

CROP RELATED PATTERN AND CROP DISEASE DETECTION USING DATA MINING AND DENSENET-121

A PROJECT REPORT

Submitted by
SAMUNDAR SINGH
(2019272033)

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DEPARTMENT OF INFORMATION SCIENCE AND TECHNOLOGY
COLLEGE OF ENGINEERING, GUINDY
ANNA UNIVERSITY
CHENNAI 600 025
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BONA FIDE CERTIFICATE

Certified that this project report titled CROP RELATED PATTERN AND CROP DISEASE DETECTION USING DATA MINING AND DENSENET-121 is the bona fide work of Samundar Singh who carried out project work under my supervision. Certified further that to the best of my knowledge and belief, the work reported herein does not form part of any other thesis or dissertation on the basis of which a degree or an award was conferred on an earlier occasion on this or any other candidate.

PLACE: CHENNAI

DATE:

DR. R. GEETHA RAMANI

PROFESSOR

PROJECT GUIDE

DEPARTMENT OF IST, CEG

ANNA UNIVERSITY

CHENNAI 600025

COUNTERSIGNED

DR. S. SRIDHAR

HEAD OF THE DEPARTMENT

DEPARTMENT OF INFORMATION SCIENCE AND TECHNOLOGY

COLLEGE OF ENGINEERING, GUINDY

ANNA UNIVERSITY

CHENNAI 600025

ABSTRACT

India is an agriculture-based economy where climate and other environmental changes have become a major threat in the agricultural field. Machine learning is an essential approach for achieving partial and effective solution for this problem. One of the significant issues in the industry is lacking an accurate way to identify the best crop that can be grown with the available soil fertility in a particular land, also If the crops are not feed the required amount of water, the crop yields are adversely affected, by over-irrigation or under-irrigation. Another aspect is identification of the plant diseases which is the key to preventing the losses in the yield and quantity of the agricultural product.

Crop produce prediction: This module predicts the produce of the crop on basis of area and location. There are 15 types of crops whose data is collected from 1966 to 2017 almost all the district of the India. In this module linear regression is model is used to predict the produce of crop on basis of area and location provided.

Fertilizer and Irrigation prediction: This module predict the various amount of fertilizer and amount of water to be used in irrigation. There are 13 crops whose data is collected from 1966 to 2017 almost all district of India. In this module after analysis of various model, the random forest module is used to predict both the fertilizer and irrigation

Plant Disease Prediction: This module aims to predict the plant disease on basis of the image of leaf. The proposed algorithm is to optimize the information from the resources available to us for the betterment of the result without any complexity. The neural network used for classification is the Dense Convolution Neural Network (DCNN). In this project, a pre-trained neural network model (densenet-121) has been used to classify the 29 different diseases for 7 plants (potato, tomato, corn, bell pepper, grape, apple and cherry). In this project, the original image is converted to HSV colour, greyscale colour and lab colour form and then the masked image is generated by thresholding and given to the proposed model for training and classification

In this project a Web application has been developed which consist of the above module. The tech stack used in this project are mentioned below:

- **Frontend:** Angular
- **Backend:** Nodejs, python
- **Libraries:** plotlyjs, scikitlearn, Pandas, Numpy.

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MCA (Self-Supporting)

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CHAPTER 1

1. INTRODUCTION

1.1 Crop related pattern (Fertilizer and irrigation):

In India there is need of advances in agriculture sector so that it meets the needs of people. agriculture is considered as the main and the foremost culture practiced in India. Nowadays, modern people don't have awareness about the cultivation of the crops in a right time and at a right place Because of these cultivating techniques the seasonal climatic conditions are also being changed against the fundamental assets like soil, water and air which lead to insecurity of food. Crop yield prediction is an important agricultural problem. Each and Every farmer is always trying to know, how much yield will get from his expectation. In the past, yield prediction was calculated by analysing farmer's previous experience on a particular crop. The Agricultural yield is primarily depending on weather conditions, pests and planning of harvest operation. Accurate information about history of crop yield is an important thing for making decisions related to agricultural risk management. The fertilizer and water irrigation plays a major role in the crop production.

1.2 Crop Disease:

Disease detection involves the steps like image acquisition, image pre-processing, image segmentation, feature extraction and classification. Disease on plants results in large reduction in both the standard and the quantity of agricultural products. The study of diseases asks the studies of visually observable patterns on the plants. In most of the cases, the symptoms of a plant disease are seen on the leaves and stem. The plant leaf for the detection of disease is taken into account which shows the disease symptoms. The disease of plant is also highly responsible for the produce of the crop.

1.3 Problem Statement:

The decrease in the production of crop due to lack of knowledge of correct amount of fertilizer that is needed to be used and proper arrangement of the irrigation facility.

Also due to lack of knowledge of plant disease the correct precaution is also not taken. There are possibilities that a particular crop may have more than one type of disease and sometimes the farmers missed this which result in low production of crops although he has taken the correct measure.

1.4 Objective:

The objective of this challenge is to build a machine learning algorithm to correctly classify if a crop is healthy, has stem rust, or has leaf rust with more accuracy.

The machine learning algorithm give accurate early stage prediction about diseases occurring in the crop so that mortality rate of crop yields can be improved. Disease development depends on some conditions like crop susceptibility to disease, viability of pathogen and environment favourability for that crop yield. After detecting the machine learning algorithm will suggest the best possible way to deal with the problem.

1.5 Project Overview

This project is basically dealing with the disease prediction and the crop pattern prediction system. The crop pattern is predicted on basis of the location and crop type. The user has to mention his state and district in order to find the amount of fertilizer and irrigation estimation. For fertilizer and irrigation prediction the harvesting area and location detail is taken from user and using Random forest model the output is predicted.

For plant disease prediction the image of leaf is taken from the user and using deep learning the disease is predicted. After the disease prediction It also suggest the remedy or cure for the predicted disease.

1.6 Organization of Report:

Chapter 2: This chapter explains about the literature survey made on the existing system, analysing the problem statements and issues with the existing system.

Chapter 3: This module consists System design of the project with its preliminary design and descriptive details about the modules.

Chapter 4: This module consists Algorithm and pseudocode related to the models with their outcomes.

CHAPTER 2

2. LITERATURE SURVEY/RELATED WORK

This Chapter explains about the literature survey made on the existing system, analysing the problem statements and issues with the existing system and proposed objectives for the new system. Role of Crop related pattern in produce and effect of disease in crop produce analysis has been conducted by many authors. Some of them are summarised here

2.1 Crop production related pattern.

In recent years, machine learning techniques have been applied in agriculture domain by various researchers. A review on the implementation of different ML techniques in the field of crop yield prediction from soil analysis from past few years is presented as under.

Bhuyar [1] proposed an approach where different classification algorithms such as J48, Naïve Bayes, and Random forest algorithm were applied to soil data set to predict its fertility. J48 algorithm gave better result with an accuracy of 98.17% than other algorithms. This analysis is done only to soil of a particular location.

Rajeshwari and Arunesh [2] performed a comparative analysis of ML algorithms i.e. Naive Bayes, JRIP and J48 for prediction of soil types. The experiments were performed on soil data consisting of 110 samples using data analytics tool R. The experimental results predicted that JRIP algorithm performed better as it gave highest accuracy of 98.18% with kappa statistic of approximate 1.0. The problem with this analysis is that the dataset is taken is very small.

Awasthi and Bansal [3] performed comparative study on two data mining techniques namely Artificial Neural Network and Support Vector Machine with the help of data analytics tool R on soil. ANN was implemented with 7 hidden nodes and this model trained for 73073 steps. It predicted accuracy of 55% with root mean square error is of 15. SVM implemented with Radial basis kernel and it achieved much better results with 74% of accuracy. This model is not ideal since the accuracy is very low

Singh et al.[4] implemented different machine learning techniques in order to predict the category of the rice crop yield based on macro-nutrients and micro- nutrients status in dataset. ANN, Naïve Bayes and KNN are applied on soil data. Decision Tree Classifier and Naïve Bayes Classifier are found to be better models for classifying the soils into categories and in the prediction of yield on the basis of Nutrient status in the soil.

Rajesh Kannan Mega lingam [5] presented IOT sensor unit which is part of the rover system consists of moisture sensor and a temperature sensor, for detection of the moisture and temperature respectively in close proximity of plants. The demerit of this project is it little costly and mostly the operation of IOT devices are not feasible at every villages of India.

2.2 Plant Disease Literature Survey

S. S. Sannakki and V. S. Raj purohit, [7], proposed a “Classification of Pomegranate Diseases Based on Back Propagation Neural Network” which mainly works on the method of Segment the defected area and color and texture are used as the features. Here they used neural network classifier for the classification. The main advantage is it Converts to L^*a^*b to extract chromaticity layers of the image and Categorisation is found to be 97.30% accurate. The main disadvantage is that it is used only for the limited crops

P. R. Rothe and R. V. Kshirsagar [8] introduced a” Cotton Leaf Disease Identification using Pattern Recognition Techniques” which Uses snake segmentation, here Hu’s moments are used as distinctive attribute. Active contour model used to limit the vitality inside the infection spot, BPNN classifier tackles the numerous class problems. The average classification is found to be 85.52%. the problem with this model is the background noise.

Godliver Owomugisha, John A. Quinn, Ernest Mwebaze and James Lwasa, proposed” [10] Automated Vision-Based Diagnosis of Banana Bacterial Wilt Disease and Black Sigatoka Disease “Color histograms are extracted and transformed from RGB to HSV, RGB to L^*a^*b . Peak components are used to create max tree, five shape attributes are used and area under the curve analysis is used for classification. They used nearest neighbors, Decision tree, random forest, extremely randomized tree, Naïve bayes and SV classifier. In seven classifiers extremely, randomized trees yield a very high score, provide real time information provide flexibility to the application.

Aakanksha Rastogi, Ritika Arora and Shanu Sharma [11] Leaf Disease Detection and Grading using Computer Vision Technology &Fuzzy Logic”. K-means clustering used to segment the defected area; GLCM is used for the extraction of texture features, Fuzzy logic is used for disease grading. They used artificial neural network (ANN) as a classifier which mainly helps to check the severity of the diseased leaf.

Uan Tian, Chunjiang Zhao, Shenglian Lu and Xinyu Guo [12] SVM-based Multiple Classifier System for Recognition of Wheat Leaf Diseases,” Color features are represented in RGB to HIS, by using GLCM, seven invariant moment are taken as shape parameter. They used SVM classifier which has MCS, used for detecting disease in wheat plant offline.

Aswin Vellaichamy , Akshay Swaminathan , C Varun and Dr. Kalaivani S [13] performed implemented the HSV technique to find the crop disease using leaf using densenet-121 and Achieved Accuracy of 97%

CHAPTER 3

SYSTEM DESIGN

This module consists system design of the project with preliminary design such as overall Architecture diagram and process flow diagram which tells about the modules integration in the project.

System Architecture

The proposed work of the system architecture is shown in Figure 3.1. The proposed system works on Crop Data Set. The target is to get data of the produce, fertilizer, irrigation and disease of the crop.

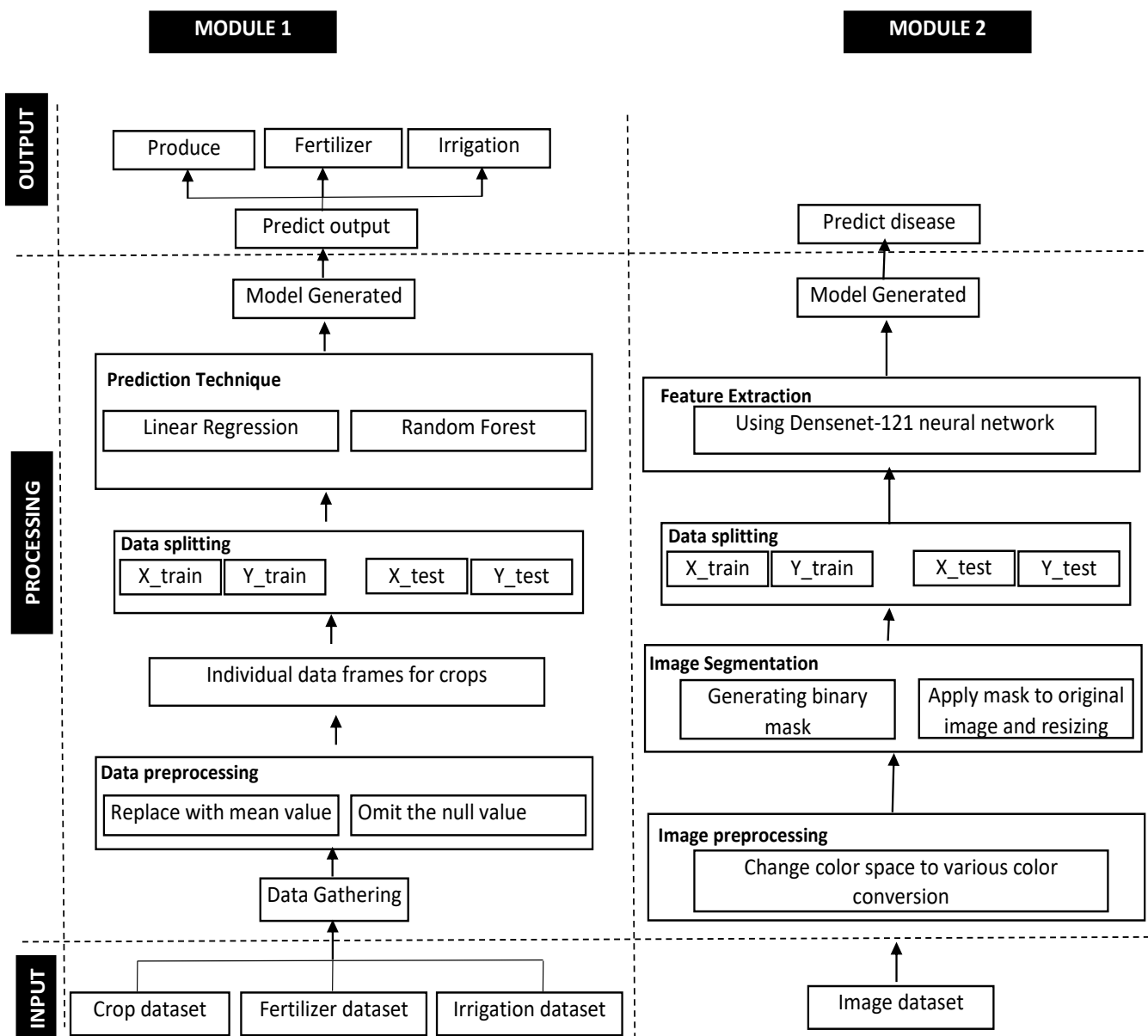


Fig 3.1 System Architecture

3.1 Crop Pattern Prediction

3.1.1 Data Gathering

There are three datasets are used in proposed model these datasets consists of the data regarding the fertilizers, irrigation and crop produce of all district in India. These datasets are collected from year 1966 – 2017. Total 14 type of Commercial crops as well as non-commercial crops. the dataset is collected from ICRISAT website. Further detail of these datasets are given below.

- The produce dataset contains 79 attributes and 16047 rows. These 79 attributes define the basic features of the data which is being forecasted like area, yield and produce. These attributes are divided into area, produce and yield of different crops. In this dataset the target value is the produce of different crop. This produce is dependent on the area.
- The fertilizer dataset contains 20 attributes and 16146 rows. These 20 attributes further divided into 3 main categories of fertilizer like Nitrogen, Phosphate, potash which is used in all district of India in between year 1966 – 2017. The target data that need to be predicted is the composition of Nitrogen, Phosphate, potash on basis of the location (various District of India).
- The irrigation dataset contains 24 attribute and 14506 columns. These attributes define the irrigated area of all the 14 crops that is mentioned in above dataset. The target data is the irrigation amount for different crop this is dependent on the location (various District of India).

At last the data set is divided as training and test data set.

3.1.2 Data Pre-processing

The pre-processing of data is a method for preparing and adapting raw data to a model of learning. This is the first and significant step to construct a machine learning model. Real-world data generally contain noise, missing values and may not be used in an unusable format especially for machine learning models. Data pre-processing needs to be performed in order to purify data and adapt it to the machine learning model of a system which also makes a machine learning model more accurate and efficient. The first thing for data preprocessing is to collect the required data set, and then check the missing values once the data set is imported. Correcting missed values is necessary, or else the data would be difficult to access and maintain. This data is unclean and contain many errors as well as bugs. Correcting missed values is necessary, or else the data would be difficult to access and maintain. Then calculate the mean of the column containing missing values to rectify the missed values, and substitute it with the measured mean. Now, this data set can be used to train a machine learning algorithm to predict values.

3.1.3 Prediction Technique

3.1.3.1 Linear Regression

Linear regression algorithm tries to predict the results by plotting the graph between an independent variable and a dependent variable that are derived from the dataset. It

is a general statistical analysis mechanism used to build machine learning models. The general equation for linear regression is

$$Z = a + b \cdot E$$

Where, Z is the dependent variable and E is independent variable.

3.1.3.2 Random Forest

Random Forest Algorithm is used to incorporate predictions from multiple decision trees into a single model. This algorithm uses bagging mechanism to create a forest of decision trees. It incorporates the predictions from multiple decision trees to give very accurate predictions. The Random Forest algorithm has two steps involved
a) Random forest formation. b) Predict by Random forest classifier generated.

3.1.4 Outcome:

After training the model on above mentioned data then 3 individual model is created whose output are mentioned below:

Output 1: Produce of the crop per hectare

Output 2: Nitrogen, Phosphate, potash amount in kg/hectare

Output 3: irrigation of water on per cubic meter per hectare

3.2 Crop Disease Detection

3.2.1 Image Acquisition

A dataset which consist of various plant disease is taken from Kaggle. This dataset contains 35779 images of size 256 x 256 from Plant Village which contains across 29 labels of diseases for 7 plants namely apple, tomato, corn, bell pepper, grape, cherry and potato. The images used for each category are mentioned in table which gives the image count for each category under various crops.

Crop Name	Image category	Number of Images
Apple	Scab	630
	Black rot	621
	Healthy	1645
Cherry	Cedar apple rust	275
	Powdery mildew	1052
	Healthy	854
Corn	Cercosporin leaf spot	513
	Common rust	1192
	Northern Leaf Blight	985
	Healthy	1162
Grape	Black rot	1180
	Esca (black measles)	1384
	Leaf blight	1076
	Healthy	423

Pepper-bell	Bacterial spot	997
	Healthy	1478
Potato	Early Blight	1000
	Late Blight	1000
	Healthy	152
Tomato	Bacterial spot	2127
	Early blight	1000
	Late blight	1909
	Healthy	1591
	Leaf mold	952
	Septoria leaf spot	1771
	Spider mites	1676
	Yellow leaf curl virus	5357
	Target spot	1404

Table 1 Total no of images for each class of disease

3.2.2 Image Pre-Processing:

The image colour is converted from RGB to other Form. The R, G and B values are divided by 255 to change the range of values from 0-255 to 0-1

$$R' = R/255$$

$$G' = G/255$$

$$B' = B/255$$

$$C_{max} = \max(R', G', B')$$

$$C_{min} = \min(R', G', B')$$

$$\Delta = C_{max} - C_{min}$$

Hue calculation:

$$H = \begin{cases} 60^\circ \times \frac{G' - B'}{\Delta} \bmod 6, C_{max} = R' \\ 60^\circ \times \frac{B' - R'}{\Delta} + 2, C_{max} = G' \\ 60^\circ \times \frac{R' - G'}{\Delta} + 4, C_{max} = B' \end{cases}$$

Saturation calculation:

$$S = \begin{cases} 0 & , \text{if } C_{max} = 0 \\ \frac{\Delta}{C_{max}} & , \text{if } C_{max} \neq 0 \end{cases}$$

Value calculation:

$$V = C_{max}$$

Where, R, G and B represent values of red, green and blue in original image and H, S and V represent hue, saturation and value after processing the original image.

3.2.3 Image Segmentation

Generating binary mask via thresholding image is defined as:

$$g(x, y) = \begin{cases} 1, & T1 \leq f(x, y) < T2 \\ 0, & \text{otherwise} \end{cases}$$

Where x and y represent the coordinal values of the pixels in image g and f . Thresholding is done where T1 is equal to 0.130 and T2 at 0.600 for H-values alone. These values are chosen after statistical analysis of images and their histograms. After this step there are still discrepancies in the image due to noise and shadows, which makes the binary map imperfect. To overcome this, the gaps is filled by morphological reconstruction. After this binary mask

is created. The area of the binary mask is highlighted by a red line along with the identification of the centroid of the image.

The obtained binary mask is then used to make masked image by,

$$\text{Masked image} = \text{binary mask} * \text{Original image}$$

Here “*” represents normal dot multiplication of corresponding values that are present in binary mask and original image. These have been displayed the sample outputs of images obtained in each of the above-described steps below. These masked images are then resized to 64 to reduce time consumption for neural network training.

3.2.4 Feature Extraction with DCNN

Dense Net is a convolutional neural organization where each layer is associated with any non-subsequent layers that are more profound in the organization, that is, the principal layer is associated with the second, third, fourth etc., the subsequent layer is associated with the third, fourth, fifth, etc. This is done to empower greatest data stream between the layers of the organization. To protect the feed-forward nature, each layer acquires contributions from every one of the past layers and gives its own element guides to every one of the layers which will come after it. So, the 'i'th' layer has 'I' data sources and comprises of highlight guides of all its first convolutional blocks. Its own element maps are given to all the following 'I-I' layers. The number 121 in densenet-121 is computed as follows.

DenseNet-121: $5 + [2 \times (6 + 12 + 24 + 16)]$

5– Convolution and Pooling Layer

3– Transition Layers (6, 12, 24)

1– Classification Layer (16)

2– Dense Block (1×1 and 3×3 conv)

3.2.5 Outcome:

As an output the model predict the type of disease related to the crop.

CHAPTER 4

ALGORITHM AND PSEUDO CODE

This section explains in detail the various modules in the system. Each module includes the input for the module, process flow for the module and output for the module in detail.

4.1 Crop Produce Prediction Using Linear Regression

Linear regression is a linear approach to modelling the relationship between a scalar response (or dependent variable) and one or more explanatory variables (or independent variables). The case of one explanatory variable is called simple linear regression. This Algorithm give the accuracy of 78 to 93 percent ranging on different crop. Machine learning, more specifically the field of predictive modelling is primarily concerned with minimizing the error of a model or making the most accurate predictions possible, at the expense of explain ability. In applied machine learning we will borrow, reuse and steal algorithms from many different fields, including statistics and use them towards these ends. As such, linear regression was developed in the field of statistics and is studied as a model for understanding the relationship between input and output numerical variables, but has been borrowed by machine learning. It is both a statistical algorithm and a machine learning algorithm. Linear Regression for Machine Learning more specifically the field of predictive modelling is primarily concerned with minimizing the error of a model or making the most accurate predictions possible, at the expense of explain ability. In applied machine learning we will borrow, reuse and steal It is both a statistical algorithm and a machine learning algorithm. He reason is because linear regression has been around for so long (more than 200 years). It has been studied from every possible angle and often each angle has a new and different name. Linear regression is a linear model, e.g. a model that assumes a linear relationship between the input variables (x) and the single output variable (y). More specifically, that y can be calculated from a linear combination of the input variables (x).

Linear Regression Algorithm to find the crop produce

```

1: Start
2: import linear regression from sklearn
3: take variable lr = Linear Regression(normalize=True)
4: get the X train = train df(area)
5: Get the Y train = train df(produce)
6: Get the x test = test df. drop (produce, axis=1). copy()
7: fit in variable lr.fit (x train, Y train)
8: lr. predict = lr. predict (X test)
9: lr. accuracy = round (lr. score(X train,Y train) * 100,2)
10: print lr. accuracy

```

OUTPUT: Got Accuracy of 78 to 93 %. Depending on type of crop.

4.2 Crop fertilizer and irrigation Prediction using Random Forest

Random Forest Classifier- Random forests or random decision forests are an ensemble learning method for classification, regression and other tasks that operate by constructing a multitude of decision trees at training time and outputting the class that is the mode of the classes (classification) or mean prediction (regression) of the individual trees. Random decision forests correct for decision trees' habit of over fitting to their training set.

Random Forest Algorithm to find the irrigation and fertilizer value

```

1: start
2: import RandomForestRegressor from sklearn.
3: take variable tree1,tree2 = RandomForestRegressor
4: split the irrigation dataframe according to the crop.
5: split the fertilizer dataframe according to the district.
6: get the x1train = train df (district name)
7: get the x2train = train df (district)
8: Get the y1 train = train df(fertilizer names)
9: Get the y2train = train df(crop irrigation data )
10: create the test data for both the data frame also using train_test_split feature
11: fit in variable tree1.fit(x1train, y1train)
12: fit in variable tree2.fit(x2train, y2train)
13: output1 = tree1.predict(x1 test)
14: output2 = tree2.predict(x2.test)
15: output1.accuracy = print round(tree1.score(X train,Y train) * 100,2)
16: output2.accuracy = print round(tree2.score(X train,Y train) * 100,2)

```

OUTPUT: Got Accuracy of 84 % to 93 %.in predicting irrigation amount, depending on type of crop.

4.3 Crop disease prediction on basis of Densenet -121

The problems arise with CNNs when they go deeper. This is because the path for information from the input layer until the output layer (and for the gradient in the opposite direction) becomes so big, that they can get vanished before reaching the other side. The dense net requires parameters than an equivalent traditional CNN, as there is no need to learn redundant feature maps, it combines the previous layer output with a future layer. In Dense net the layer is connected densely to all layer. Details of the DenseNet-121 is following: 5—convolution and pooling layers, 3—transition layers (6,12,24), 1—Classification layer (16) and 2—dense block (1×1 and 3×3 conv).

Generally, traditional CNNs calculate the output layers (l th) using a non-linear transformation $H_l(.)$ to the output of the previous layer X_{l-1}

$$X_l = H_l(X_{l-1}).$$

DenseNets do not really sum up the layer output functionality maps with the inputs but concatenate them. DenseNet offers an easy communication model for improving

information flow between layers: the l th layer receives inputs from the features of all previous levels: The equation is then again transformed into

$$X_l = H_l [(X_0, X_1, X_2, \dots, X_{l-1})],$$

where $[X_0, X_1, X_2, \dots, X_{l-1}]$ is a single tensor formed by the concatenation of the output maps of previous layers. Out of the functions, $H_l(.)$ represents a non-linear transformation function. This function consists of three major operations, batch normalization (BN), activation (ReLU) and pooling and convolution (CONV)

Dense net Algorithm to find the disease of plant.

```

1: start
2: image Acquisition.
3: Convert the RGB to HSV, lab colour and greyscale format.
4: Generate Binary Mask after filling gaps.
5: Apply mask to the original image and Resizing it.
6: import Densenet 121 from keras.application .
7: get xtrain = np.zeros((train.shape[0], IMAGE_SIZE, IMAGE_SIZE, 3))
8: Normalise the data by dividing it 255.
9: get ytrain=train['DiseaseID'].values.
10: ytrain = to_categorical(y_train, num_classes=15).
11: xtest=pick an image and normalize it and feed to model.
12: use ImageDataGenerator and fit the imane into it.
13: get Accuracy by model.fit_generator function

```

OUTPUT: Got Accuracy of 96.56%.

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