

# **B.Tech Project Work Report (CSIR32)**



## **Title:-**

Land Cover Classification on Hyper-spectral Imagery using CNN - SVM Hybrid

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## CERTIFICATE

I hereby certify that the work which is being presented in this B.Tech Project Work (**CSIR32**) report entitled “**Land Cover Classification on Hyper-spectral imagery using CNN-SVM Hybrid**”, in partial fulfillment of the requirements for the award of the **B.Tech (Computer Engineering)** is an authentic record of my own work carried out during a period from January 1<sup>st</sup>, 2020 to May 30<sup>th</sup>, 2020 under the supervision of **Ritu Garg, Assistant Professor, Computer Engineering Department.**

The matter presented in this project report has not been submitted for the award of any other degree elsewhere.

*Signature of Candidate*

**Himanshu Garg (11710589)**

This is to certify that the above statement made by the candidates is correct to the best of my knowledge.

*Signature with date*

*Faculty Mentor*

**Ritu Garg**

*Asst. Professor*

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# 1. Introduction

The Physical material present on the surface of the earth is known as Land Cover. Land covers include grass, trees, water, crops, etc. The process to capture information is important to identify the region each land cover holds. The process of capturing information on the land cover can be done in two ways Remote Sensing, Field Survey. The methodology used in this paper to classify land cover is based on remote sensing. Remote sensing is the science of obtaining information about objects or areas from a distance, typically from aircraft or satellites. The information collected is the amount of data reflected by the earth's surface. The collection of data is done by remote sensors. Several types of sensors are used to collect the data some are LANDSAT, MODIS, etc. These Sensors transmit the electromagnetic waves to the surface of the earth and then collect the number of waves that are reflected. These sensors transmit waves in different continuous ranges of the electromagnetic spectrum known as bands. Each band is represented by its wavelength and bandwidth. The scale of each band is grey-scaled. All these bands make a satellite image. The information of these bands can be used for the identification of materials on the surface of the earth. In this project, these images are used for land cover classification. These bands are used to calculate indices such as ENVI, NDWI, NDVI, etc. These indices are used to define the land cover by using machine learning algorithms. These indices are applied on pixel-wise or on a certain area of the satellite image. These indices are widely used in the field of crop type classification.

Many machine learning algorithms are used for hyperspectral image classification. Large dimensions/bands can be a problem for the classification problem. Therefore, Dimensionality Reduction (DR) is used to reduce the dimension of HSI. In this project, the Principal Component Analysis (PCA) algorithm is used to extract the best features. Several kinds of research were conducted on the Hyperspectral images using SVM for classification purposes. But, currently, there are so many algorithms available that can classify HSI with better accuracy than SVM. Newly researched deep learning models (Ex:-CNN) also results in better accuracy than SVM. CNN uses feature maps and applies Neural Networks on them.

In this project, a CNN-SVM based hybrid model is used which uses the feature maps extracted by CNN layers and then classifies them using SVM. The features extracted by CNN results better because CNN generates all possible maps by applying different filters and select only better features. This results in better accuracy than SVM or even CNN. This model is applied to the INDIAN-PINES dataset.

The result with increasing accuracy can be seen as the accuracy obtained by SVM is 83.67(%), the accuracy obtained by only CNN is 85.48(%) and the accuracy obtained by the hybrid model of CNN-SVM is 92.31(%).

## 2. Motivation

Accurate land cover classification is very important for environmental sustainability. Manual classification of land cover is a very difficult task because it is not possible to classify every place on earth. Most of the existing land cover approaches are pixel-based single-date multi-spectral images using conventional techniques like random forests (RFs), neural networks (NNs), and support vector machines (SVMs). They all provide classification with an unacceptable accuracy

Nowadays, there is a significant improvement in satellite systems and sensors that acquire data with improved spectral, spatial, radiometric, and temporal characteristics. Recently, several major satellites remotely sensed datasets become more affordable and feasible with much higher spatial resolutions acquired by the Landsat systems, ASTER (Advanced Space-borne Thermal Emission and Reflection Radiometer), and Sentinels, exploiting multi-temporal information for land cover classification. However, working with such higher-resolution multi-temporal, multi-spectral imagery datasets is facing some crucial challenges mainly caused by the frequent occurrences of pixels contaminated by clouds or shadows.

In the last decade, there is rapid development in deep learning algorithms like CNN. Compared to traditional classifiers like random forest and support vector machine, DCNN does not need extraction and selection of hand-crafted features. This motivated the researchers in the remote sensing community to investigate its usefulness for remote sensing image analysis. Moreover, convolutional neural networks (CNN) are not considered to be fully connected neural nets. CNN's have convolution and pooling layers and fully connected layers, whereas ANN has only fully connected layers, which is a key difference.

Therefore, in this project, the focus is to exploit multispectral remote sensing data by using a hybrid model based on convolutional neural networks and support vector machine to have land cover classification with improved accuracy. A system is designed that classifies the image using SVM based on the features extracted from the last layer of CNN. This hybrid model results in greater accuracy than CNN or SVM.

### 3. Literature Survey

In the field of land cover mapping using remote sensing, many works had already been carried out. Many researches were carried out using different machine learning algorithms. Some of the researches carried out are -

**1. Land cover identification using CNN on LandSat Imagery :-** In this work, land cover is mapped using CNN(Convolutional Neural Networks). The accuracy obtained using only CNN is 83%.

**Reference:-**

X. Zhao, L. Gao, Z. Chen, B. Zhang and W. Liao, "CNN-based Large Scale Landsat Image Classification," *2018 Asia-Pacific Signal and Information Processing Association Annual Summit and Conference (APSIPA ASC)*, Honolulu, HI, USA, 2018, pp. 611-617, doi: 10.23919/APSIPA.2018.8659654.

**2. Land cover identification using SVM :-** In this work, land cover identification is done using only SVM(Support Vector Machine). The accuracy Obtained using only SVM is much lower as compare to that from using only CNN.

**Reference:-**

B. Baassou, M. He and S. Mei, "An accurate SVM-based classification approach for hyperspectral image classification," *2013 21st International Conference on Geoinformatics*, Kaifeng, 2013, pp. 1-7, doi: 10.1109/Geoinformatics.2013.6626036.

Based on the results obtained from researches using only CNN and SVM it can be shown that the accuracy from CNN is better than SVM in most cases. So, In this project a hybrid model of both CNN and SVM is used. Which uses features from CNN and train on SVM.

The accuracy comparison of only CNN, only SVM and CNN-SVM hybrid on indian pines dataset can be seen as:-

Model Approach	Accuracy Obtained on training model on indian pines dataset
Only SVM	83.67479674796748 (%)
Only CNN	85.48575639724731 (%)
CNN-SVM Hybrid	92.31369488880219 (%)

## 4. Dataset Description

The dataset used in this project is the **Indian Pines Hyper-Spectral Dataset**. Indian pines dataset contains images with mainly two resolutions.

- **Spatial Resolution:** - It shows how many areas a pixel in image occupies. A spatial resolution of 10m means the 1-pixel cover ground area of 10m x 10m.
- **Spectral Resolution:** - Spectral resolution describes the ability of a sensor to define fine wavelength intervals. The finer the spectral resolution, the narrower the wavelength ranges for a particular channel or band.

Hyperspectral imagery with a resolution of  $M \times N \times B$  can be seen as follows:

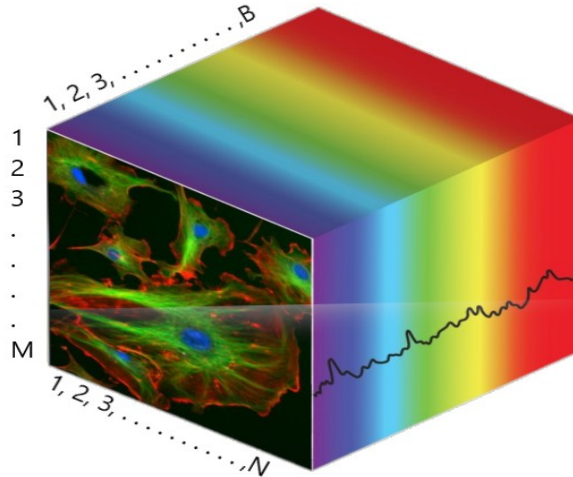


Fig: - Hyper-Spectral image with dimensions  $N \times M \times B$

The Indian pines dataset contains 145x145 pixels with 200 bands and ground truth resolution of 20m. Each pixel of a band corresponds to the spectral characteristics of a particular area in the frequency range corresponding to the sensor range. This dataset contains a total of 16 classes. The ground truth contains a total of 10249 pixels. The pixels in the ground truth corresponding to these classes are not evenly distributed. So, we randomly select 75% of these pixels for training purposes and the remaining 25% for testing these pixels on the machine learning model. These pixels are firstly scaled in the range from  $[-1, 1]$ . After that

patches are created using the ground truth corresponding to each pixel. The pixel distribution in Indian Pines is as follows:

#	Class	Samples	No. of Train Samples	No. of Test Samples
1	Alfalfa	46	35	11
2	Corn-notill	1428	1071	357
3	Corn-mintill	830	622	208
4	Corn	237	178	59
5	Grass-pasture	483	362	121
6	Grass-trees	730	547	183
7	Grass-pasture-m	28	21	7
8	Hay-windrowed	478	358	120
9	Oats	20	15	5
10	Soybean-notill	972	729	243
11	Soybean-mintill	2455	1841	614
12	Soybean-clean	593	445	148
13	Wheat	205	154	51
14	Woods	1265	949	316
15	Buildings-Grass-	386	289	97
16	Stone-Steel-Tow	93	70	23
	<b>TOTAL</b>	<b>10249</b>	<b>7686</b>	<b>2563</b>

In this dataset bands that are affected by atmospheric absorption are removed. So, the training is done only on 200 bands of the image.



## 5. Classification Algorithms

There are two classification algorithms used in this project. First, we use CNN (Convolutional Neural Network); this algorithm is used to extract the important features needed for classification. Then the features are extracted before the last layer. These features are then used to train the dataset on the second model. The second model used for the final classification is SVM (Support Vector Machine). SVM trains dataset based on features from CNN. Detailed explanations of these algorithms are given as follows:

### 5.1 CNN (Convolutional Neural Networks)

A Convolutional Neural Network (CNN) is a Deep Learning algorithm that takes an image as an input and then can classify them by assigning labels accordingly. It is a type of feed-forward artificial neural network. The pre-processing required in CNN is

lower as compared to other classification algorithms. CNN has wide applications in the field of natural language processing, audio and video recognition, recommender systems, etc.

CNN is comprised of one or more layers. CNN is generally made of two types of layers: convolutional layers, pooling layers. These layers generate random patches corresponding to the input patch. This patch is passed to the convolution layer to generate feature maps. These feature maps can be pooled or can be used to generate even more feature maps. Generally, the activation function used in the convolution layer is **ReLU Activation Function**.

$$y = \max(0, x)$$

After the preprocessing phase (generation of feature maps and classification) is done, these features can be used to classify the patches. Generally, ANN (Artificial Neural Networks) is used as a classifier in the last layer of the CNN model for classification. The basic architecture of CNN can be seen as:

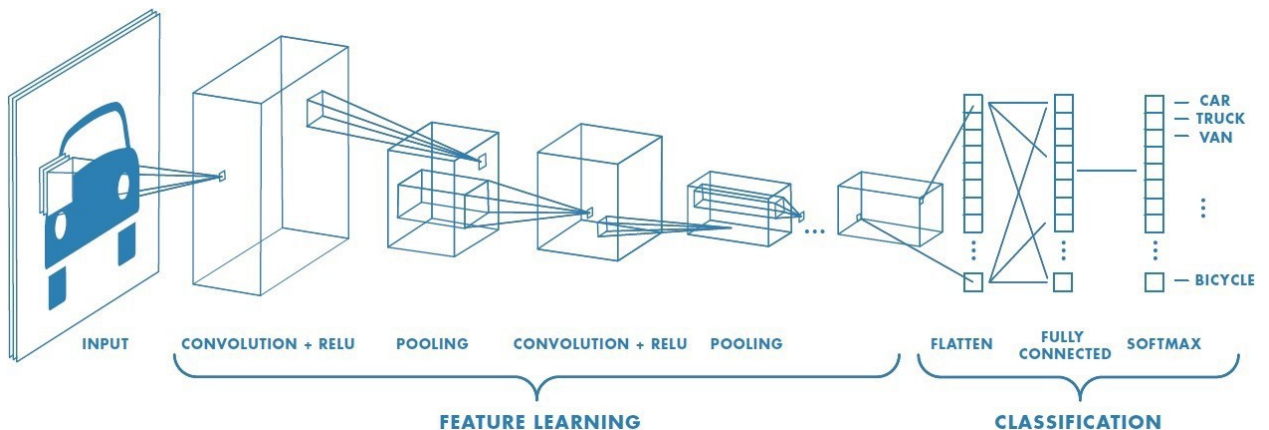


Fig: - The Convolutional Neural Network (CNN) architecture

Layers of CNN can be explained as:

1. **Input layer:** - This is the first layer of CNN which takes an image as an input. In this case of the colored image, the size of the input is  $M \times N \times 3$ , where 3 represents no of channels/bands (R, G, and B).
2. **Convolutional Layer:** - The element involves in carrying the convolution operation in this layer is known as kernel/filter. The main objective of this layer is to extract high-level features such as edges etc. This layer applies the kernel to generate the feature maps. These maps then again pooled to extract some high-level features.
3. **Pooling Layer:** - This layer works almost the same as the convolution layer to decrease the spatial size of the convolved feature. This reduction is done to reduce the computational power required to process the data. Furthermore, it is useful for extracting dominant features that are rotational and positional invariant, thus maintaining the process of effectively training of the model. The most common pooling done is max pooling. In this pooling, the max value is taken from the portion of the image on which filter is applied.
4. **Flatten Layer:** - This layer is used to flatten the resulted feature maps from the max-pooling layer to provide input to the fully connected layer.
5. **Fully Connected Layer:** - This layer is used to train the model on input data by extracting dominating features for that image. This layer provides the input flattened by the flattening layer to the feed-forward neural network and the backpropagation is applied to every image. After training of the model is done images taken by the input layer can be classified easily using the weights obtained after training. For dividing or classifying this image the **Softmax function** is used which divides the image according to the probability of which class it belongs.

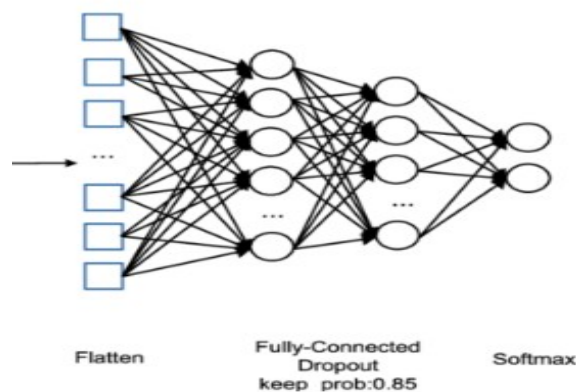
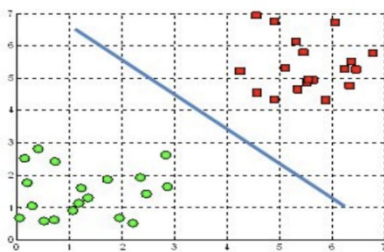


Fig: - Fully Connected layer with input from pooling layer

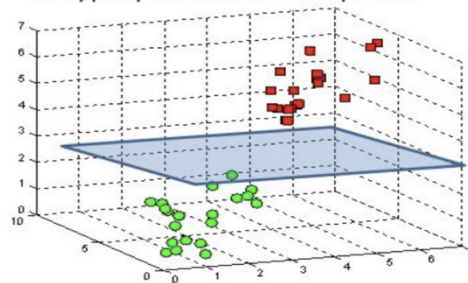
## 5.2 Support Vector Machine(SVM)

SVM is a type of classifier that is used to classify an input in a certain class. The objective of the support vector machine algorithm is to find a hyperplane in N-dimensional space (N - the number of features) that distinctly classifies the data points. Hyperplanes are decision boundaries that help in classify input. The area on either side of a hyperplane can be considered as a different class. Support vectors are the data points that are closer to other points. Support vectors help in defining the position and orientation of the hyperplane. A hyperplane can be seen as:

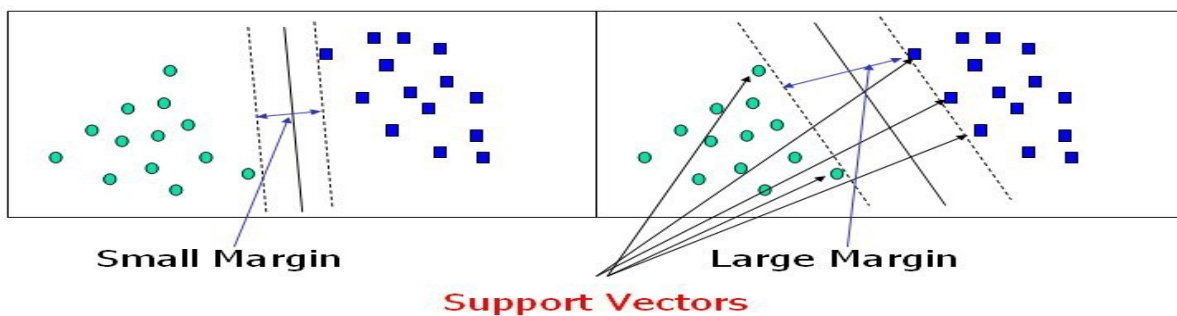
A hyperplane in  $\mathbb{R}^2$  is a line



A hyperplane in  $\mathbb{R}^3$  is a plane



(a)



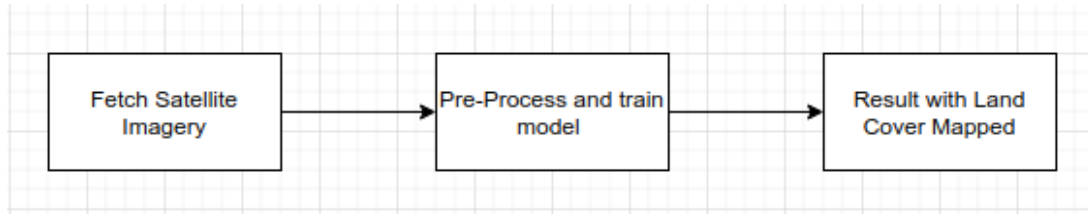
(b)

Fig: - (a) Hyper-plane to divide two classes, (b) Support Vectors

## 6. Data Flow Diagram

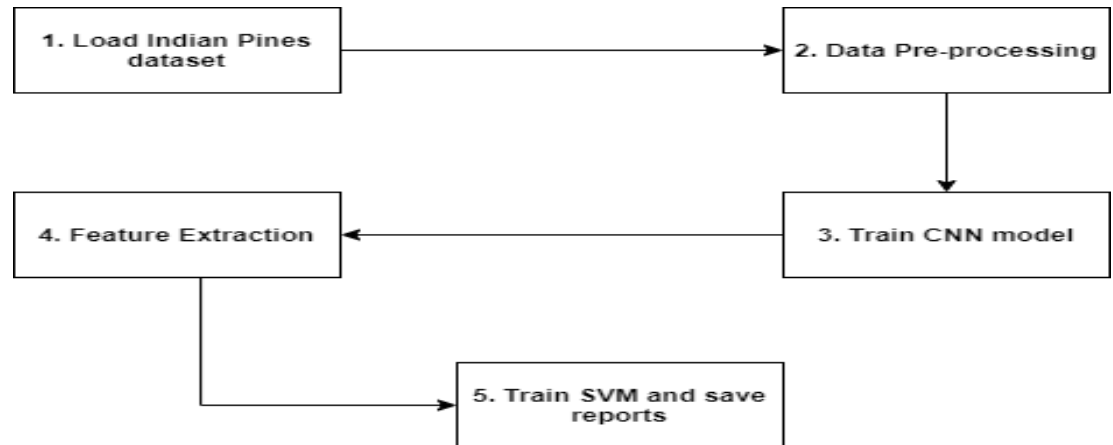
### 6.1 Level 0 DFD

This Level of flow diagram represents the abstraction view of model :-



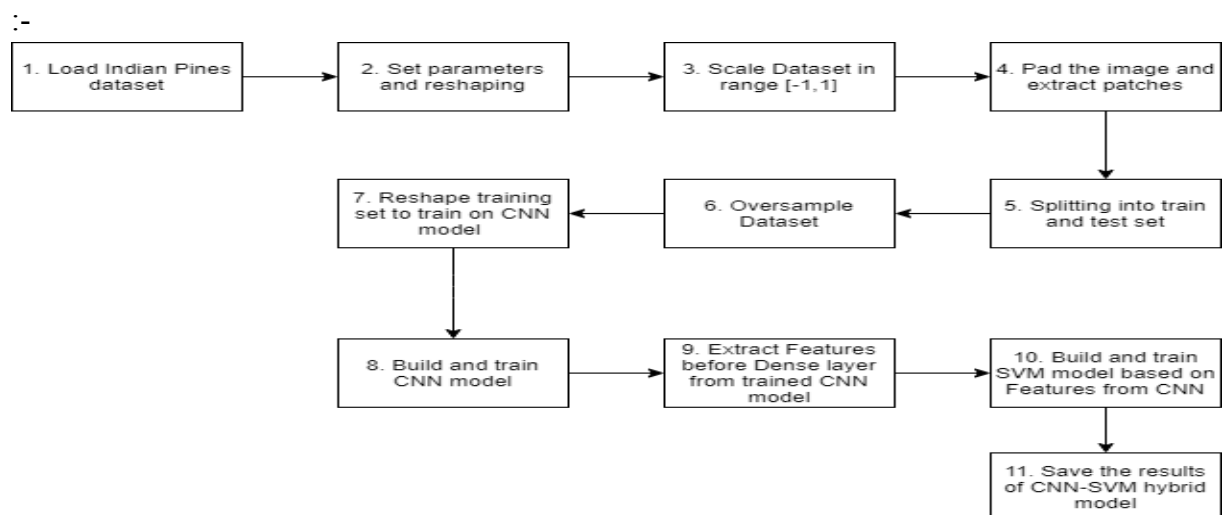
### 6.2 Level 1 DFD

This Level of flow diagram high level 0 -level DFD is breakdown into sub-process :-



### 6.3 Level 2 DFD

This Level of flow diagram high level 1-level DFD is breakdown into sub-process :-

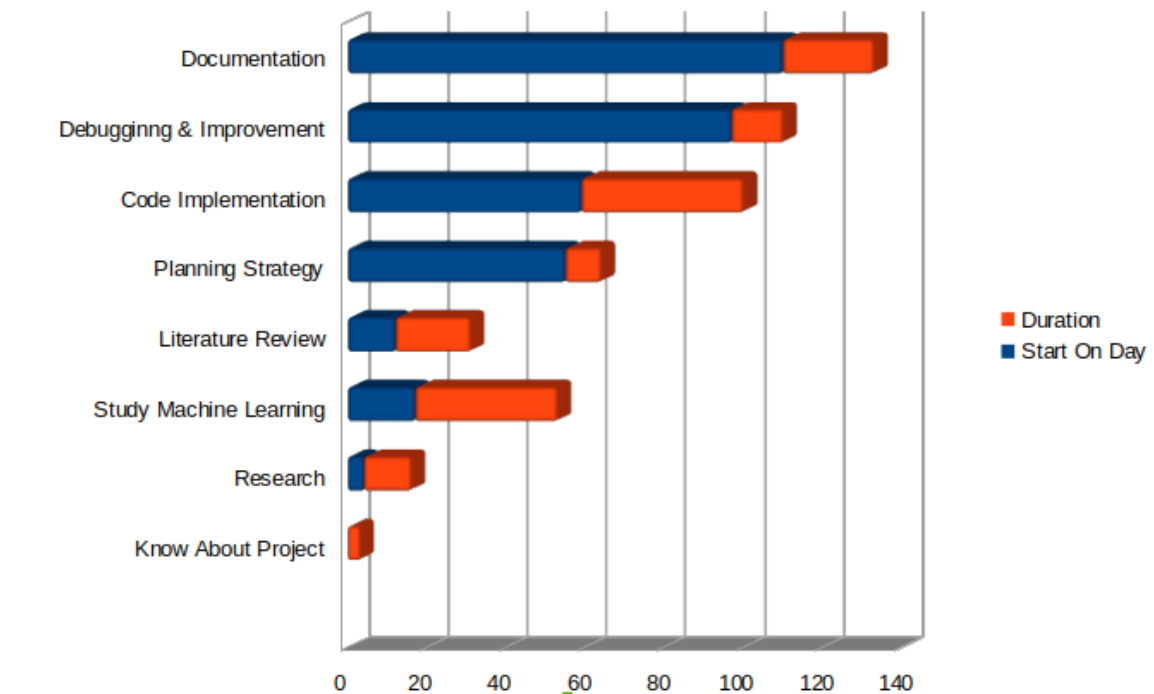


## 7. Gantt Chart

This Gantt Chart represents the duration that each section in this project takes :-

1. Know About Project
2. Research
3. Study Machine Learning
4. Literature Review
5. Planning & Studying
6. Implementation & Research
7. Debugging & Improvement
8. Documentation

Machine Learning is the core section for this project. In this project, CNN and SVM is used as machine learning algorithms.



**Fig: - Gantt Chart Representing Duration of Each Task**

## 8. Implementation Details

Classification of land cover is a very crucial task. But, recently many works are carrying out on this topic. So, hyperspectral images are used for classification. The classification can be done by applying machine learning models to images. Since hyperspectral images are large in dimensions and can have large no of bands, this can make the training of image a lot costly. Normal computers with no GPU or GPU with less computing power cannot work effectively for this purpose because training can take a lot more time. So, for training purposes, we need a GPU with at least 20GB RAM. The model used in this project is a hybrid of CNN and SVM. The methodology used in this project can be seen with the help of the chart as follows:

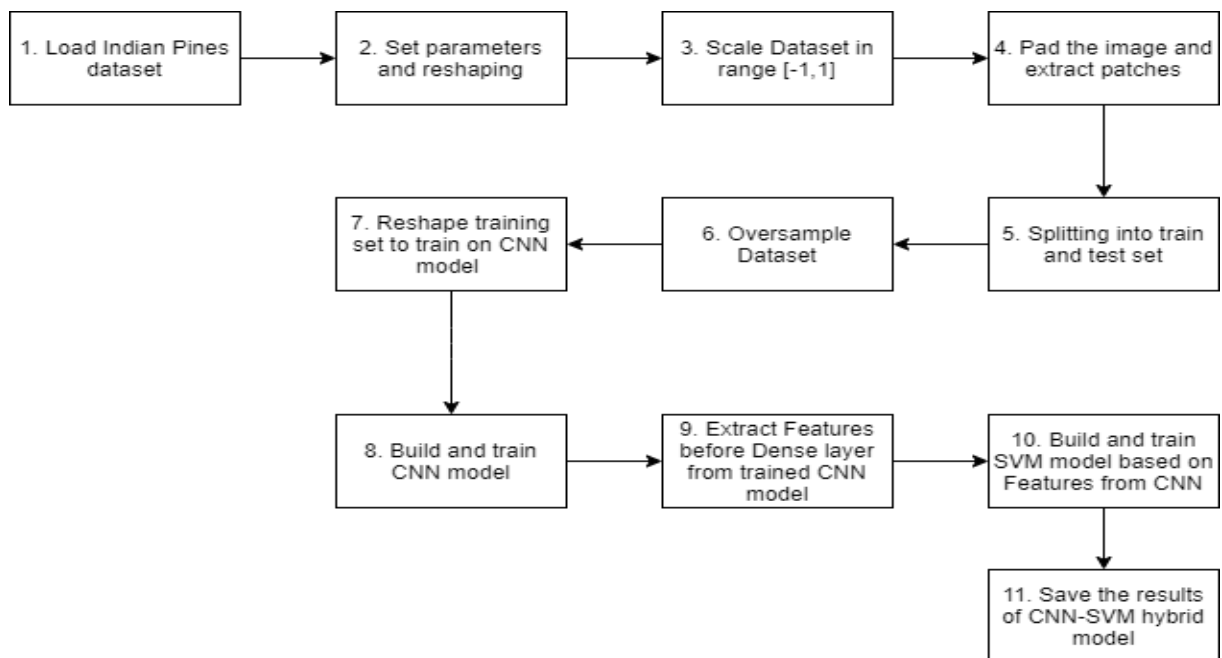


Fig: - Flow chart of CNN-SVM hybrid model

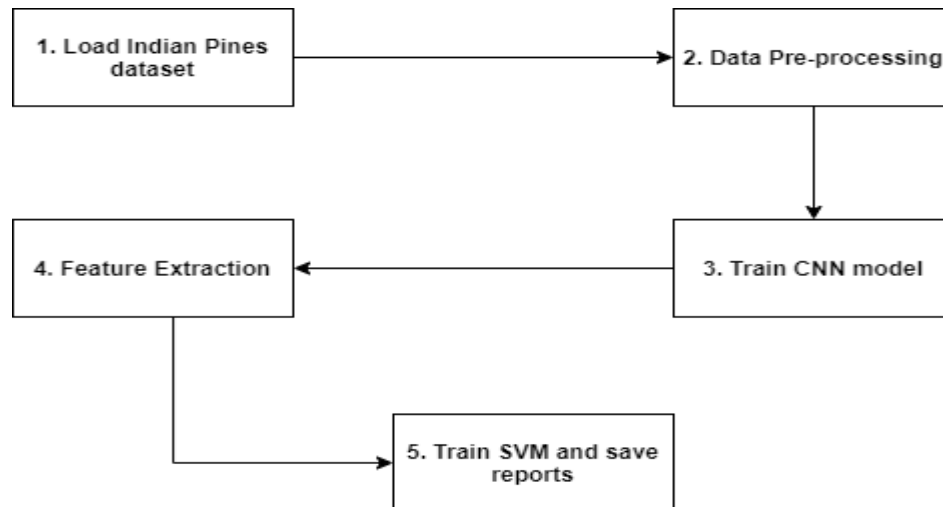


Fig: - Brief of CNN-SVM hybrid

In this project, a hybrid model of CNN and SVM is used. First, a CNN model is built and trains then features are extracted, and then the SVM model is trained. The steps can be seen in the above flowchart. These steps are divided into 5 parts:

- 1) **Import dataset:** In this step, the Indian pines dataset is loaded to train and built our model.
- 2) **Data pre-processing:**
  1. **Defining parameters:** In this step, parameters are decided to pre-process the dataset. Like height, width, no. of bands, oversampling, etc.  
Parameters used are: height=145, width=145, bands=200
  2. **Reshaping Dataset:** In this step, the dataset is reshaped and converted into a NumPy array. All other steps are performed on this array.
  3. **Scaling:** Above created NumPy array is scaled into a range of  $[-1, 1]$  because a hyperspectral image contains more than one band and each band may have a different range of values.
  4. **Pad the image:** In this step, the image is padded in all sides with values. These values can be zero or cornered values. This is done because the patch-size can be greater than 1. And to extract patch corresponding to each pixel padding must be done.
  5. **Extract Patches:** Patches are extracted and for each patch, its ground-truth value is matched with the center pixel of the patch.

Ex: Let the patch-size=3 x 3

Then after padding shape will be  $146 \times 146$

So, in  $padded_x[x][y]$ ;  $0 \leq x < 3 \wedge 0 \leq y < 3$

$padded_x[1][1]$  Corresponds to  $ground_{truth}[0][0]$

6. **Splitting into train and test set:** After extracting patches the dataset is divided into training and test set. The distributed dataset contains only those pixels for which ground-truth value is available. The total no of pixels with ground truth available is 10249. The no samples in the training set are 7686 and no. of samples in the test set is 2563 with test ratio is 0.25.
7. **Oversampling:** This is done if no of samples for a certain class is much lower than the max no of samples of any class. Then to reduce the difference between no. of samples of classes oversampling is done. This process creates new patches by making copies, rotate, etc. previous samples.

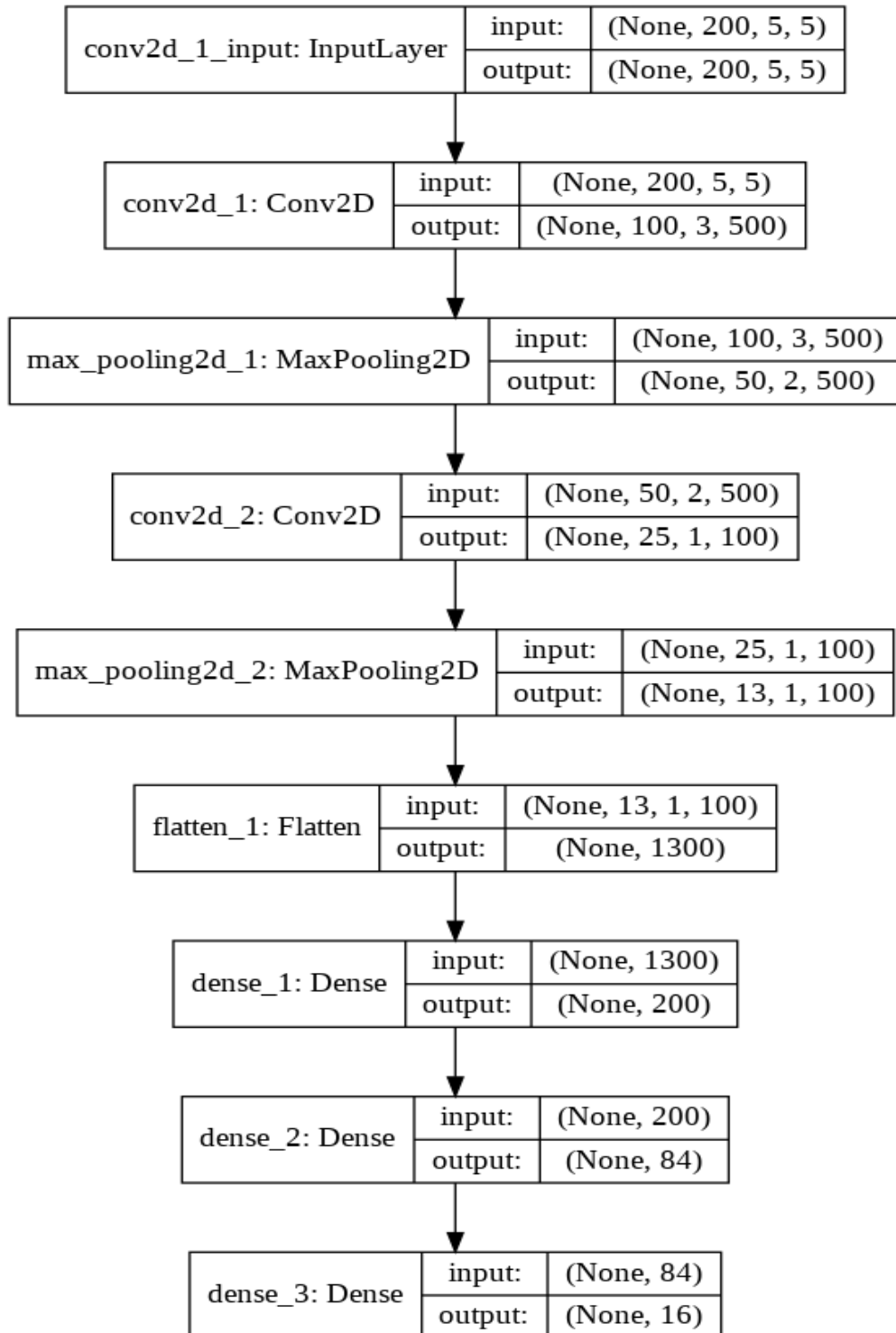
### 3) Build and Train CNN Model

1. **Building CNN model:** In this step, the CNN model is built. This model is built because recent works prove that CNN results in a greater change in accuracy than SVM. In this project, CNN is used just only for feature extraction.
2. **Training CNN model:** After the Successful building of the CNN model its training is done on the dataset. This training is done for feature extraction.
- 4) **Feature Extraction:** In this step features from the layer before fully-connected are extracted. This step is done because when our model trains on CNN it selected the best features by itself that can result in a better classification when passed to MLP (Multi-level Perceptron). So, the best features selected by CNN can also be passed to SVM rather than MLP.
- 5) **Build and train the SVM model:** Features from CNN can be passed to any classifier. In this project, SVM is used as a classifier while CNN is used as a feature extractor. Based on the features from CNN, the SVM model is built and trained. This whole model is known as the CNN-SVM hybrid model.
- 6) **Save Results:** After Training is done various reports are saved.
  1. Accuracy of Only SVM on test samples
  2. Accuracy of Only CNN on test samples
  3. Accuracy of CNN-SVM hybrid model on test samples
  4. Overall Accuracy Comparison

After this step, it is found that the CNN-SVM hybrid results in greater accuracy than both of these models.



The overall CNN model architecture is:



## 9. Results and Observations

The training of the INDIAN PINES dataset on the hybrid model of CNN and SVM results better as compared to either of them. All the models are trained over 7686 samples with available ground truth and tested over 2563 samples and the predicted results are compared with ground truth values corresponding to these pixels. The accuracy can be seen as:

No. of train samples: 7686

No. of test samples: 2563

Accuracy on Test Samples Using Only CNN: 85.48575639724731 (%)

Accuracy on Test Samples Using Only SVM: 83.67479674796748 (%)

Accuracy on Test Samples Using CNN-SVM: 92.31369488880219 (%)

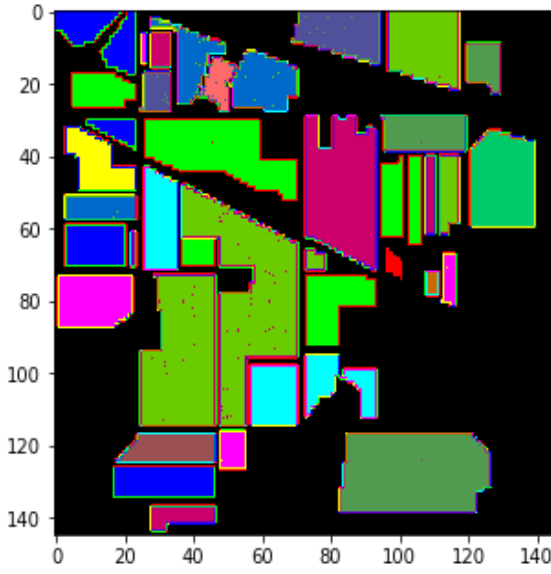
Classification result:				
	precision	recall	f1-score	support
0	1.00	1.00	1.00	11
1	0.88	0.88	0.88	357
2	0.87	0.95	0.91	208
3	0.95	1.00	0.98	59
4	1.00	0.97	0.98	121
5	0.99	0.99	0.99	183
6	0.70	1.00	0.82	7
7	0.99	1.00	1.00	120
8	1.00	1.00	1.00	5
9	0.84	0.95	0.89	243
10	0.95	0.85	0.90	614
11	0.89	0.93	0.91	148
12	1.00	1.00	1.00	51
13	0.98	0.94	0.96	316
14	0.81	0.95	0.88	97
15	1.00	1.00	1.00	23
accuracy			0.92	2563
macro avg	0.93	0.96	0.94	2563
weighted avg	0.93	0.92	0.92	2563

(a)

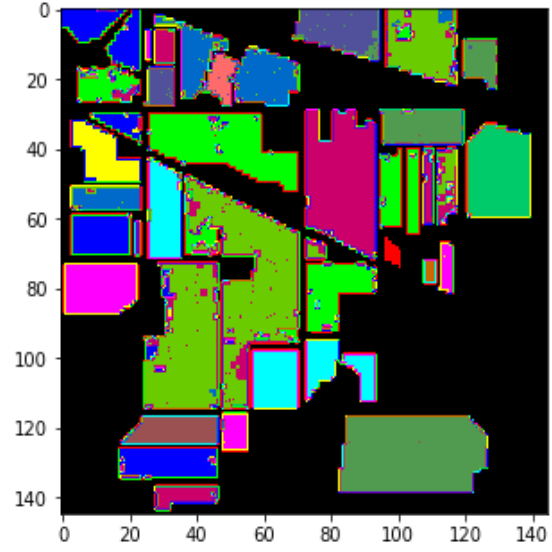
confusion matrix:														
[	[	11	0	0	0	0	0	0	0	0	0	0	0	0]
[	0	313	11	1	0	0	0	0	8	17	7	0	0	0]
[	0	0	198	0	0	0	0	0	3	4	3	0	0	0]
[	0	0	0	59	0	0	0	0	0	0	0	0	0	0]
[	0	0	0	0	117	0	3	1	0	0	0	0	0	0]
[	0	0	0	0	0	181	0	0	0	0	0	0	1	1]
[	0	0	0	0	0	0	7	0	0	0	0	0	0	0]
[	0	0	0	0	0	0	0	120	0	0	0	0	0	0]
[	0	0	0	0	0	0	0	0	5	0	0	0	0	0]
[	0	3	1	0	0	0	0	0	0	232	7	0	0	0]
[	0	34	16	2	0	1	0	0	32	522	7	0	0	0]
[	0	5	1	0	0	0	0	0	2	1	138	0	0	1]
[	0	0	0	0	0	0	0	0	0	0	0	51	0	0]
[	0	0	0	0	0	0	0	0	0	0	0	0	297	19]
[	0	0	0	0	0	0	0	0	0	0	0	0	5	92]
[	0	0	0	0	0	0	0	0	0	0	0	0	0	23]]]

(b)

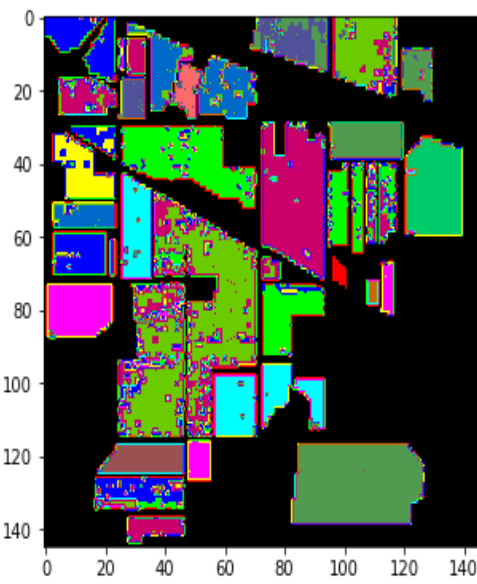
Fig: - (a) Classification report, (b) Confusion Matrix of CNN-SVM on test samples.



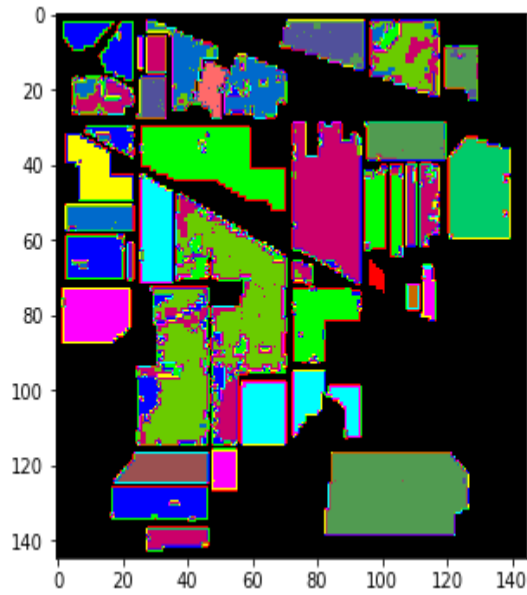
(a)



(b)



(c)



(d)

Fig: - (a) Indian pines ground truth, (b) predicted CNN-SVM hybrid image, (c) Predicted SVM image, (d) Predicted CNN image

From the above figures, it can be easily seen that CNN-SVM hybrid results in better accuracy than both of them.

## 10. References

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8. F. Melgani and L. Bruzzone, "Classification of hyperspectral remote sensing images with support vector machines," IEEE Transactions on Geoscience and Remote Sensing, vol. 42, no. 8, pp. 1778–1790, 2004, IEEE

## 11. Complete Source Code

- Github Link:- [https://github.com/darkshadow013/IP\\_ML](https://github.com/darkshadow013/IP_ML)
- Training SVM on indian pines:-
  1. IP\_SVM.ipynb:-  
[https://github.com/darkshadow013/IP\\_ML/blob/master/INDIAN\\_PINES\\_SVM\\_TRY/IP\\_SVM.ipynb](https://github.com/darkshadow013/IP_ML/blob/master/INDIAN_PINES_SVM_TRY/IP_SVM.ipynb)
- Training CNN on indian pines:-
  1. IP\_CNN\_3200\_epoch100.ipynb:-  
[https://github.com/darkshadow013/IP\\_ML/blob/master/INDIAN\\_PINES\\_CNN\\_TRY/IP\\_CNN\\_3200\\_epoch100.ipynb](https://github.com/darkshadow013/IP_ML/blob/master/INDIAN_PINES_CNN_TRY/IP_CNN_3200_epoch100.ipynb)
- Training CNN-SVM Hybrid on Indian Pines :-
  1. IP\_CNN\_SVM\_HYBRID.ipynb :-  
[https://github.com/darkshadow013/IP\\_ML/blob/master/IP\\_CNN\\_SVM/IP\\_CNN\\_SVM\\_HYBRID.ipynb](https://github.com/darkshadow013/IP_ML/blob/master/IP_CNN_SVM/IP_CNN_SVM_HYBRID.ipynb)
- Results of hybrid model on Indian Pines :-  
[https://github.com/darkshadow013/IP\\_ML/tree/master/IP\\_CNN\\_SVM/Files\\_CNN](https://github.com/darkshadow013/IP_ML/tree/master/IP_CNN_SVM/Files_CNN)

### IP\_SVM.py :-

```
#!/usr/bin/env python
# coding: utf-8

# # **This project is done on Google Colab**
# ### **Note :- To run this project on platform other than colab change the directories
# locations in this project**

# # **1. Load the data from Github**

# In[1]:

get_ipython().system('git clone -l -s git://github.com/darkshadow013/Land-cover-ML
himanshu-garg')
get_ipython().run_line_magic('cd', 'himanshu-garg')
get_ipython().system('ls')
```

```

# # **2. Data Pre-processing**

# In[2]:

get_ipython().system('pip install spectral')
from spectral import *
import scipy.io as si      # for inputing matlab files
import numpy as np        # Linear Algebra tools
from random import shuffle # for shuffling dataset
import pandas as pd       # for csv files and dataframes
import matplotlib.pyplot as plt #for plotting graphs
from sklearn.model_selection import train_test_split
from sklearn.metrics import confusion_matrix
from sklearn.metrics import accuracy_score
from sklearn.metrics import classification_report
from sklearn import svm
import time
import os

# In[ ]:

mat_x = si.loadmat('/content/himanshu-garg/Data/Indian_pines_corrected.mat')
['indian_pines_corrected'] # shape 145*145*220
mat_y = si.loadmat('/content/himanshu-garg/Data/Indian_pines_gt.mat')['indian_pines_gt']
# shape 145*145

# In[ ]:

h, w, b = 145, 145, 200 # height,width and band of image
P_S = 1 # patch size (window size)
oversampling = 1 # Turn true to make distribution of samples across various classes
to be uniform
no_of_patch = 200 # if oversampling is true enter no of patches for each class

# In[ ]:

x_np, y_np = np.array(mat_x), np.array(mat_y)
x, y = x_np.reshape(h, w, b), y_np.reshape(h, w) # x and y are equal to x_np and y_np

# In[ ]:

x = x.astype(float)
for i in range(b):
    x[:, :, i] = x[:, :, i] / np.max(x[:, :, i])

```

```
# In[ ]:
```

```
pad_width=int((P_S-1)/2)
padded_x=np.pad(x,[(pad_width,pad_width),(pad_width,pad_width),(0,0)],'edge')
```

```
# In[ ]:
```

```
X,Y=[],[]
for i in range(h):
    for j in range(w):
        if(y[i][j]!=0):
            patch=padded_x[i:i+P_S,j:j+P_S,:]
            X.append(patch)
            Y.append(y[i][j]-1)
```

```
# In[ ]:
```

```
x_train,x_test,y_train,y_test=train_test_split(X,Y,test_size=0.3,stratify=Y)
```

```
# In[12]:
```

```
u_c=np.unique(y_train,return_counts=True)
print(u_c)
plt.bar(u_c[0],u_c[1],align='center',alpha=0.5)
plt.title('Distribution of samples before resampling')
plt.show()
```

```
# In[13]:
```

```
output_classes=len(np.unique(y_np))-1
classes=[]
if(oversampling):
    for i in range(output_classes):
        classes.append([])

    for i in range(len(y_train)):
        classes[y_train[i]].append(x_train[i])

    for c in range(output_classes):
        temp=classes[c]
        for i in range(int(no_of_patch/len(classes[c]))):
            classes[c]+=temp
        shuffle(classes[c])
```

```

        classes[c] = classes[c][0:no_of_patch]

x_train,y_train= [], []
for c in range(output_classes):
    x_train.extend( classes[c] )
    for i in range(len(classes[c])):
        y_train.append(c)

randomize=np.arange(len(x_train))
np.random.shuffle(randomize)
x_train=[x_train[i] for i in randomize]
y_train=[y_train[i] for i in randomize]

u_c=np.unique(y_train,return_counts=True)
print(u_c)
plt.bar(u_c[0],u_c[1], align='center', alpha=0.5)
plt.title('Distribution of samples after resampling')
plt.show()

# In[ ]:

x_train,x_test,y_train,y_test=
np.array(x_train),np.array(x_test),np.array(y_train),np.array(y_test)

# In[ ]:

patch1=np.zeros((x_train.shape[0],P_S*P_S*b))
patch2=np.zeros((x_test.shape[0],P_S*P_S*b))

# In[ ]:

for i in range(patch1.shape[0]):
    patch1[i,:]=x_train[i,:,:,:].flatten('C')
for i in range(patch2.shape[0]):
    patch2[i,:]=x_test[i,:,:,:].flatten('C')

x_train=patch1
x_test=patch2

# # **3. Training Begins**

# In[3]:

import pickle
if(os.path.exists('/content/himanshu-garg/IP_SVM_BEST.sav')):
    # load the model

```



```

filename = '/content/himanshu-garg/IP_SVM_BEST.sav'
clf_best = pickle.load(open(filename, 'rb'))
print('The optimal value of accuracy is: {}'.format(clf.score(x_test, y_test)*100))
else:
    sigma = [0.01, 0.03, 0.1, 0.3, 1, 3, 10, 30]
    C = [0.01, 0.03, 0.1, 0.3, 1, 3, 10, 30]
    sigma = [0.3]
    C = [30]
    acc = list()
    sigma_c = list()
    max_acc = 0.0
    for each in sigma:
        for each_c in C:
            start = time.time()
            clf = svm.SVC(kernel='rbf', gamma = 1/(2*(each**2)), C=each_c,
decision_function_shape='ovr')
            clf.fit(x_train, y_train)
            accuracy = clf.score(x_test, y_test)
            acc.append(accuracy)
            sigma_c.append((each, each_c))
            print('Sigma: %f, C: %f, accuracy: %f, Time: %f sec' % (each, each_c, accuracy,
time.time()-start))
            index = np.argmax(acc)
            sigma_max, c_max = sigma_c[index]
            print('The optimal value of sigma is: {}'.format(sigma_max))
            print('The optimal value of C is: {}'.format(c_max))
            print('The optimal value of accuracy is: {}'.format(acc[index]))

```

# In[4]:

```

sigma = 1
gamma = 1/(2 * sigma**2)
print(gamma)

```

# In[ ]:

```

y_pred = clf_best.predict(x_test)

```

# In[28]:

```

results = confusion_matrix(y_test, y_pred)
print('Confusion Matrix :')
print(results)
print('Accuracy Score :', accuracy_score(y_test, y_pred))
print('Report : ')
print(classification_report(y_test, y_pred))

```

# # \*\*4. Predict Results\*\*

# In[ ]:

```

X1=np.array(X)
X2=np.zeros((X1.shape[0],P_S*P_S*b))
for i in range(X2.shape[0]):
    X2[i,:]=X1[i,:,:,:].flatten('C')

# In[ ]:

# Function to extract patche at h_index,w_index

def patch_at_index(h_index,w_index):
    patch=padded_x[h_index:h_index+P_S,w_index:w_index+P_S,:]
    patch=patch.flatten('C')
    patch=patch.reshape(1,patch.shape[0])
    return patch

# In[ ]:

y_hat=np.zeros((h,w),dtype=int)

for i in range(h):
    for j in range(w):
        if(y[i][j]):
            patch=patch_at_index(i,j)
            y_hat[i][j]=clf_best.predict(patch)+1
        else:
            y_hat[i][j]=0

# In[32]:

ground_truth=imshow(classes=y,figsize=(5,5))
predicted_image=imshow(classes=y_hat,figsize=(5,5))

```

```

# In[ ]:

```

### IP\_CNN\_3200\_epoch100.py :-

```

#!/usr/bin/env python
# coding: utf-8

# In[1]:

get_ipython().system('git clone -l -s git://github.com/darkshadow013/Land-cover-ML
himanshu-garg')
get_ipython().run_line_magic('cd', 'himanshu-garg')

```

```

get_ipython().system('ls')

# In[2]:

# Import the necessary libraries
get_ipython().system('pip install spectral')
from spectral import *
import scipy
import scipy.io as sio
import scipy.ndimage
from sklearn.model_selection import train_test_split, StratifiedShuffleSplit
from sklearn import preprocessing
from sklearn.metrics import classification_report, confusion_matrix
from sklearn.decomposition import PCA
from skimage.transform import rotate
#from spectral import *
from keras.models import Sequential, load_model
from keras.layers import Dense, Dropout, Flatten, Conv2D, MaxPooling2D
from keras.optimizers import SGD
from keras.callbacks import ReduceLROnPlateau, ModelCheckpoint
from keras import backend as K
from keras.utils import np_utils
import itertools
import os
import random
from random import shuffle
import numpy as np
import matplotlib.pyplot as plt

# In[ ]:

mat_x = sio.loadmat('/content/himanshu-garg/Data/Indian_pines_corrected.mat')
['indian_pines_corrected']      # shape 145*145*220
mat_y = sio.loadmat('/content/himanshu-garg/Data/Indian_pines_gt.mat')['indian_pines_gt']
# shape 145*145

# In[ ]:

h, w, b = 145, 145, 200      # height,width and band of image
P_S = 5                      # patch size (window size)
oversampling = 1             # Turn true to make distribution of samples across various classes
                              # to be uniform
no_of_patch = 200            # if oversampling is true enter no of patches for each class

# In[ ]:

x_np, y_np = np.array(mat_x), np.array(mat_y)
x, y = x_np.reshape(h,w,b), y_np.reshape(h,w)      # x and y are equal to x_np and y_np

# In[ ]:

x = x.astype(float)

```

```

for i in range(b):
    x[:, :, i] = x[:, :, i] / np.max(x[:, :, i])

# In[ ]:

pad_width = int( (P_S-1)/2 )
padded_x = np.pad(x, [(pad_width,pad_width), (pad_width,pad_width)], (0,0), 'edge')

# In[ ]:

X,Y = [], []
for i in range(h):
    for j in range(w):
        if(y[i][j] != 0):
            patch = padded_x[i:i+P_S, j:j+P_S, :]
            X.append(patch)
            Y.append(y[i][j]-1)

# In[ ]:

x_train, x_test, y_train, y_test = train_test_split( X, Y, test_size=0.25, stratify=Y,
random_state=10)

# In[11]:

u_c = np.unique(y_train, return_counts=True)
print(u_c)
plt.bar( u_c[0], u_c[1], align='center', alpha=0.5)
plt.title('Distribution of samples before resampling')
plt.show()

# In[12]:

output_classes = len(np.unique(y_np))-1
classes = []
if(oversampling):
    for i in range(output_classes):
        classes.append([])

    for i in range(len(y_train)):
        classes[y_train[i]].append(x_train[i])

    for c in range(output_classes):
        temp = classes[c]
        for i in range( int( no_of_patch / len(classes[c]) ) ):
            classes[c] += temp
            shuffle(classes[c])
            classes[c] = classes[c][0:no_of_patch]

    x_train, y_train = [], []
    for c in range(output_classes):

```

```

        x_train.extend( classes[c] )
        for i in range(len(classes[c])):
            y_train.append(c)

    randomize = np.arange(len(x_train))
    np.random.shuffle(randomize)
    x_train = [x_train[i] for i in randomize]
    y_train = [y_train[i] for i in randomize]

    u_c = np.unique(y_train,return_counts=True)
    print(u_c)
    plt.bar( u_c[0],u_c[1], align='center', alpha=0.5)
    plt.title('Distribution of samples after resampling')
    plt.show()

# In[ ]:

x_train,x_test,y_train,y_test =
np.array(x_train),np.array(x_test),np.array(y_train),np.array(y_test)

# In[14]:

BAND = 200
BATCH_SIZE = 100
LEARNING_RATE = 0.01
CONV_1_CHANNELS = 500
CONV_2_CHANNELS = 100
FC_1_UNITS = 200
FC_2_UNITS = 84
FC_3_UNITS = 16

#Kernel Sizes
CONV_K_S = 5
POOL_K_S = 2

STRIDES = 2
x_train = np.reshape(x_train, (x_train.shape[0], x_train.shape[3],
                               x_train.shape[1], x_train.shape[2]))
x_test = np.reshape(x_test, (x_test.shape[0],x_test.shape[3],
                             x_test.shape[1], x_test.shape[2]))
y_train = np_utils.to_categorical(y_train)
y_test = np_utils.to_categorical(y_test)
print(x_train.shape[0],x_train.shape[1],x_train.shape[2],x_train.shape[3])

# In[ ]:

input_shape= x_train[0].shape

# In[ ]:

# Define the model structure
model = Sequential()

```

```

model.add(Conv2D(500, (3,3) , activation='relu', input_shape=input_shape, strides = STRIDES,
padding = 'same'))
model.add(MaxPooling2D(pool_size = POOL_K_S, padding = 'same', strides = STRIDES))
model.add(Conv2D(100, (3,3) , activation='relu',strides = STRIDES, padding = 'same'))
model.add(MaxPooling2D(pool_size = POOL_K_S, padding = 'same', strides = STRIDES))
model.add(Flatten())
model.add(Dense(200, activation='relu'))
model.add(Dense(84, activation='relu'))
model.add(Dense(16,activation='softmax'))

# In[ ]:

reduce_lr = ReduceLROnPlateau(monitor='val_acc', factor=0.9, patience=25,
min_lr=0.000001, verbose=1)
checkpointer = ModelCheckpoint(filepath="checkpoint.hdf5", verbose=1,
save_best_only=True)
sgd = SGD(lr=0.001, decay=1e-6, momentum=0.9, nesterov=True)
model.compile(loss='categorical_crossentropy', optimizer=sgd,
metrics=['accuracy'])

# In[19]:

from keras.utils import plot_model
if not os.path.exists('/content/Files_CNN'):
    os.makedirs('/content/Files_CNN')
plot_model(model, to_file='/content/Files_CNN/CNN_model.png', show_shapes=True)

# In[ ]:

if(os.path.exists('/content/himanshu-garg/IP_CNN_samples3200_epoch200.h5')):
    # load the model architecture and weights
    model.load_weights("/content/himanshu-garg/IP_CNN_samples3200_epoch200.h5")
else:
    # Start to train model
    history = model.fit(x_train, y_train,
        batch_size=32,
        epochs=200,
        verbose=1,
        validation_data=(x_test, y_test),
        callbacks=[reduce_lr, checkpointer],
        shuffle=True)
    # save the model with h5py
    import h5py
    model.save('/content/himanshu-garg/IP_CNN_samples3200_epoch200.h5')

# In[21]:

score = model.evaluate(x_test, y_test, batch_size=32)

# In[22]:

Test_Loss = score[0]*100

```

```

Test_accuracy = score[1]*100
print(Test_Loss,Test_accuracy)

# In[ ]:

height = y.shape[0]
width = y.shape[1]
PATCH_SIZE = 5
numComponents = 30

def Patch(data,height_index,width_index):
    height_slice = slice(height_index, height_index+PATCH_SIZE)
    width_slice = slice(width_index, width_index+PATCH_SIZE)
    patch = data[height_slice, width_slice, :]
    return patch

# calculate the predicted image
outputs = np.zeros((height,width))
for i in range(height-PATCH_SIZE+1):
    for j in range(width-PATCH_SIZE+1):
        p = int(PATCH_SIZE/2)
        target = y[i+p][j+p]
        if target == 0 :
            continue
        else :
            image_patch=Patch(x,i,j)
            X_test_image =
image_patch.reshape(1,image_patch.shape[2],image_patch.shape[0],image_patch.shape[1]).astype
pe('float32')
            prediction = (model.predict_classes(X_test_image))
            outputs[i+p][j+p] = prediction+1

# In[25]:

predict_image = imshow(classes=outputs.astype(int), figsize=(5, 5))

```

### IP\_CNN\_SVM\_HYBRID.py :-

```

#!/usr/bin/env python
# coding: utf-8

# # **This project is done on Google Colab**
# ### **Note :- To run this project on platform other than colab change the directories
locations in this project**

# # **1. Load the data from Github**

# In[119]:

get_ipython().system('git clone -l -s git://github.com/darkshadow013/Land-cover-ML
himanshu-garg')
get_ipython().run_line_magic('cd', 'himanshu-garg')
get_ipython().system('ls')

# # **2. Data Pre-processing**

```

```

# In[ ]:

# Import the necessary libraries
get_ipython().system('pip install spectral')
from spectral import *
import scipy
import scipy.io as sio
import scipy.ndimage
from sklearn.model_selection import train_test_split, StratifiedShuffleSplit
from sklearn import preprocessing
from sklearn.metrics import classification_report, confusion_matrix, accuracy_score
from sklearn.decomposition import PCA
from skimage.transform import rotate
#from spectral import *
from keras.models import Sequential, load_model
from keras.layers import Dense, Dropout, Flatten, Conv2D, MaxPooling2D
from keras.optimizers import SGD
from keras.callbacks import ReduceLROnPlateau, ModelCheckpoint
from keras import backend as K
from keras.utils import np_utils
import itertools
import os
import random
from random import shuffle
import numpy as np
import matplotlib.pyplot as plt
from sklearn.svm import SVC
from sklearn.model_selection import GridSearchCV
import parameters as p
import pickle

# In[ ]:

#Files_CNN is used to store the reports of this project
if not os.path.exists('/content/Files_CNN'):
    os.makedirs('/content/Files_CNN')

# In[ ]:

mat_x = sio.loadmat('/content/himanshu-garg/Data/Indian_pines_corrected.mat')
['indian_pines_corrected'] # shape 145*145*220
mat_y = sio.loadmat('/content/himanshu-garg/Data/Indian_pines_gt.mat')['indian_pines_gt']
# shape 145*145

# In[ ]:

h, w, b = p.h, p.w, p.b # height,width and band of image
patch_size = p.patch_size # patch size (window size)
oversampling = p.oversampling # Turn true to make distribution of samples across
various classes to be uniform
no_of_patch = p.no_of_patch # if oversampling is true enter no of patches for each
class

target_names = ['Alfalfa', 'Corn-notill', 'Corn-mintill', 'Corn']

```



```

        'Grass-pasture', 'Grass-trees', 'Grass-pasture-mowed',
        'Hay-windrowed', 'Oats', 'Soybean-notill', 'Soybean-mintill',
        'Soybean-clean', 'Wheat', 'Woods', 'Buildings-Grass-Trees-Drives',
        'Stone-Steel-Towers']

# In[ ]:

x_np, y_np = np.array(mat_x), np.array(mat_y)
x, y = x_np.reshape(h,w,b), y_np.reshape(h,w)    # x and y are equal to x_np and y_np

# In[ ]:

def normalize_x(x):
    x = x.astype(float)
    for i in range(b):
        x[:, :, i] = x[:, :, i] / np.max(x[:, :, i])
    return x

# In[ ]:

def create_padded_x(x, patch_size):
    pad_width = int((patch_size-1)/2)
    padded_x = np.pad(x, [(pad_width, pad_width), (pad_width, pad_width), (0, 0)], 'edge')
    return padded_x

# In[ ]:

x = normalize_x(x)

# In[ ]:

padded_x = create_padded_x(x, patch_size)

# In[ ]:

def ground_truth_patches_labels(h, w, padded_x, patch_size):
    X, Y = [], []
    for i in range(h):
        for j in range(w):
            if (y[i][j] != 0):
                patch = padded_x[i:i+patch_size, j:j+patch_size, :]
                X.append(patch)
                Y.append(y[i][j]-1)
    return X, Y

# In[ ]:

X, Y = ground_truth_patches_labels(h, w, padded_x, patch_size)

```

```

# In[ ]:

x_train, x_test, y_train, y_test = train_test_split(X, Y, test_size=0.25, stratify=Y,
random_state=10)

# In[132]:

u_c = np.unique(y_train, return_counts=True)
print(u_c)
plt.bar(u_c[0], u_c[1], align='center', alpha=0.5)
plt.title('Distribution of samples before resampling')
plt.show()

# In[133]:

output_classes = len(np.unique(y_np)) - 1
classes = []
if(oversampling):
    for i in range(output_classes):
        classes.append([])

    for i in range(len(y_train)):
        classes[y_train[i]].append(x_train[i])

    for c in range(output_classes):
        temp = classes[c]
        for i in range(int(no_of_patch / len(classes[c]))):
            classes[c] += temp
        shuffle(classes[c])
        classes[c] = classes[c][0:no_of_patch]

    x_train, y_train = [], []
    for c in range(output_classes):
        x_train.extend(classes[c])
        for i in range(len(classes[c])):
            y_train.append(c)

    randomize = np.arange(len(x_train))
    np.random.shuffle(randomize)
    x_train = [x_train[i] for i in randomize]
    y_train = [y_train[i] for i in randomize]

    u_c = np.unique(y_train, return_counts=True)
    print(u_c)
    plt.bar(u_c[0], u_c[1], align='center', alpha=0.5)
    plt.title('Distribution of samples after resampling')
    plt.show()

# In[ ]:

```

```

x_train,x_test,y_train,y_test=
np.array(x_train),np.array(x_test),np.array(y_train),np.array(y_test)

# # **3. Build and train CNN model on DATASET**

# In[ ]:

CONV_1_CHANNELS=p.CONV_1_CHANNELS
CONV_2_CHANNELS=p.CONV_2_CHANNELS
FC_1_UNITS=200
FC_2_UNITS=84
FC_3_UNITS=16

#Kernel Sizes
CONV_K_S=5
POOL_K_S=2

STRIDES=2
# Before x_train.shape= {number,height,width,bands}
# Aftet x_train.shape= {number,bands,height,width}
x_train=np.reshape(x_train, (x_train.shape[0], x_train.shape[3],
                             x_train.shape[1], x_train.shape[2]))
x_test=np.reshape(x_test, (x_test.shape[0],x_test.shape[3],
                            x_test.shape[1], x_test.shape[2]))
p1=y_train
q1=y_test
y_train=np_utils.to_categorical(y_train)
y_test=np_utils.to_categorical(y_test)
a1=x_test
b1=y_test

# In[ ]:

input_shape=x_train[0].shape

# In[ ]:

# Define the model structure
model=Sequential()
model.add(Conv2D(500, (3,3) , activation='relu', input_shape=input_shape, strides=STRIDES,
padding= 'same'))
model.add(MaxPooling2D(pool_size= POOL_K_S, padding= 'same', strides= STRIDES))
model.add(Conv2D(100, (3,3) , activation='relu',strides= STRIDES, padding= 'same'))
model.add(MaxPooling2D(pool_size= POOL_K_S, padding= 'same', strides= STRIDES))
model.add(Flatten())
model.add(Dense(200, activation='relu'))
model.add(Dense(84, activation='relu'))
model.add(Dense(16,activation='softmax'))

# In[ ]:

reduce_lr=ReduceLROnPlateau(monитор='val_acc', factor=0.9, patience=25,
                             min_lr=0.000001, verbose=1)
checkpointer=ModelCheckpoint(filepath="checkpoint.hdf5", verbose=1,

```

```

        save_best_only=True)
sgd = SGD(lr=0.001, decay=1e-6, momentum=0.9, nesterov=True)
model.compile(loss='categorical_crossentropy', optimizer=sgd,
              metrics=['accuracy'])

# In[139]:

from keras.utils import plot_model
plot_model(model, to_file='/content/Files_CNN/CNN_model.png', show_shapes=True)

# In[140]:

if(os.path.exists('/content/himanshu-garg/IP_CNN_samples3200_epoch200.h5')):
    # load the model architecture and weights
    model.load_weights("/content/himanshu-garg/IP_CNN_samples3200_epoch200.h5")
else:
    # Start to train model
    history = model.fit(x_train, y_train,
                      batch_size=32,
                      epochs=200,
                      verbose=1,
                      validation_data=(x_test, y_test),
                      callbacks=[reduce_lr, checkpointer],
                      shuffle=True)
    # save the model with h5py
    import h5py
    model.save('/content/himanshu-garg/IP_CNN_samples3200_epoch200.h5')

for l in range(len(model.layers)):
    print(l, model.layers[l])

# # **4. Feature Extraction from second last layer of CNN**

# In[ ]:

# feature extraction layer
getFeature = K.function([model.layers[0].input, K.learning_phase()],
                       [model.layers[6].output])
# classification layer
getPrediction = K.function([model.layers[7].input, K.learning_phase()],
                           [model.layers[7].output])

# In[ ]:

exTrain3000 = getFeature([x_train, 0])[0]
exTest1000 = getFeature([x_test, 0])[0]

# In[ ]:

y_train=p1
y_test=q1
y_train3000 = y_train

```

```

y_test1000=y_test

# In[144]:

print(exTrain3000.shape, exTest1000.shape, y_train3000.shape, y_test1000.shape)

# # **5. Build and Train SVM based on Features from CNN**

# In[ ]:

if(os.path.exists('/content/himanshu-garg/IP_CNN_SVM_BEST.sav')):
    # load the model
    filename = '/content/himanshu-garg/IP_CNN_SVM_BEST.sav'
    svmclf=pickle.load(open(filename, 'rb'))
else:
    C = [0.01, 0.03, 0.1, 0.3, 1, 3, 10, 30]
    sigma = [0.01, 0.03, 0.1, 0.3, 1, 3, 10, 30]
    gamma=[]
    for i in range(0,len(sigma)):
        gamma.append((1/(2 * sigma[i]**2)))
    """parameters = {'kernel':['rbf'],
                    'C':C,
                    'gamma':gamma}"""
    parameters = {'kernel':['rbf'],
                  'C':[30],
                  'gamma':[0.005]}
    clf=GridSearchCV(SVC(), parameters)
    clf.fit(exTrain3000, y_train3000)
    #By this we obtain optimal C=30, gamma=0.005

    svmclf=clf.best_estimator_
    svmclf.fit(exTrain3000, y_train3000)

    filename = '/content/himanshu-garg/IP_CNN_SVM_BEST.sav'
    pickle.dump(svmclf, open(filename, 'wb'))

# # **6. Save the reports**

# In[ ]:

def save_report_of_CNN_on_test_samples(y_test, test_accuracy):
    file_name = "/content/Files_CNN/Report_cnn_on_TEST_SAMPLES.txt"
    with open(file_name, 'w') as x_file:
        x_file.write('Number of Test Samples : {} '.format(y_test.shape[0]))
        x_file.write('\n')
        x_file.write('Test accuracy : {} (%)'.format(test_accuracy))

# In[ ]:

def
save_report_of_CNN_SVM_on_train_samples(y_train3000, train_accuracy, classification, confusio
n):
    file_name = "/content/Files_CNN/Report_cnn_svm_on_TRAIN_SAMPLES.txt"
    with open(file_name, 'w') as x_file:

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x_file.write('Number of train Samples : {} '.format(y_train3000.shape[0]))
x_file.write('\n')
x_file.write('Train accuracy : {} (%)'.format(train_accuracy))
x_file.write('\n')
x_file.write('\n')
x_file.write("Classification result: \n")
x_file.write('{} '.format(classification))
x_file.write('\n')
x_file.write(" confusion matrix: \n")
x_file.write('{} '.format(confusion))

# In[ ]:

def
save_report_of_CNN_SVM_on_test_samples(y_test1000,Test_accuracy,classification,confusion):
    file_name = "/content/Files_CNN/Report_cnn_svm_on_TEST_SAMPLES.txt"
    with open(file_name, 'w') as x_file:
        x_file.write('Number of Test Samples : {} '.format(y_test1000.shape[0]))
        x_file.write('\n')
        x_file.write('Test accuracy : {} (%)'.format(Test_accuracy))
        x_file.write('\n')
        x_file.write('\n')
        x_file.write("Classification result: \n")
        x_file.write('{} '.format(classification))
        x_file.write('\n')
        x_file.write(" confusion matrix: \n")
        x_file.write('{} '.format(confusion))

# In[ ]:

def
save_overall_comparision_of_models(Y,h,w,y_train,y_test,test_accuracy,train_accuracy,Test_
accuracy):
    file_name = "/content/Files_CNN/Overall_Accuracy_Comparision.txt"
    with open(file_name, 'w') as x_file:
        x_file.write('\n\t1. Number of Pixels with Ground Truth: {}
'.format(np.array(Y).shape[0]))
        x_file.write('\n\n')
        x_file.write('\t2. Number of Pixels without Ground Truth: {} '.format(h*w -
np.array(Y).shape[0]))
        x_file.write('\n\n')
        x_file.write('\t3. Number of Train Samples : {} '.format(y_train.shape[0]))
        x_file.write('\n\n')
        x_file.write('\t4. Number of Test Samples : {} '.format(y_test.shape[0]))
        x_file.write('\n\n\n\
n-----OVERALL
COMPARISION OF ACCURACY-----\
n\n\n\n')
        x_file.write('\t1. Accuracy on Test Samples Using Only CNN :- {}
(%) '.format(test_accuracy))
        x_file.write('\n\n')
        x_file.write('\t2. Accuracy on Test Samples Using Only SVM :- {}
(%) '.format(83.67479674796748))
        x_file.write('\n\n')
        x_file.write('\t3. Accuracy on Train Samples Using CNN-SVM :- {}
(%) '.format(train_accuracy))
        x_file.write('\n\n')

```

```

        x_file.write('\t4. Accuracy on Test Samples Using CNN-SVM :- {}'.format(Test_accuracy))

# In[ ]:

y_testSVM=svmclf.predict(exTest1000)
confusion=confusion_matrix(y_test1000,y_testSVM)
Test_accuracy=100*accuracy_score(y_test1000,y_testSVM)
classification=classification_report(y_test1000,y_testSVM)

save_report_of_CNN_SVM_on_test_samples(y_test1000,Test_accuracy,classification,confusion)

# In[ ]:

y_trainSVM=svmclf.predict(exTrain3000)
confusion=confusion_matrix(y_train3000,y_trainSVM)
train_accuracy=100*accuracy_score(y_train3000,y_trainSVM)
classification=classification_report(y_train3000,y_trainSVM)

save_report_of_CNN_SVM_on_train_samples(y_train3000,train_accuracy,classification,confusion)

# In[152]:

model=load_model("/content/himanshu-garg/IP_CNN_samples3200_epoch200.h5")
score=model.evaluate(a1,b1,batch_size=32)
Test_Loss=score[0]*100
test_accuracy=score[1]*100

# In[ ]:

save_report_of_CNN_on_test_samples(y_test,test_accuracy)

# In[ ]:

save_overall_comparision_of_models(Y,h,w,y_train,y_test,test_accuracy,train_accuracy,Test_accuracy)

# # **7. Generate Predicted Image**

# In[ ]:

x_np,y_np=np.array(mat_x),np.array(mat_y)
x,y=x_np.reshape(h,w,b),y_np.reshape(h,w)

# In[ ]:

x=normalize_x(x)

```

```

# In[ ]:

padded_x=create_padded_x(x,patch_size)

# In[ ]:

X,Y=ground_truth_patches_labels(h,w,padded_x,patch_size)

# In[ ]:

X=np.array(X)
x_test=np.reshape(X, (X.shape[0],X.shape[3],
                      X.shape[1], X.shape[2]))

# In[160]:

print(x_test.shape)

# In[ ]:

pred=[]
for i in range(0,x_test.shape[0]):
    x_TEST=x_test[i]
    x_TEST=np.reshape(x_TEST, (1,x_test.shape[1],x_test.shape[2],x_test.shape[3]))
    x_TEST=getFeature([x_TEST, 0])[0]
    pre=svmclf.predict(x_TEST)
    pred.append(pre[0])

# In[ ]:

output=np.zeros((h,w))
k=0
for i in range(h):
    for j in range(w):
        if(y[i][j]!=0):
            output[i][j]=pred[k]+1
            k=k+1
        else:
            output[i][j]=y[i][j]

# In[172]:

ground_truth=imshow(classes=y,figsize=(5,5))

# In[175]:

```



```
predict_image = imshow(classes = output.astype(int), figsize = (5, 5))
```