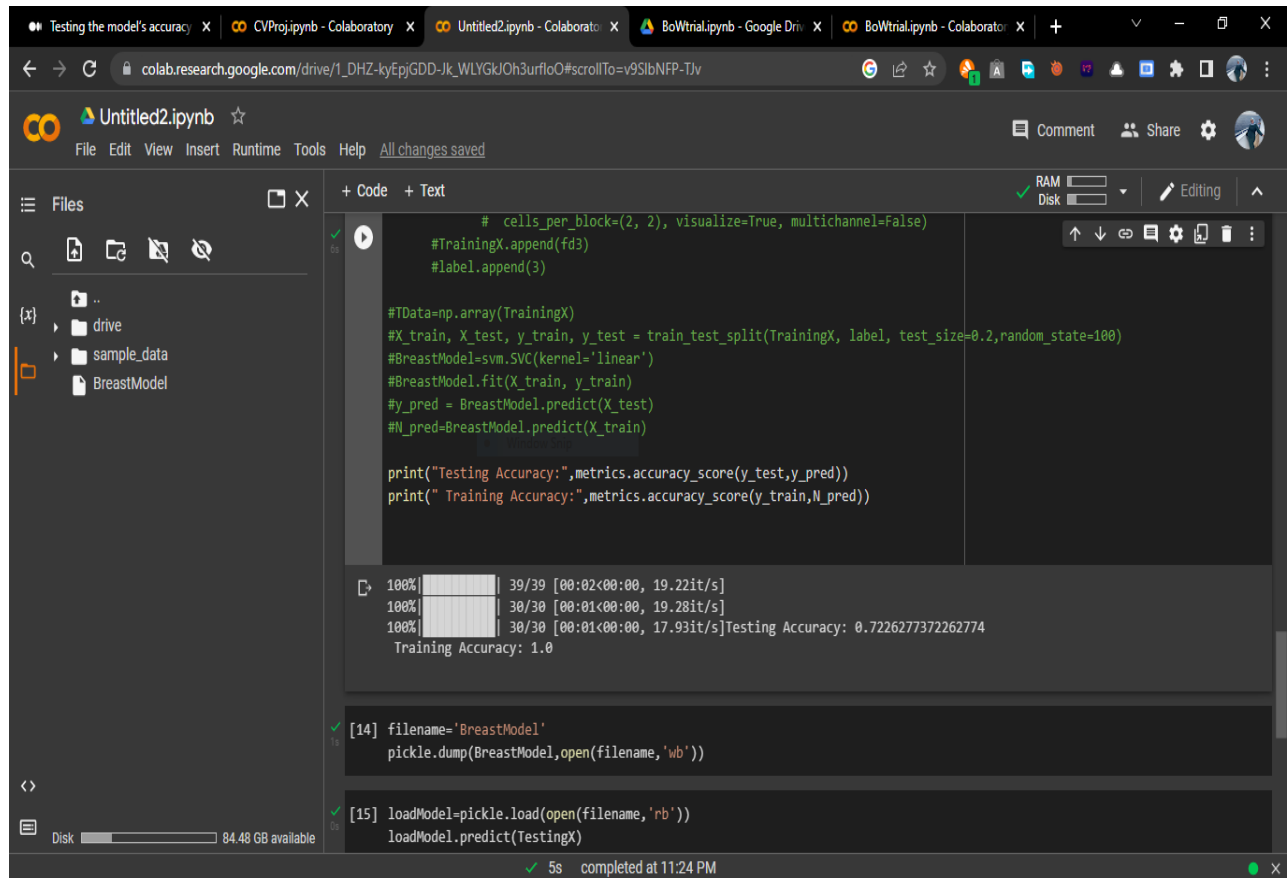
```
100%|██████████| 100/100 [00:04<00:00, 21.86it/s]
100%|██████████| 100/100 [00:05<00:00, 19.66it/s]
100%|██████████| 500/500 [00:22<00:00, 22.10it/s]
100%|██████████| 501/501 [00:22<00:00, 22.72it/s]
Testing Accuracy: 0.9036544850498339
Precision: 0.8766233766233766
Recall: 0.9310344827586207
Training Accuracy: 1.0
```

```
[10] filename='BrainModel'
pickle.dump(BrainModel, open(filename, 'wb'))
```

Disk 84.69 GB available

57s completed at 11:05 PM

Breast Model and Testing Time:



The screenshot displays a Google Colab notebook interface. The top bar shows the notebook title 'Untitled2.ipynb' and various menu options like File, Edit, View, Insert, Runtime, Tools, and Help. The left sidebar shows the file explorer with a directory structure including 'drive', 'sample_data', and 'BreastModel'. The main code area contains the following Python code:

```
# cells_per_block=(2, 2), visualize=True, multichannel=False)
#TrainingX.append(fd3)
#label.append(3)

#TData=np.array(TrainingX)
#X_train, X_test, y_train, y_test = train_test_split(TrainingX, label, test_size=0.2, random_state=100)
#BreastModel=svm.SVC(kernel='linear')
#BreastModel.fit(X_train, y_train)
#y_pred = BreastModel.predict(X_test)
#N_pred=BreastModel.predict(X_train)

print("Testing Accuracy:",metrics.accuracy_score(y_test,y_pred))
print(" Training Accuracy:",metrics.accuracy_score(y_train,N_pred))
```

The output of the code execution is shown in the bottom panel, displaying progress bars and accuracy scores:

```
100%|██████████| 39/39 [00:02<00:00, 19.22it/s]
100%|██████████| 30/30 [00:01<00:00, 19.28it/s]
100%|██████████| 30/30 [00:01<00:00, 17.93it/s]Testing Accuracy: 0.7226277372262774
Training Accuracy: 1.0
```

Below the output, the code continues with saving and loading the model:

```
[14] filename='BreastModel'
pickle.dump(BreastModel,open(filename,'wb'))

[15] loadModel=pickle.load(open(filename,'rb'))
loadModel.predict(TestingX)
```

The bottom status bar indicates that the code was completed at 11:24 PM.

BOW Histogram From Brain and Breast Scan Model :

