



**VIT**<sup>®</sup>  
**Vellore Institute of Technology**  
(Deemed to be University under section 3 of UGC Act, 1956)

# **CSE3013- ARTIFICIAL INTELLIGENCE** **PROJECT FINAL REVIEW**

**SUBMITTED BY:**

**Arya Dubey (20BCE0908)**

**Prachurya Priyadarshini (20BCT0155)**

**Prakhar Kandpal (20BCE2117)**

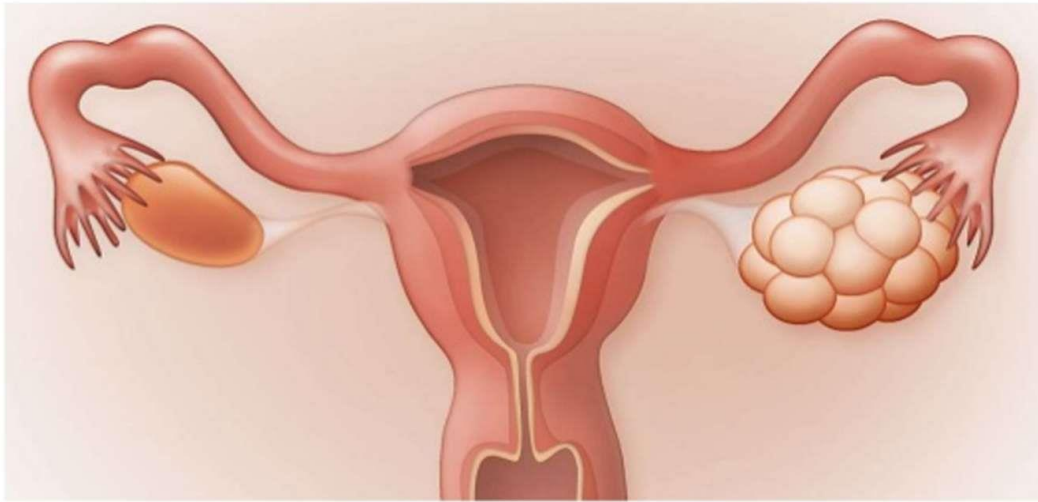
**Abhinav Bijith (20BCE2149)**

**UNDER THE GUIDANCE OF:**

**PROF. GUNAVATHI C**

# **Polycystic Ovarian Syndrome Detecting** **Analyser**

Polycystic Ovarian Syndrome (PCOS)



## **CERTIFICATE**

**This is to certify that the project entitled “Polycystic Ovarian Syndrome Detecting Analyzer”, submitted by Arya Dubey (20BCE0908), Prachurya Priyadarshini (20BCT0155), Prakhar Kandpal (20BCE2117), Abhinav Bijith (20BCE2149), VIT, for the award of the degree of Bachelor of Technology in Programme, is a record of bonafide work carried out by them under my supervision during the period, 02. 08. 2021 to 10.12.2021, as per the VIT code of academic and research ethics.**

**The contents of this report have not been submitted and will not be submitted either in part or in full, for the award of any other degree or diploma in this institute or any other institute or university. The thesis fulfills the requirements and regulations of the University and in my opinion meets the necessary standards for submission.**

**Place : Vellore**

**Date :**

**Signature of the Guide**

**Internal Examiner**

**External Examiner**

**Head of the Department Programme**

## **DECLARATION**

**We thereby declare that the project entitled “Polycystic Ovarian Syndrome Detecting Analyser” submitted by us, for the award of the degree of Bachelor of Technology in Programme to VIT is a record of bonafide work carried out by us under the supervision of Prof. Gunavathi C. We further declare that the work reported in this project has not been submitted and will not be submitted, either in part or in full, for the award of any other degree or diploma in this institute or any other institute or university.**

## **EXECUTIVE SUMMARY**

In the past few decades, technology has revolutionized our universe and affected our lives, making them easier day by day. Emerging technologies are reshaping mankind in a lot of ways. These days, machine learning, a field of study that gives computers to learn without being explicitly programmed, is playing a key role in the healthcare sector. Machine learning can deal with obscenely huge datasets, convert analysed data into clinical insights and help in the diagnosis of various ailments. Polycystic Ovary Syndrome (PCOS) is a medical condition which causes hormonal disorder in women in their childbearing years. PCOS occurs as a result of hormonal imbalances. In this disorder, the ovaries develop small collections of fluids called follicles (cysts) and fail to release eggs, which is why women suffering from PCOS tend to have complications in conceiving . A lot of women have PCOS, but do not get diagnosed with it at an earlier stage. In a study, 69 to 70 percent of women did not have a pre-existing diagnosis.

While the actual causes of PCOS remain a mystery, studies say that it is generally inherited. It is a very unpredictable condition as the cure is uncertain since there is no observable trend for this medical condition.

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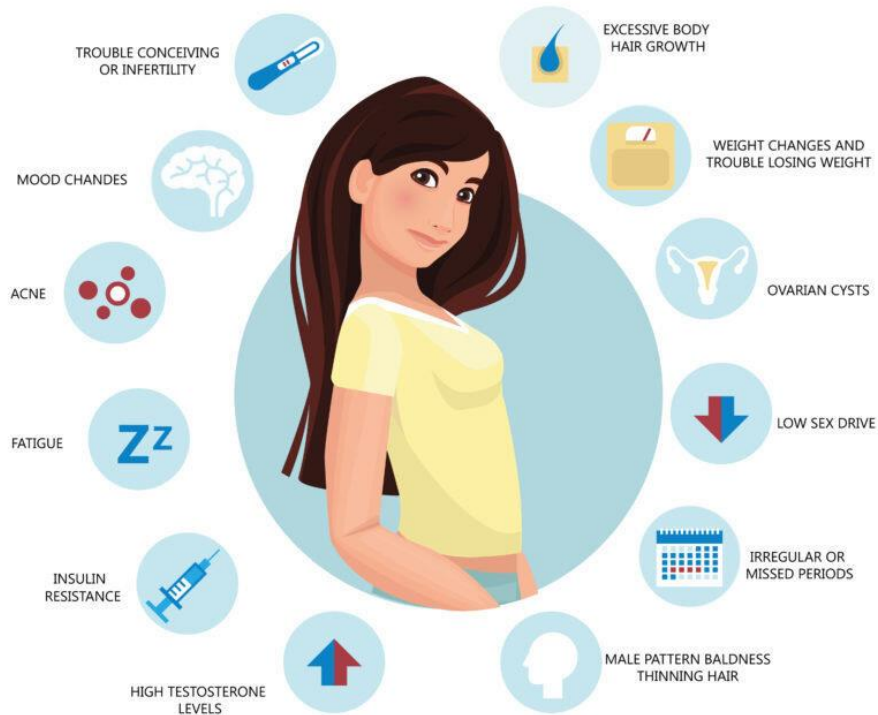
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## **INTRODUCTION:**

Polycystic ovary syndrome (PCOS) is a condition in which the ovaries produce an abnormal amount of androgens, male hormones that are usually present in women in small amounts. The name polycystic ovary syndrome describes the numerous small cysts (fluid-filled sacs) that form in the ovaries. The basic symptoms of PCOS are:

- Cysts in ovaries.
- High levels of hormone: androgen.
- Irregular Periods
- Excessive body hair growth
- Weight gain especially around the belly

# PCOS SYMPTOMS



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## **PROBLEM STATEMENT**

Several factors cause infertility; one of which has been the irregular number and the dimension of follicle development in the ovulation phase. Such abnormality has been the first symptom of PCOS.

Women having PCOS are at risk of improving a diversity of symptoms/intricacies involving variant systems, i.e., gynecological, metabolic, physical, cardiac, and psychological. Perfect finding of the PCOS is important for the treatment.

Real-time analysis of PCOS is a major task as follicles contain different sizes and are highly connected with tissues and blood vessels which results in errors. In contemporary years, transabdominal and/or transvaginal ultrasound have turned into the most generally utilized diagnostic techniques for the recognition of PCOS.

Manual identification of follicles may cause several problems, such as more time is needed for the follicles' identification, inter and intra-observer inconsistency and sometimes they could seriously



affect women's health. Additionally, there is no assessment on the use of a better diagnosis for PCOS awareness and management. Thus, a more general machine learning procedure is needed for the better diagnosis of PCOS.



## **OBJECTIVE**

- This project is an attempt to develop a real-time object detection using machine learning concepts. The project aims to deliver an implemented approach to detect PCOS. In the modern era, there are several new technologies available to diagnose various diseases and one of them is Machine learning algorithms because they are exposed to new data. These algorithms learn from past experiences to produce reliable and repeatable decisions. In this article, Machine learning algorithms are used to identify the important features to diagnose PCOS.
- From 1 in 10 women suffering from PCOS worldwide to currently 3-4 in 10 women, PCOS is now exponentially increasing among women due to an unhealthy lifestyle. The literature says that 1 in every 5 women in India suffers from PCOS. PCOS symptoms differ in every patient.
- The major diagnosis includes scanning for follicles, their number and sizes using Ultrasound imaging.

- Even though it is called Polycystic Ovary Syndrome, it is not essentially described by ovarian cysts. It is defined by examining at least two of three diagnostic criteria.
- In 2012, the 2003 Rotterdam criteria were endorsed by NIH for PCOS.
- For an accurate PCOS diagnosis, disorders that have specific signs and symptoms that match with those of PCOS must be dismissed.

# POLYCYSTIC OVARY SYNDROME (PCOS): SYMPTOMS, DIAGNOSIS AND TREATMENT

Polycystic ovary syndrome (PCOS) is a condition that affects a woman's hormone levels.

## SYMPTOMS



Missed periods,  
irregular periods



Large ovaries



Excess body hair



Weight gain



Acne or oily skin



Male-pattern baldness



Infertility

## CAUSES

The exact cause of PCOS isn't known.

### DIAGNOSIS



Physical exam



Ultrasound



Blood tests

### TREATMENT



Medications



Birth control pills



Surgery

## HOME REMEDIES



Eating a healthy diet



Regular physical  
activity



Maintaining a  
healthy weight



Avoid smoking

## **PROJECT DESCRIPTION**

- The team has used different machine learning algorithms like K-nearest neighbour (KNN), decision tree and SVM with different kernel functions to predict PCOS from the identification of new genes.
- It has used machine learning algorithms like Logistic Regression (LR) and Random Classifier to develop an automated system that will act as an assisted tool for the doctor for saving considerable time in examining the patients and hence reducing the delay in diagnosing the risk of PCOS by using metabolic and clinical factors in a feature vector.

In the modern era, there are several new technologies available to diagnose PCOS and one of them is Machine learning algorithms because they are exposed to new data. These algorithms learn from past experiences to produce reliable and repeatable decisions. In this article, Machine learning algorithms are used to identify the important features to diagnose PCOS.

## **TECHNICAL SPECIFICATIONS**

Machine Learning Algorithms like K-nearest neighbour (KNN), decision tree and SVM with different kernel functions are used in this project to predict PCOS from the identification of new genes. Other ML Algorithms like Logistic Regression (LR) and Random Classifier are implemented to develop an automated system that will act as an assisted tool for the doctor for saving considerable time in examining the patients and hence reducing the delay in diagnosing the risk of PCOS by using metabolic and clinical factors in a feature vector.

Here's how we will do it:

- The first step to building our model is to import our libraries and datasets into our Google Colab notebook.
- pandas: The most popular python library that is used for data manipulation and analysis. In this project, it is primarily useful for dataframe manipulation.
- NumPy: A python library that provides support for large, multi-dimensional arrays and matrices, and has high-level mathematical functions to help operate on and manipulate these arrays.
- matplotlib.pyplot and seaborn: Used for data visualization.

- We can start the project by making sure we have installed the latest version of seaborn, which will be used for data visualization.

Then, we can proceed on to the architecture model of our project.

Below is the table for the Literature Survey which is used for distilling information on thesis and research papers mentioned below to scrutinize the problem.

## **LITERATURE SURVEY**

A literature survey is a piece of discursive prose, not a list describing or summarizing one piece of literature after another. It is an iterative process, assessing and distilling information. One of the key purposes of the literature survey is to investigate a problem that no one else has addressed.

Authors & Year Reference	Title Study	Journal/ Conference	Concept & Analysis	Limitations
C. Gopalkrishnan & M. Iyapparaja	Detection of PCOS from Ultrasound Images of Ovaries	International Journal of Recent Technology and Engineering September 2019	Description & Mechanism of PCOS, Ultrasound & Ultrasonography	It is difficult to implement the process of detecting PCOS from ultrasound image ovaries in accurate manner. New hybrid techniques have to develop for its improvement.
B. Bhanu	Follicle Diagnosis in PCOS	IJRTC 1995	Genetic Algorithm (GA)	Computational requirements are varied by the number of parameters
T. Chiang and Y. Q. Zhang	Delay in Period Cycle Measure	IJRTC 1997	Quadratic-Rate distortion method	Detection of minimum is failed
G. Vasavi & Dr. S. Jyothi	PCOS Detection Using Various Machine Learning Methods	Adv Research Journal in Dynamical & Control Systems, 05- Special Issue, July 2017	K-Nearest Neighbour Technique Based Classification of Biomedical Objects	Only few techniques have presented effective performance. Although the implementation of such techniques has proffered good quality outcomes but a great number of tasks are still to

be enhanced in future.

Anita Raj & Remya George

Prediction System of PCOS using ML

Tencon 2019- IEEE Region 10 Conference

Classification of PCOS with the feature set transformed with Principal component Analysis (PCA)

Still difficult to diagnose or detect but give 90% of the best results.

Usman

Texture Features & Artificial Neural Network for Follicle Detection

JARDCS 2011

Automatic characterization of ovaries amidst complete female cycle

Necessity for enhancing on the denoising methodology as medical images is fraught with impair noise

# DESIGN APPROACH AND DETAILS

```
In [ ]: !pip install seaborn --upgrade
```

```
Requirement already up-to-date: seaborn in /usr/local/lib/python3.7/dist-packages (0.11.1)
Requirement already satisfied, skipping upgrade: pandas>=0.23 in /usr/local/lib/python3.7/dist-packages (from seaborn) (1.1.5)
Requirement already satisfied, skipping upgrade: numpy>=1.15 in /usr/local/lib/python3.7/dist-packages (from seaborn) (1.19.5)
Requirement already satisfied, skipping upgrade: matplotlib>=2.2 in /usr/local/lib/python3.7/dist-packages (from seaborn) (3.2.2)
Requirement already satisfied, skipping upgrade: scipy>=1.0 in /usr/local/lib/python3.7/dist-packages (from seaborn) (1.4.1)
Requirement already satisfied, skipping upgrade: pytz>=2017.2 in /usr/local/lib/python3.7/dist-packages (from pandas>=0.23->seaborn) (2018.9)
Requirement already satisfied, skipping upgrade: python-dateutil>=2.7.3 in /usr/local/lib/python3.7/dist-packages (from pandas>=0.23->seaborn) (2.8.1)
Requirement already satisfied, skipping upgrade: pyparsing!=2.0.4,!=2.1.2,!=2.1.6,>=2.0.1 in /usr/local/lib/python3.7/dist-packages (from matplotlib>=2.2->seaborn) (2.4.7)
Requirement already satisfied, skipping upgrade: cycler>=0.10 in /usr/local/lib/python3.7/dist-packages (from matplotlib>=2.2->seaborn) (0.10.0)
Requirement already satisfied, skipping upgrade: kiwisolver>=1.0.1 in /usr/local/lib/python3.7/dist-packages (from matplotlib>=2.2->seaborn) (1.3.1)
Requirement already satisfied, skipping upgrade: six>=1.5 in /usr/local/lib/python3.7/dist-packages (from python-dateutil>=2.7.3->pandas>=0.23->seaborn) (1.15.0)
```

```
In [ ]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
%matplotlib inline
import seaborn as sns
sns.__version__
```

```
Out[ ]: '0.11.1'
```

```
In [ ]: from google.colab import files
uploaded = files.upload()
pcos = pd.read_csv('PCOS_no_infertility.csv')
pcos.head(15)
```

Upload widget is only available when the cell has been executed in the current browser session. Please rerun this cell to enable.  
Saving PCOS\_no\_infertility.csv to PCOS\_no\_infertility (1).csv

```
In [ ]: from google.colab import files
uploaded = files.upload()
pcos = pd.read_csv('PCOS_no_infertility.csv')
pcos.head(15)
```

Upload widget is only available when the cell has been executed in the current browser session. Please rerun this cell to enable.  
Saving PCOS\_no\_infertility.csv to PCOS\_no\_infertility (1).csv

```
Out[ ]: Unnamed: 0 Unnamed: 1 Unnamed: 2 Unnamed: 3 Unnamed: 4 Unnamed: 5 Unnamed: 6 Unnamed: 7 Unnamed: 8 Unnamed: 9 Unnamed: 10 Unnamed: 11 Unnamed: 12 Unnamed: 13 Unnamed: 14
```

	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14
	PCOS (Y/N)	Age (yrs)	Weight (Kg)	Height	BMI	Blood Group	Pulse rate(bpm)	RR (breaths/min)	Hb(g/dl)	Cycle(R/I)	Cycle length(days)	Marraige Status (Yrs)	Pregnant(Y/N)	No. of absorptions	HCG(n)
0	0	28	44.6	152	19.3	15	78	22	10.48	2	5	7	0	0	
2	0	36	65	161.5	24.92116266	15	74	20	11.7	2	5	11	1	0	
3	1	33	68.8	165	25.27089073	11	72	18	11.8	2	5	10	1	0	
4	0	37	65	148	29.67494522	13	72	20	12	2	5	4	0	0	
5	0	25	52	161	20.06095444	11	72	18	10	2	5	1	1	0	
6	0	36	74.1	165	27.21763085	15	78	28	11.2	2	5	8	1	0	
7	0	34	64	156	26.29848784	11	72	18	10.9	2	5	2	0	0	
8	0	33	58.5	159	23.13990744	13	72	20	11	2	5	13	1	2	
9	0	32	40	158	16.02307323	11	72	18	11.8	2	5	8	0	1	
10	0	36	52	150	23.11111111	15	80	20	10	4	2	4	0	0	
11	0	20	71	163	26.72287252	15	80	20	10	2	5	4	1	2	
12	0	26	49	160	19.140625	13	72	20	9.5	2	5	3	0	1	
13	1	25	74	152	32.02908587	17	72	18	11.7	4	2	7	1	0	
14	0	38	50	152	21.64127424	13	74	20	12.1	2	5	15	0	0	



```
In [ ]: df = pcos
```

```
In [ ]: #Use first row of data as df header
new_header = df.iloc[0]
df = df[1:]
df.columns = new_header
```

```
In [ ]: df
```

Out[ ]:

	PCOS (Y/N)	Age (yrs)	Weight (Kg)	Height	BMI	Blood Group	Pulse rate(bpm)	RR (breaths/min)	Hb(g/dl)	Cycle(R/I)	Cycle length(days)	Marriage Status (Yrs)	Pregnant(Y/N)	No. of abortions	I beta- HCG(mIU/mL)	II HCG(mIU/mL)
1	0	28	44.6	152	19.3	15	78	22	10.48	2	5	7	0	0	1.99	
2	0	36	65	161.5	24.92116286	15	74	20	11.7	2	5	11	1	0	60.8	
3	1	33	68.8	165	25.27089073	11	72	18	11.8	2	5	10	1	0	494.08	
4	0	37	65	148	29.67494522	13	72	20	12	2	5	4	0	0	1.99	
5	0	25	52	161	20.06095444	11	72	18	10	2	5	1	1	0	801.45	
...	...	...	...	...	...	...	...	...	...	...	...	...	...	...	...	...
537	0	35	50	164.592	18.5	17	72	16	11	2	5	8	0	1	1.99	
538	0	30	63.2	158	25.3	15	72	18	10.8	2	5	4	1	1	80.13	
539	0	36	54	152	23.4	13	74	20	10.8	2	6	8	0	0	1.99	
540	0	27	50	150	22.2	15	74	20	12	4	2	2	0	0	292.92	
541	1	23	82	165	30.1	13	80	20	10.2	4	7	2	0	0	1.99	

541 rows × 42 columns

```
In [ ]: #Checking for null values
df.isnull().sum()
```

```
In [ ]: df
```

Out[ ]:

	PCOS (Y/N)	Age (yrs)	Weight (Kg)	Height	BMI	Blood Group	Pulse rate(bpm)	RR (breaths/min)	Hb(g/dl)	Cycle(R/I)	Cycle length(days)	Marriage Status (Yrs)	Pregnant(Y/N)	No. of abortions	I beta- HCG(mIU/mL)	II HCG(mIU/mL)
1	0	28	44.6	152	19.3	15	78	22	10.48	2	5	7	0	0	1.99	
2	0	36	65	161.5	24.92116286	15	74	20	11.7	2	5	11	1	0	60.8	
3	1	33	68.8	165	25.27089073	11	72	18	11.8	2	5	10	1	0	494.08	
4	0	37	65	148	29.67494522	13	72	20	12	2	5	4	0	0	1.99	
5	0	25	52	161	20.06095444	11	72	18	10	2	5	1	1	0	801.45	
...	...	...	...	...	...	...	...	...	...	...	...	...	...	...	...	...
537	0	35	50	164.592	18.5	17	72	16	11	2	5	8	0	1	1.99	
538	0	30	63.2	158	25.3	15	72	18	10.8	2	5	4	1	1	80.13	
539	0	36	54	152	23.4	13	74	20	10.8	2	6	8	0	0	1.99	
540	0	27	50	150	22.2	15	74	20	12	4	2	2	0	0	292.92	
541	1	23	82	165	30.1	13	80	20	10.2	4	7	2	0	0	1.99	

539 rows × 42 columns

```
In [ ]: df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
Int64Index: 539 entries, 1 to 541
Data columns (total 42 columns):
#   Column                Non-Null Count  Dtype
---  -
0   PCOS (Y/N)            539 non-null    object
1   Age (yrs)             539 non-null    object
2   Weight (Kg)           539 non-null    object
3   Height                539 non-null    object
4   BMI                   539 non-null    object
5   Blood Group           539 non-null    object
6   Pulse rate(bpm)       539 non-null    object
7   RR (breaths/min)      539 non-null    object
8   Hb(g/dl)              539 non-null    object
```

```
In [ ]: # Create a loop to convert all values into numeric
        for column in df:

            # Select column contents by column
            # name using [] operator
            columnSeriesObj = df[column]
            df[column] = pd.to_numeric(df[column], errors='coerce')
```

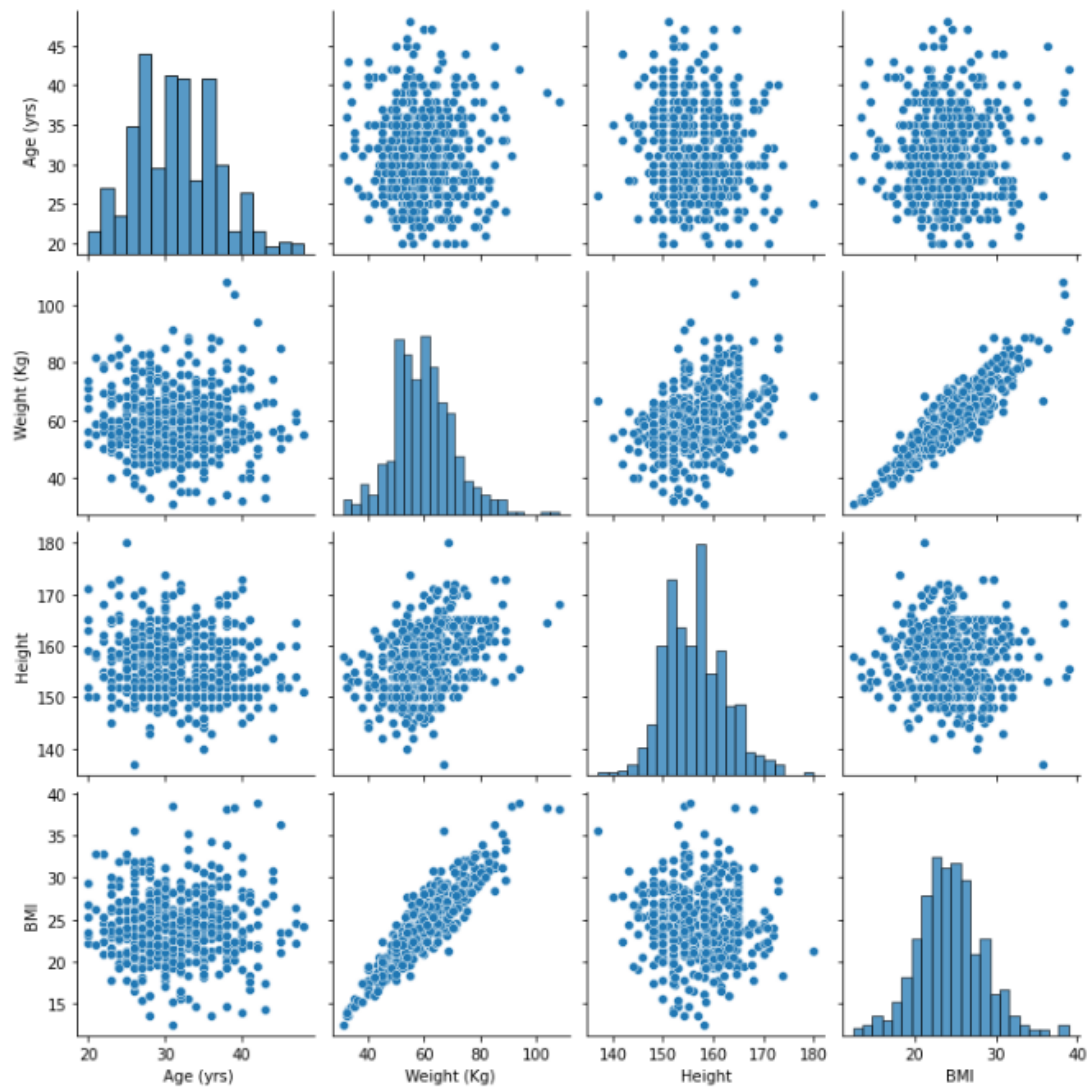
/usr/local/lib/python3.7/dist-packages/ipykernel\_launcher.py:7: SettingWithCopyWarning:  
A value is trying to be set on a copy of a slice from a DataFrame.  
Try using .loc[row\_indexer,col\_indexer] = value instead

See the caveats in the documentation: [https://pandas.pydata.org/pandas-docs/stable/user\\_guide/indexing.html#returning-a-view-versus-a-copy](https://pandas.pydata.org/pandas-docs/stable/user_guide/indexing.html#returning-a-view-versus-a-copy)  
import sys

```
In [ ]: df.info()
```

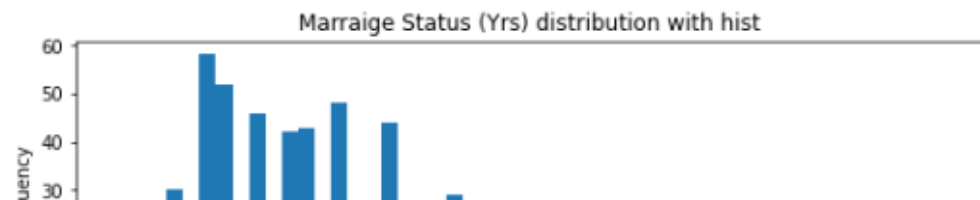
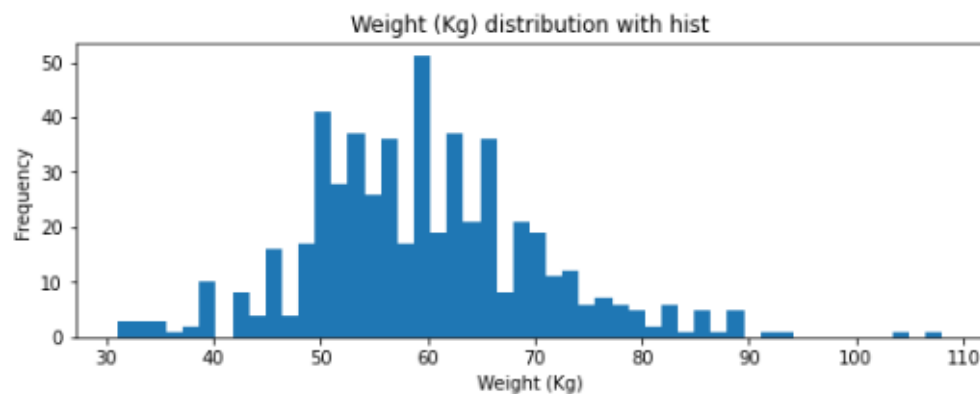
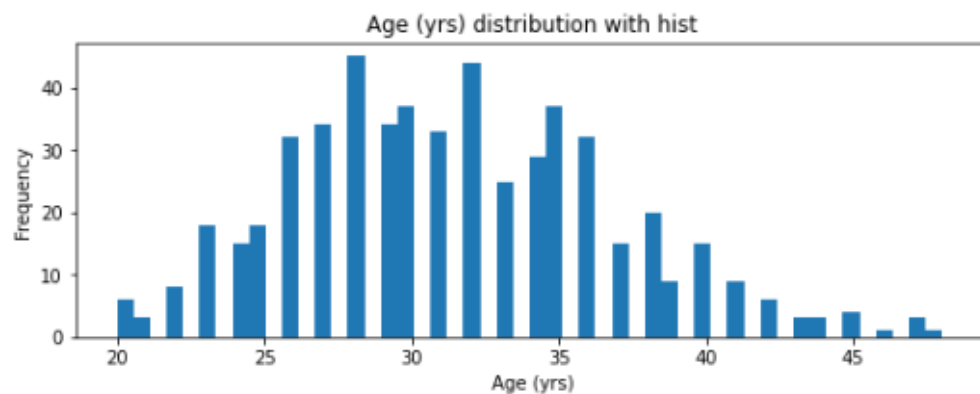
```
<class 'pandas.core.frame.DataFrame'>
Int64Index: 539 entries, 1 to 541
Data columns (total 42 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   PCOS (Y/N)                            539 non-null    int64
1   Age (yrs)                             539 non-null    int64
2   Weight (Kg)                           539 non-null    float64
3   Height                                539 non-null    float64
4   BMI                                    539 non-null    float64
5   Blood Group                           539 non-null    int64
6   Pulse rate(bpm)                       539 non-null    int64
7   RR (breaths/min)                      539 non-null    int64
8   Hb(g/dl)                              539 non-null    float64
9   Cycle(R/I)                            539 non-null    int64
10  Cycle length(days)                    539 non-null    int64
11  Marraige Status (Yrs)                  539 non-null    float64
12  Pregnant(Y/N)                          539 non-null    int64
13  No. of abortions                       539 non-null    int64
14  I   beta-HCG(mIU/mL)                   539 non-null    float64
15  II  beta-HCG(mIU/mL)                    538 non-null    float64
16  FSH(mIU/mL)                           539 non-null    float64
17  LH(mIU/mL)                            539 non-null    float64
18  FSH/LH                                539 non-null    float64
19  Hip(inch)                             539 non-null    int64
20  Waist(inch)                           539 non-null    int64
21  Waist:Hip Ratio                        539 non-null    float64
22  TSH (mIU/L)                           539 non-null    float64
23  AMH(ng/mL)                            538 non-null    float64
24  PRL(ng/mL)                            539 non-null    float64
25  Vit D3 (ng/mL)                        539 non-null    float64
```

```
Out[ ]: <seaborn.axisgrid.PairGrid at 0x7ff7b0e0b910>
```



```
In [ ]: #Plot histogram
def plot_hist(variable):
    plt.figure(figsize = (9,3))
    plt.hist(df[variable], bins = 50)
    plt.xlabel(variable)
    plt.ylabel("Frequency")
    plt.title("{} distribution with hist".format(variable))
    plt.show()
```

```
In [ ]: numericVar = [" Age (yrs)", "Weight (Kg)","Marraige Status (Yrs)"]
for n in numericVar:
    plot_hist(n)
```



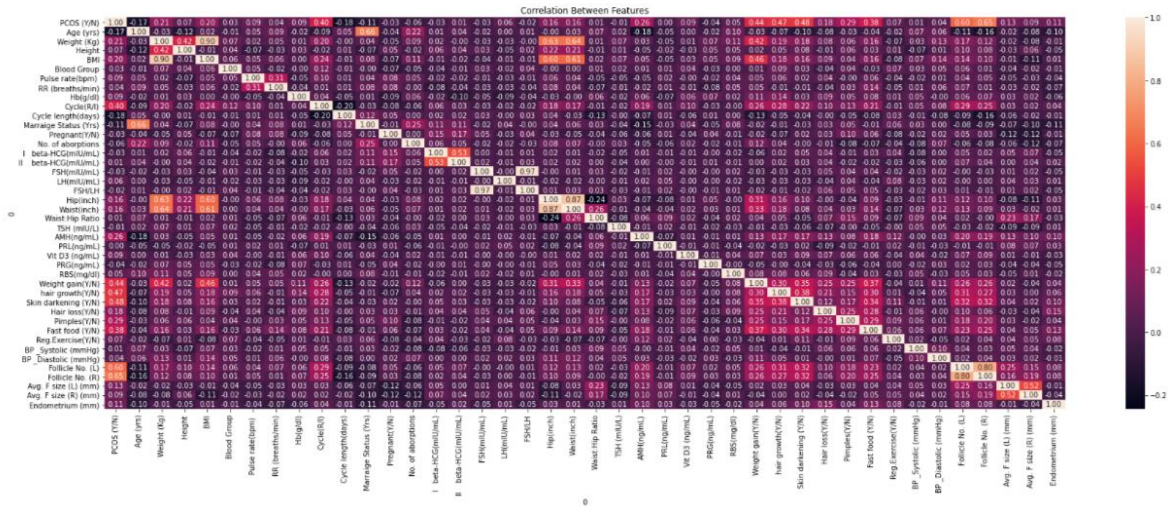
In [ ]: df.corr()

Out[ ]:

	PCOS (Y/N)	Age (yrs)	Weight (Kg)	Height	BMI	Blood Group	Pulse rate(bpm)	RR (breaths/min)	Hb(g/dl)	Cycle(R/I)	Cycle length(days)	Marriage Status (Yrs)	Pregnant(Y/N)	No. of abortion
0														
PCOS (Y/N)	1.000000	-0.172436	0.210280	0.066361	0.198638	0.034244	0.092437	0.038031	0.086934	0.402312	-0.183781	-0.113701	-0.026586	-0.056741
Age (yrs)	-0.172436	1.000000	-0.031582	-0.121124	0.019868	-0.012655	0.045910	0.089509	-0.022466	-0.086540	0.052985	0.661834	-0.044235	0.221169
Weight (Kg)	0.210280	-0.031582	1.000000	0.420005	0.901719	0.071601	0.019995	0.045048	0.008979	0.201862	-0.004477	0.043238	-0.051148	0.093612
Height	0.066361	-0.121124	0.420005	1.000000	-0.006869	0.039981	-0.073801	-0.031093	0.025084	-0.020910	0.007512	-0.066508	0.048157	-0.024771
BMI	0.198638	0.019868	0.901719	-0.006869	1.000000	0.061174	0.050277	0.064197	0.002533	0.235702	-0.007615	0.083298	-0.074875	0.109512
Blood Group	0.034244	-0.012655	0.071601	0.039981	0.061174	1.000000	0.047719	-0.023064	-0.002153	0.123744	-0.008485	-0.002247	-0.070714	-0.053812
Pulse rate(bpm)	0.092437	0.045910	0.019995	-0.073801	0.050277	0.047719	1.000000	0.305751	-0.052581	0.103183	0.006831	0.038505	0.081828	0.045612
RR (breaths/min)	0.038031	0.089509	0.045048	-0.031093	0.064197	-0.023064	0.305751	1.000000	-0.038149	0.012243	0.005788	0.079823	0.081720	-0.004812
Hb(g/dl)	0.086934	-0.022466	0.008979	0.025084	0.002533	-0.002153	-0.052581	-0.038149	1.000000	0.039702	-0.052484	0.006273	-0.094530	0.059510
Cycle(R/I)	0.402312	-0.086540	0.201862	-0.020910	0.235702	0.123744	0.103183	0.012243	0.039702	1.000000	-0.204648	-0.032283	-0.077656	-0.055237
Cycle length(days)	-0.183781	0.052985	-0.004477	0.007512	-0.007615	-0.008485	0.006831	0.005788	-0.052484	-0.204648	1.000000	0.117289	0.050248	0.004512
Marriage Status (Yrs)	-0.113701	0.661834	0.043238	-0.066508	0.083298	-0.002247	0.038505	0.079823	0.006273	-0.032283	0.117289	1.000000	-0.007629	0.246122
Pregnant(Y/N)	-0.026586	-0.044235	-0.051148	0.048157	-0.074875	-0.070714	0.081828	0.081720	-0.094530	-0.077656	0.050248	-0.007629	1.000000	0.002085
No. of abortions	-0.056741	0.221169	0.093612	-0.024771	0.109512	-0.053812	0.045612	-0.004812	0.059510	-0.055237	0.004583	0.246122	0.002085	1.000000
I beta-HCG(mIU/mL)	-0.027820	0.007928	0.015838	0.062524	-0.010223	-0.035430	-0.020625	-0.084756	-0.017163	0.064315	0.020100	0.111390	0.150637	0.057712
II beta-HCG(mIU/mL)	0.013413	0.042787	-0.000924	0.036754	-0.015905	-0.011020	-0.016383	-0.038889	-0.095008	0.028944	0.018823	0.112525	0.170652	0.046712
FSH(mIU/mL)	-0.030300	-0.017804	-0.025759	0.031094	-0.040789	0.028154	-0.013149	-0.032302	-0.047566	-0.025800	0.029796	-0.023549	0.052763	-0.018212
LH(mIU/mL)	0.064136	0.000479	-0.029878	-0.045381	-0.013384	-0.019520	-0.032415	-0.031081	-0.089329	-0.020987	-0.001587	0.035410	-0.034381	-0.018912
FSH/LH	-0.018535	0.012367	-0.004925	0.021976	-0.012138	0.036223	-0.013089	-0.043382	-0.039832	-0.016318	0.025847	-0.003001	0.044657	-0.026412
Hip(inch)	0.161700	-0.003565	0.633911	0.216065	0.596729	-0.001977	0.062698	0.075094	-0.025845	0.175864	0.039425	0.037672	-0.032181	0.078912
Waist(inch)	0.163143	0.033705	0.638154	0.208676	0.607384	0.003385	0.038037	0.038064	0.001473	0.168933	0.026418	0.056751	0.048564	0.073812

## PROJECT DEMONSTRATION

```
In [ ]: #Plot correlation matrix
corr_matrix = df.corr()
plt.subplots(figsize=(30,10))
sns.heatmap(corr_matrix, annot = True, fmt = ".2f");
plt.title("Correlation Between Features")
plt.show()
```



```
In [ ]: #Split the dataset into 70% training data and 30% testing data
from sklearn.model_selection import train_test_split
X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size = 0.3 , random_state = 0)
```

```
In [ ]: #Scale the data (Feature Scaling)
from sklearn.preprocessing import StandardScaler
sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.fit_transform(X_test)
```

```
In [ ]: # Create a function for the models
def models(X_train, Y_train):
    #Logistic Regression
    from sklearn.linear_model import LogisticRegression
    log = LogisticRegression(random_state = 0)
    log.fit(X_train, Y_train)

    #Random Forest Classifier
    from sklearn.ensemble import RandomForestClassifier
    forest = RandomForestClassifier(n_estimators = 50, criterion = 'entropy', random_state = 0)
    forest.fit(X_train, Y_train)

    #Print the models' accuracy on the training data
    print('Logistic Regression Training Accuracy:', log.score(X_train, Y_train))
    print('Random Forest Classifier:', forest.score(X_train, Y_train))

    return log, forest
```

```
In [ ]: model = models(X_train, Y_train)
```

Logistic Regression Training Accuracy: 0.92  
Random Forest Classifier: 1.0

## **SUMMARY**

The purpose of this project was to build a PCOS detecting Analyser. So, through our analysis and research, we've successfully made this possible to have built our project which would benefit when put to use. There's no test to definitively diagnose PCOS. The doctor is likely to start with a discussion of the patient's medical history, including the menstrual periods and weight changes. A physical exam will include checking for signs of excess hair growth, insulin resistance and acne. Awareness helps the public understand the above-mentioned symptoms which are not something to be ignored and getting it checked is necessary. Therefore, there's a need to have a device which would properly detect PCOS with accurate results. Hence, this analyser is designed which would serve for it as it gives 85-90% accuracy. We got to learn a lot of applications of Artificial Intelligence with great experiences.

## **REFERENCES**

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