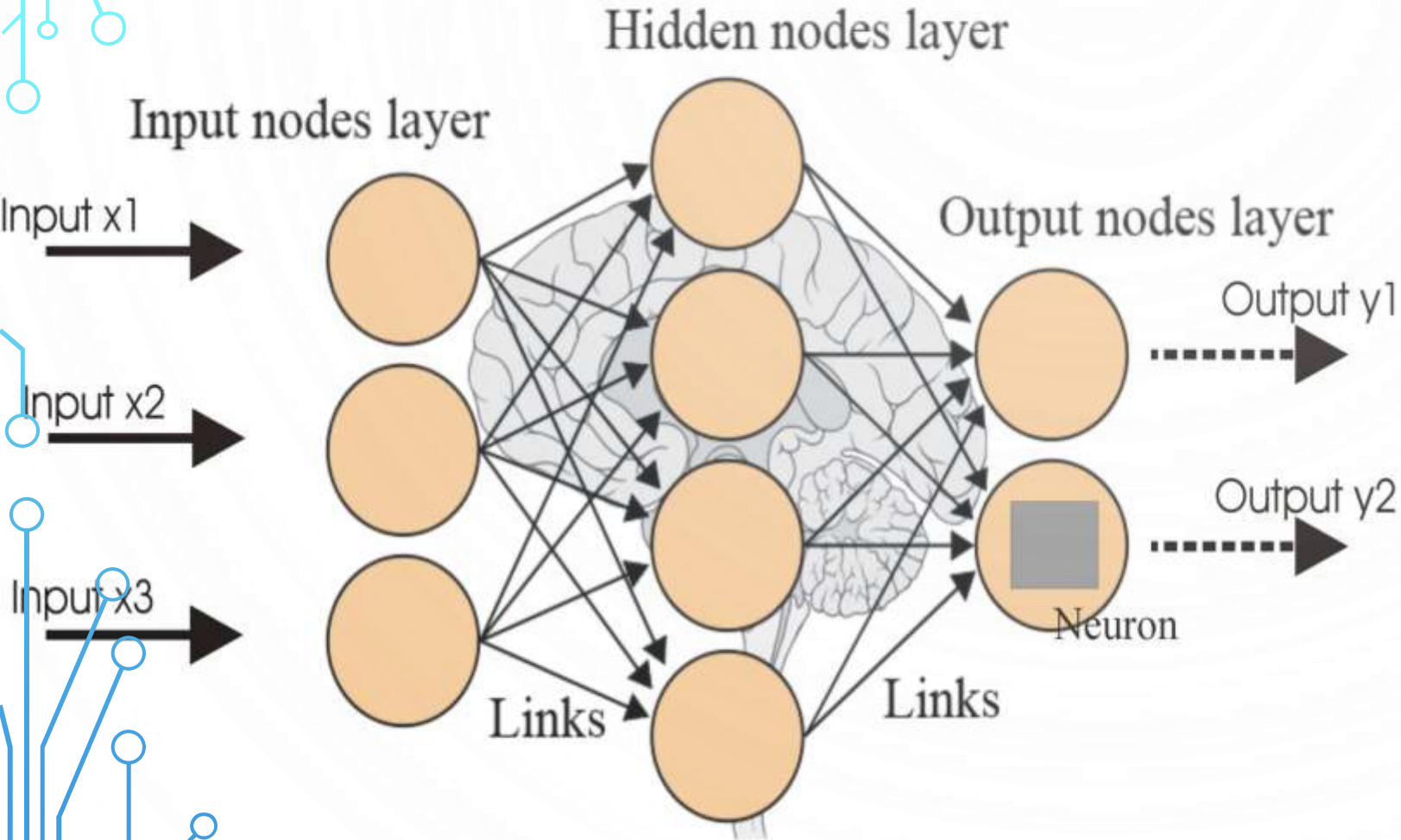


NEURAL NETWORK AND FUZZY CONTROL

- **Project: Predicting Diabetes with multilayer perceptron**
- **Under the guidance of: Dr. Sathish Kumar K**
- **Slot: D2**

Team Members	Registration number
Shreya Biswas	20BEI0072
Pratik Jain	20BEI0091
Pulkit Saraf	20BEI0092

ACKNOWLEDGEMENT



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INTRODUCTION

Diabetes is a chronic medical condition that is associated with elevated blood sugar levels in the body. Diabetes often leads to cardiovascular disease, stroke, kidney damage and long term damage to the extremities (i.e. limbs and eyes). It is estimated that there are 415 million people in the world suffering from diabetes, with up to 5 million deaths every year attributed to diabetes-related complications. Clearly, diabetes is a cause of concern to the wellbeing of modern society. Diabetes can be divided into two subtypes: type 1 and type 2

ABSTRACT

- **Type 1 diabetes results from the body's inability to produce enough insulin. It is relatively rare compared to type 2 diabetes, and it only accounts for 5% of diabetes. Unfortunately, the exact cause of type 1 diabetes is unknown and therefore, it is difficult to prevent the onset of type 1 diabetes. Type 2 diabetes results from the body's gradual resistance to insulin. Type 2 diabetes is the prevalent form of diabetes in the world, and it is caused by excessive body weight, irregular exercise, and a poor diet. Fortunately, the onset of type 2 diabetes can be prevented and reversed if diagnosed early. One of the barriers for early detection and diagnosis of diabetes is that the early stages of diabetes are often non-symptomatic. People who are on the path to diabetes (also known as prediabetes) often do not know they have diabetes until it is too late.**

THE DIABETES MELLITUS DATASET

The screenshot displays a Microsoft Excel spreadsheet titled "Pregnancies". The spreadsheet contains 30 rows of data, indexed from 1 to 30 in column A. The columns are labeled as follows:

- A: Pregnancies
- B: Glucose
- C: BloodPres
- D: SkinThickn
- E: Insulin
- F: BMI
- G: DiabetesP
- H: Age
- I: Outcome

The data represents various physiological measurements for pregnant women, used to predict the outcome (0 or 1). The interface shows the standard Excel ribbon with tabs for Home, Insert, Page Layout, Formulas, Data, Review, and View. The Font section is expanded, showing settings for Calibri font, size 11, bold, italic, underline, and color options.

Pregnancies	Glucose	BloodPres	SkinThickn	Insulin	BMI	DiabetesP	Age	Outcome
6	148	72	35	0	33.6	0.627	50	1
1	85	66	29	0	26.6	0.351	31	0
8	183	64	0	0	23.3	0.672	32	1
1	89	66	23	94	28.1	0.167	21	0
0	137	40	35	168	43.1	2.288	33	1
5	116	74	0	0	25.6	0.201	30	0
3	78	50	32	88	31	0.248	26	1
10	115	0	0	0	35.3	0.134	29	0
2	197	70	45	543	30.5	0.158	53	1
8	125	96	0	0	0	0.232	54	1
4	110	92	0	0	37.6	0.191	30	0
10	168	74	0	0	38	0.537	34	1
10	139	80	0	0	27.1	1.441	57	0
1	189	60	23	846	30.1	0.398	59	1
5	166	72	19	175	25.8	0.587	51	1
7	100	0	0	0	30	0.484	32	1
0	118	84	47	230	45.8	0.551	31	1
7	107	74	0	0	29.6	0.254	31	1
1	103	30	38	83	43.3	0.183	33	0
1	115	70	30	96	34.6	0.529	32	1
3	126	88	41	235	39.3	0.704	27	0
8	99	84	0	0	35.4	0.388	50	0
7	196	90	0	0	39.8	0.451	41	1
9	119	80	35	0	29	0.263	29	1
11	143	94	33	146	36.6	0.254	51	1
10	125	70	26	115	31.1	0.205	41	1
7	147	76	0	0	39.4	0.257	43	1
1	97	66	15	140	23.2	0.487	22	0

[illegible]

OVERVIEW OF THE GIVEN DATASET

1. Pregnancies – Number of previous pregnancies

2. Glucose – Plasma glucose concentration

3. Blood Pressure – Diastolic blood pressure

4. Skin Thickness – Skin fold thickness measured from the triceps

5. Insulin – Blood serum insulin concentration

6. BMI – Body Mass Index

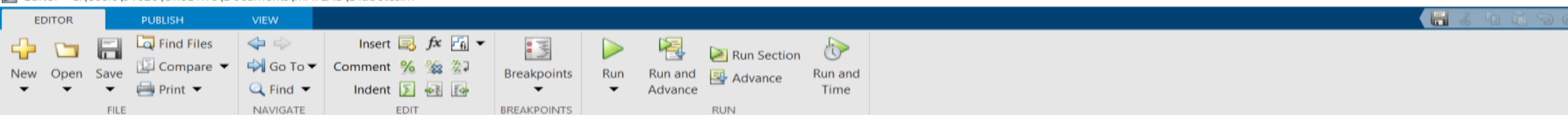
7. Diabetes Pedigree Function – A summarized score that indicates the genetic predisposition of the patient of diabetes, as extrapolated from the patient's family record for diabetes

8. Age – Age in years

9. Outcome – The target variable we are trying to predict, 1 for patients that developed diabetes within five years of the initial measurement, and 0 otherwise.

















CODE

Editor - C:\Users\91626\OneDrive\Documents\MATLAB\Diabetes.m



Diabetes.m

```
1 %Read data
2 clc
3 clear all
4 close all
5 warning off
6 data=readtable('Dia.csv');
7 %%
8 %Checking if null value present or not
9 a=sum(ismissing(data));
10 %%
11 %Feature Scaling
12 data.Pregnancies=(data.Pregnancies-min(data.Pregnancies))/(max(data.Pregnancies)-min(data.Pregnancies));
13 data.Glucose=(data.Glucose-min(data.Glucose))/(max(data.Glucose)-min(data.Glucose));
14 data.BloodPressure=(data.BloodPressure-min(data.BloodPressure))/(max(data.BloodPressure)-min(data.BloodPressure));
15 data.SkinThickness=(data.SkinThickness-min(data.SkinThickness))/(max(data.SkinThickness)-min(data.SkinThickness));
16 data.Insulin=(data.Insulin-min(data.Insulin))/(max(data.Insulin)-min(data.Insulin));
17 data.BMI=(data.BMI-min(data.BMI))/(max(data.BMI)-min(data.BMI));
18 data.DiabetesPedigreeFunction=(data.DiabetesPedigreeFunction-min(data.DiabetesPedigreeFunction))/(max(data.DiabetesPedigreeFunction)-min(data.DiabetesPedigreeFunction));
19 data.Age=(data.Age-min(data.Age))/(max(data.Age)-min(data.Age));
20 %%
21 %Split data into train test
22 cv = cvpartition(size(data,1),'HoldOut',0.3);
23 idx = cv.test;
24 % Separate to training and test data
25 dataTrain = data(~idx,:);
26 dataTest = data(idx,:);
27 %%
28 %Handle the 0 values for some features of training set
29 g=(dataTrain.Glucose==0);
30 k=(dataTrain.BloodPressure==0);
31 l=(dataTrain.SkinThickness==0);
32 m=(dataTrain.Insulin==0);
33 n=(dataTrain.BMI==0);
```

















EDITOR			PUBLISH		VIEW													
																		
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FILE			NAVIGATE			EDIT			BREAKPOINTS			RUN						

Diabetes.m

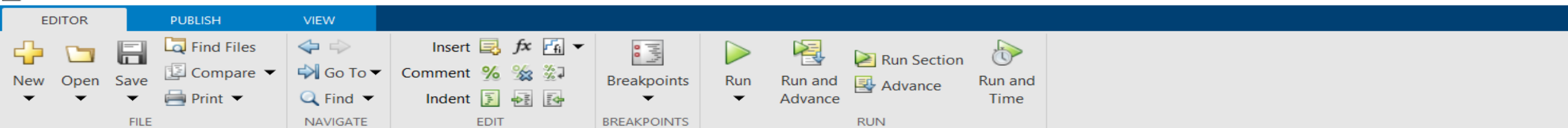
```

34 - o=(dataTrain.DiabetesPedigreeFunction==0);
35 - p=(dataTrain.Age==0);
36 - impractical=[sum(g) sum(k) sum(l) sum(m) sum(n) sum(o) sum(p)];
37 - %%
38 - %Replace those with average of the corresponding columns of training data
39 - g=(dataTrain.Glucose==0);
40 - msa=(dataTrain.Outcome==1);
41 - ks=mean(dataTrain.Glucose(~g & msa));
42 - dataTrain.Glucose(g & msa)=ks;
43 - ks=mean(dataTrain.Glucose(~g & ~msa));
44 - dataTrain.Glucose(g & ~msa)=ks;
45 - g=(dataTrain.BloodPressure==0);
46 - ks=mean(dataTrain.BloodPressure(~g & msa));
47 - dataTrain.BloodPressure(g & msa)=ks;
48 - ks=mean(dataTrain.BloodPressure(~g & ~msa));
49 - dataTrain.BloodPressure(g & ~msa)=ks;
50 - g=(dataTrain.SkinThickness==0);
51 - ks=mean(dataTrain.SkinThickness(~g & msa));
52 - dataTrain.SkinThickness(g & msa)=ks;
53 - ks=mean(dataTrain.SkinThickness(~g & ~msa));
54 - dataTrain.SkinThickness(g & ~msa)=ks;
55 - g=(dataTrain.Insulin==0);
56 - ks=mean(dataTrain.Insulin(~g & msa));
57 - dataTrain.Insulin(g & msa)=ks;
58 - ks=mean(dataTrain.Insulin(~g & ~msa));
59 - dataTrain.Insulin(g & ~msa)=ks;
60 - g=(dataTrain.BMI==0);
61 - ks=mean(dataTrain.BMI(~g & msa));
62 - dataTrain.BMI(g & msa)=ks;
63 - ks=mean(dataTrain.BMI(~g & ~msa));
64 - dataTrain.BMI(g & ~msa)=ks;
65 - g=(dataTrain.DiabetesPedigreeFunction==0);
66 - ks=mean(dataTrain.DiabetesPedigreeFunction(~g & msa));

```


EDITOR			PUBLISH		VIEW										
 New	 Open	 Save	 Find Files	 Compare	 Go To	 Find	 Insert	 Comment	 Indent	 Breakpoints	 Run	 Run and Advance	 Run Section	 Advance	 Run and Time
FILE					NAVIGATE		EDIT		BREAKPOINTS		RUN				

Diabetes.m	
67 -	dataTrain.DiabetesPedigreeFunction(g & msa)=ks;
68 -	ks=mean(dataTrain.DiabetesPedigreeFunction(~g & ~msa));
69 -	dataTrain.DiabetesPedigreeFunction(g & ~msa)=ks;
70 -	dataTrain.Age(p)=mean(dataTrain.Age(~p));
71 -	%%
72 -	%Handle the 0 values for some features of test set
73 -	g=(dataTest.Glucose==0);
74 -	k=(dataTest.BloodPressure==0);
75 -	l=(dataTest.SkinThickness==0);
76 -	m=(dataTest.Insulin==0);
77 -	n=(dataTest.BMI==0);
78 -	o=(dataTest.DiabetesPedigreeFunction==0);
79 -	p=(dataTest.Age==0);
80 -	impractical=[sum(g) sum(k) sum(l) sum(m) sum(n) sum(o) sum(p)];
81 -	%%
82 -	%Replace those with average of the corresponding columns of test data
83 -	g=(dataTest.Glucose==0);
84 -	msa=(dataTest.Outcome==1);
85 -	ks=mean(dataTest.Glucose(~g & msa));
86 -	dataTest.Glucose(g & msa)=ks;
87 -	ks=mean(dataTest.Glucose(~g & ~msa));
88 -	dataTest.Glucose(g & ~msa)=ks;
89 -	g=(dataTest.BloodPressure==0);
90 -	ks=mean(dataTest.BloodPressure(~g & msa));
91 -	dataTest.BloodPressure(g & msa)=ks;
92 -	ks=mean(dataTest.BloodPressure(~g & ~msa));
93 -	dataTest.BloodPressure(g & ~msa)=ks;
94 -	g=(dataTest.SkinThickness==0);
95 -	ks=mean(dataTest.SkinThickness(~g & msa));
96 -	dataTest.SkinThickness(g & msa)=ks;
97 -	ks=mean(dataTest.SkinThickness(~g & ~msa));
98 -	dataTest.SkinThickness(g & ~msa)=ks;
99 -	g=(dataTest.Insulin==0);



```

Diabetes.m
91 - dataTest.BloodPressure(g & msa)=ks;
92 - ks=mean(dataTest.BloodPressure(~g & ~msa));
93 - dataTest.BloodPressure(g & ~msa)=ks;
94 - g=(dataTest.SkinThickness==0);
95 - ks=mean(dataTest.SkinThickness(~g & msa));
96 - dataTest.SkinThickness(g & msa)=ks;
97 - ks=mean(dataTest.SkinThickness(~g & ~msa));
98 - dataTest.SkinThickness(g & ~msa)=ks;
99 - g=(dataTest.Insulin==0);
100 - ks=mean(dataTest.Insulin(~g & msa));
101 - dataTest.Insulin(g & msa)=ks;
102 - ks=mean(dataTest.Insulin(~g & ~msa));
103 - dataTest.Insulin(g & ~msa)=ks;
104 - g=(dataTest.BMI==0);
105 - ks=mean(dataTest.BMI(~g & msa));
106 - dataTest.BMI(g & msa)=ks;
107 - ks=mean(dataTest.BMI(~g & ~msa));
108 - dataTest.BMI(g & ~msa)=ks;
109 - g=(dataTest.DiabetesPedigreeFunction==0);
110 - ks=mean(dataTest.DiabetesPedigreeFunction(~g & msa));
111 - dataTest.DiabetesPedigreeFunction(g & msa)=ks;
112 - ks=mean(dataTest.DiabetesPedigreeFunction(~g & ~msa));
113 - dataTest.DiabetesPedigreeFunction(g & ~msa)=ks;
114 - dataTest.Age(p)=mean(dataTest.Age(~p));
115 - %%
116 - %Train model using training sets
117 - classification_model=fitcsvm(dataTrain,'Outcome~Pregnancies+Glucose+BloodPressure+SkinThickness+Insulin+BMI+DiabetesPedigreeFunction+Age');
118 - %%
119 - %Accuracy
120 - gs=dataTest(:,1:8);
121 - mk=predict(classification_model,gs);
122 - accuracy_check=(sum(mk==table2array(dataTest(:,9))))/size(dataTest,1)*100;
123 - disp(accuracy_check);
  
```


OUTPUT

MATLAB R2020b - academic use

HOME PLOTS APPS

New Script New Live Script New Open Find Files Compare Import Data Save Workspace New Variable Open Variable Clear Workspace Favorites Analyze Code Run and Time Clear Commands Simulink Layout Preferences Set Path Add-Ons Help Community Request Support Learn MATLAB

FILE VARIABLE CODE SIMULINK ENVIRONMENT RESOURCES

C:\Users\91626\OneDrive\Documents\MATLAB

73.0435

fx >> |

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80

fx >>

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81.3043

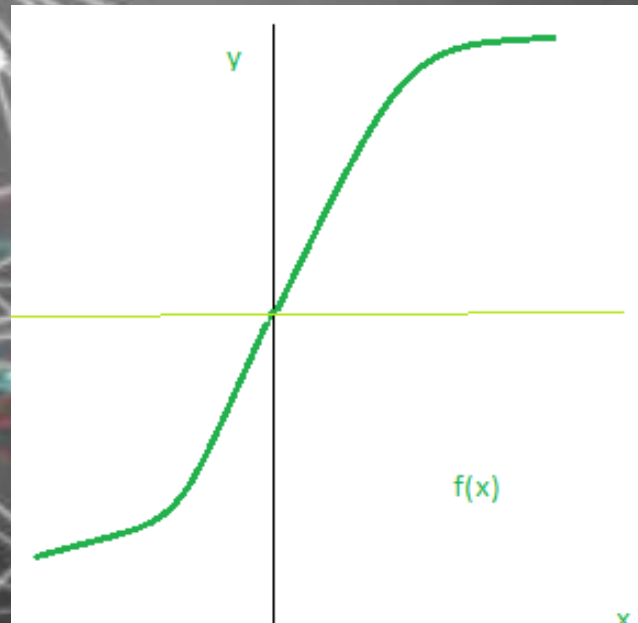
fx >>

ACTIVATION FUNCTION

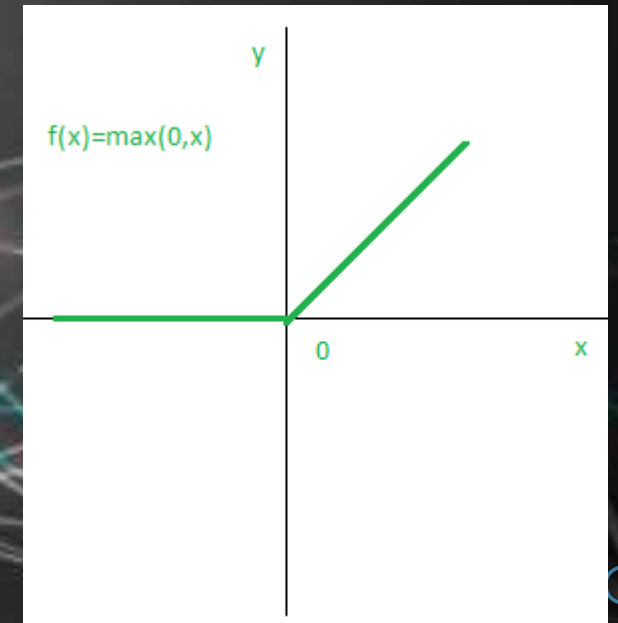
In this project, we will use the rectified linear unit (ReLU) and the sigmoid as our activation functions. ReLU is always used as the activation function for our intermediate hidden layers.

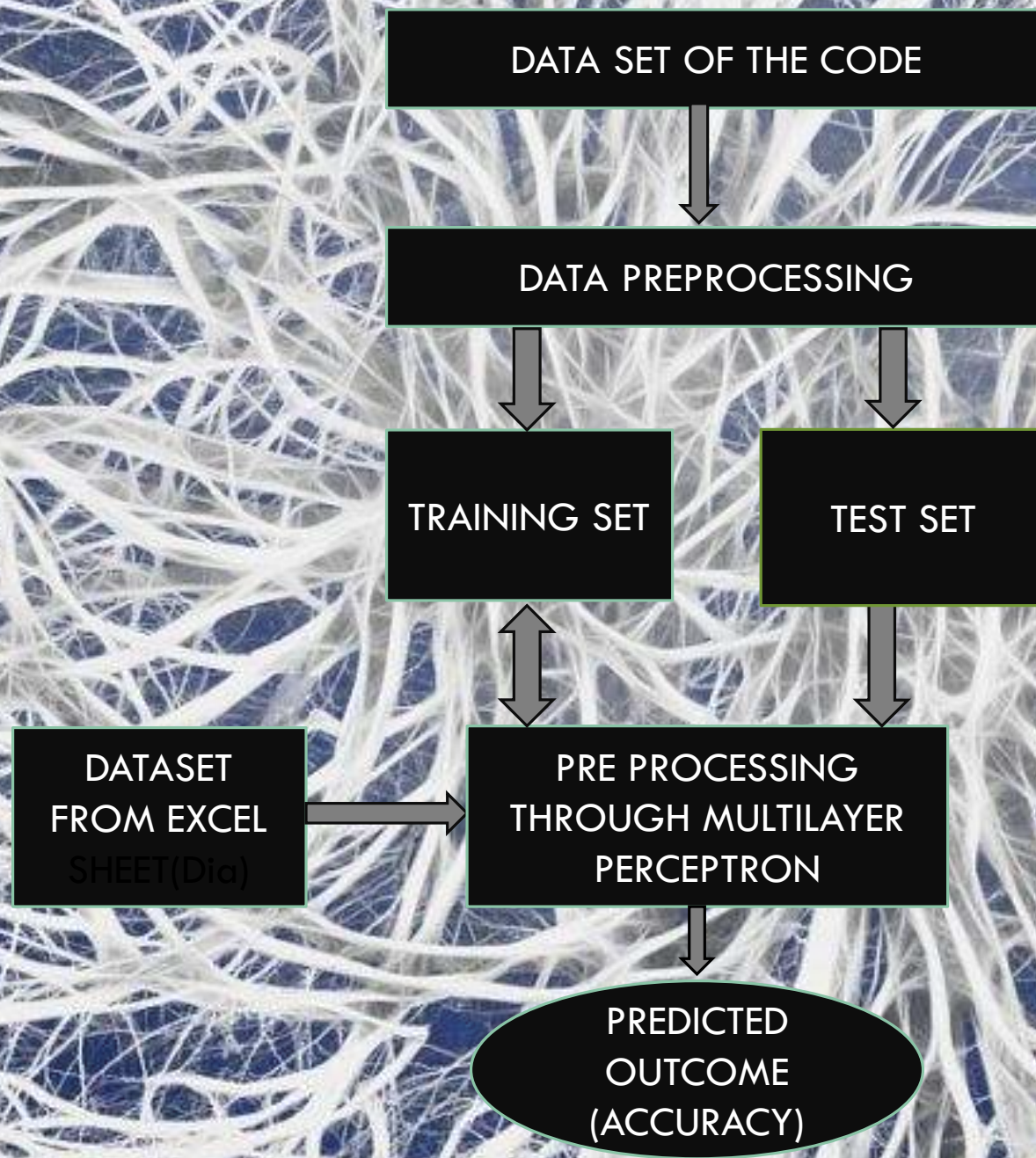
Mathematically, it can be represented as: $f(x) = \max(0, x)$
Sigmoid is used as the activation function for the output layer.
Mathematically, it can be represented as: $F(x) = 1/(1+e^{-x})$

SIGMOID FUNCTION



ReLU FUNCTION





REFERENCE

1. Anjali Negi and Varun Jaiswal, “A first attempt to develop a diabetes prediction method based on different global datasets.”
2. Jatin N Bagrecha, Chaithra G S and Jeevitha S, “Diabetes Disease Prediction using Neural Networks.”
3. Sumi Alice Saji and K Balachandran, “Performance Analysis of training algorithms of multilayer perceptrons in diabetes prediction.”
4. Talha Mehboob Alam, Muhammad Atif Iqbal, Yasir Ali, “A model for early prediction for diabetes.”
5. James Loy, “Neural Networks projects with Python.”



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Y₄ O₁ U₁