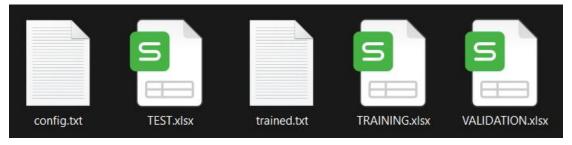
# **How This ANN works**

'train.py' for training the ANN and 'function.py' for using trained ANN >> all other files are modules made to be used

All usage related files are to be placed in '//FILES' sub-folder



#### **FILES contains:**



config.txt: contains info on the ANN structure and Functions

TEST, TRAINING, VALIDATION: contain sample data

trained.txt: contains the trained model saved by the training process

Prerequisites: numpy, matplotlib, openpyxl

### **CONFIG FILE:**

```
f = open('files\\config.txt', "r")

L = int(f.readline())
N = int(f.readline())
Af = int(f.readline())
I = int(f.readline())
0 = int(f.readline())

B = int(f.readline())
EP = int(f.readline())
Irate = float(f.readline())
mom = float(f.readline())
reg = float(f.readline())
```

L - Number of Hidden Layers

N - Neurons per Hidden Layer

Af - Activation Function (1: tanh, 2: logistic, 3: ReLU)

I - Number of inputs

O - Number of Outputs

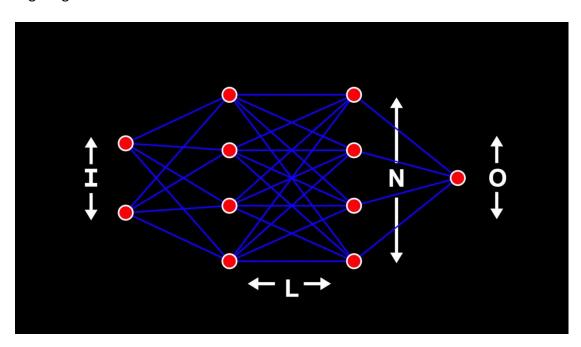
B - Batch size

EP - Number of Epochs

Irate - learning rate parameter

mom - momentum

reg - regularization constant



## **TRAINING (DATA) FILEs:**

4	Α	В	С	D	Е
1	14.96	41.76	1024.07	73.17	463.26
2	25.18	62.96	1020.04	59.08	444.37
3	5.11	39.4	1012.16	92.14	488.56
4	20.86	57.32	1010.24	76.64	446.48
5	10.82	37.5	1009.23	96.62	473.9
6	26.27	59.44	1012.23	58.77	443.67
7	15.89	43.96	1014.02	75.24	467.35

- > The training data needs to be in **xlsx** files without any labels
- > The first I (from config file) elements must be inputs
- > The last **O** (from config file) elements must be outputs

# e.g. data sample from CCPP dataset

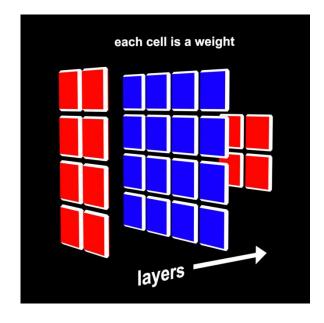
14.96	41.76	1024.07	73.17	463.26
			The second second second	THE DESIGNATION OF SECURITY

First 4 value(s) are input Last 1 value(s) is output

## How 'train.py' works:

#### **Initialization:**

Initialize Neurons, Weights, Biases, Delta Weights and Delta Biases



Weights are stored as such: A list of matrices (2D arrays)

Biases/Neuron activations are stored as such: A list of 1D arrays

## **Loading samples:**

Training and Validation data is loaded via a inputreader module The samples are min-max mapped between (0,1)

name I contains inputs, and name o contains outputs

## **Training:**

```
for ep in range(EP):
```

Runs for as many epochs as specified

```
seed = random.random()

random.Random(seed).shuffle(train_i)
random.Random(seed).shuffle(train_o)
```

Shuffles data at every epoch

```
BAn[i%B] = fwd.output(W, An, O, Af, v, L, Bias)
BErr[i%B] = fwd.erroratout(train_o[i], BAn[i%B][L+1])
```

Performs function calculations and then finds error for a batch

```
if(i%B == (B-1)):
    An = aavg.batchaverage(BAn, An, B)
    Err = aavg.batcherr(BErr, Err, B)

    (dW, dB) = backp.backprop(dW, dB, W, Bias, L, An, Err, 1rate, mom, Af, I, N, 0, reg)

    (W, Bias) = backp.upW(W, Bias, dW, dB, L)
```

At the end of each batch

- >Average activations and Average errors are calculated
- >Change in Weights and Biases is calculated (back-prop)
- >Weights and Biases are updated
- >Weights corresponding to the **Lowest validation error** are saved

Error% vs Epochs is plotted

#### **Batch Average:**

Activation:

For all batches:

nodeAvg = sum(nodes)/batchSize

Error:

For all batches:

errorAvg = sum(errors)/batchSize

#### **Activation Functions:**

Activations are calculated by the activation module Which has 3 main features:

> actfunc | activation function

> iactfunc | inverse activation function

> dactfunc | derivative of activation function

All three take 2 parameters:

1. Value | value to be calculated upon

2. Func | which function to use

#### **Function Calculations:**

Function calculations are done in the 'forward.py' module >in output function as follow

For layers in [0, **L+2**):

If(layer = input layer):

Activations = Input

Else:

Activations = act(WxA[I-1] + B) #act is activation function

Error is calculated by vector subtraction

>in **erroratout** function

#### **BackProp Calculations:**

Local Gradient is calculated for all nodes using equations written in (A)

if 
$$(L \text{ is outlayerr})$$

$$\left\{ e_{L} = d_{L} - Y_{L} \right\}$$
else
$$\left\{ e_{L} = \left\{ \int_{L+1}^{L+1} W_{L+1}(L) \right\} \right\}$$

Once local gradient is calculated, change In Weights and Biases is calculated

(Momentum and Regularization implemented in code)

M = np.size(dW[i])

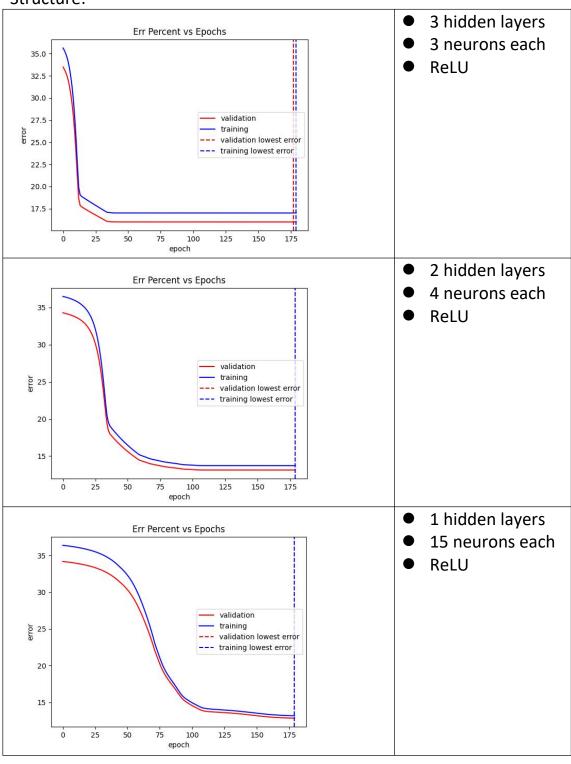
```
\frac{dW[i][j][k] = mom*dW[i][j][k] + (1-mom)*lrate*An[i][j]*L6[i+1][k] - (reg*dW[i][j][k]/M)}{dW[i][j][k] + (1-mom)*lrate*An[i][j][k] + (1-mom)*lrate*An[i][k] + (1-mom)*lrate*An
```

```
M = np.size(dB[i])
dB[i][j] = mom*dB[i][j] + (1-mom)*lrate*L6[i][j] - (reg*dB[i][j]/M)
```

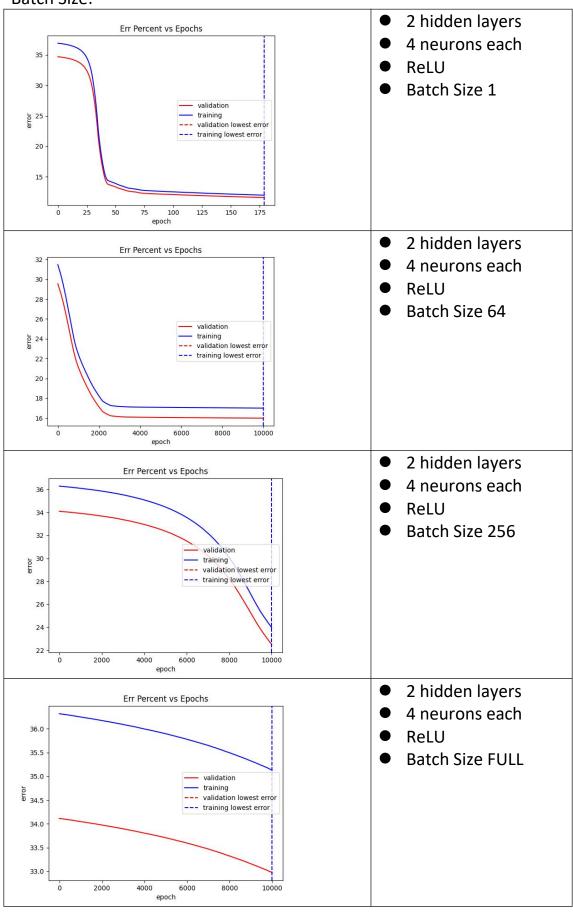
Saving Data:
Data is separated by new line and is labelled:
Data in same layer is placed between [ ]
Connection every neuron has with next layer neurons is between (only for weights) {}

# Impact of parameters:

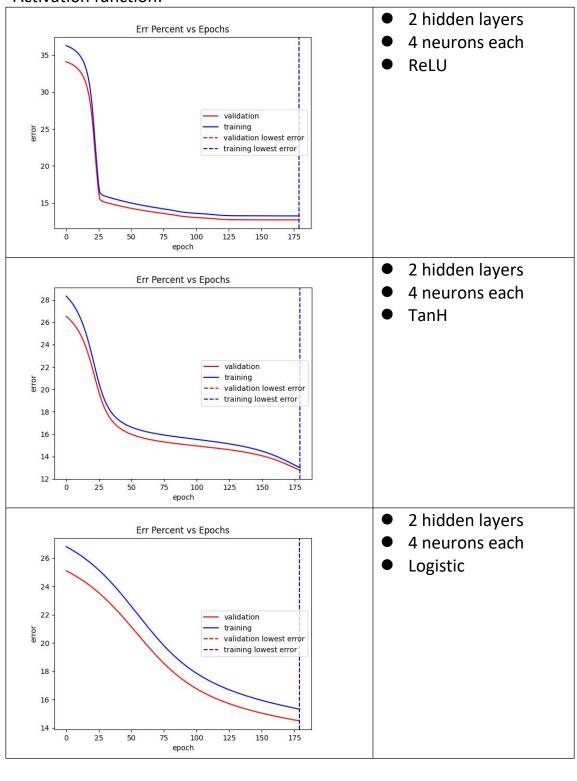
#### Structure:



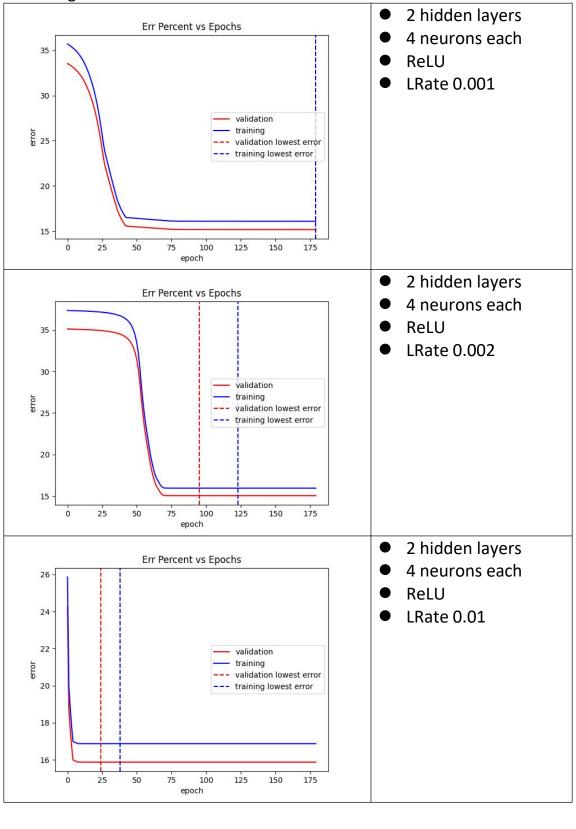
#### Batch Size:



### Activation function:



### Learning Rate:



### Momentum:

