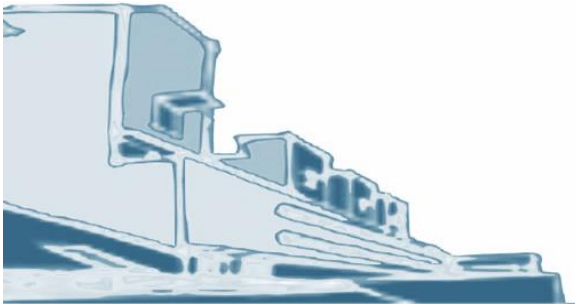


P4b – Multilayer neural networks

Jorge Henriques

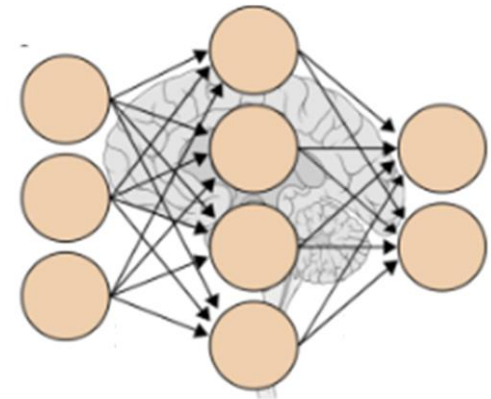
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informática



Contents

- 1 | Objectives
- 2 | Datasets
- 3 | Tasks
- 4 | Conclusions

- MLNN - Multi Layer neural networks
 - Concepts
 - Implement **BackPropagation** from scratch
 - Regression - Approximate a non-linear function
 - Use **python/scikit learning** functionalities
 - Classification
 - Regression
 - Evaluation the performance of the MLNN classifier/regressor
 - SE, SP, F1score (*classification*)
 - MSE, R2 (*regression*)

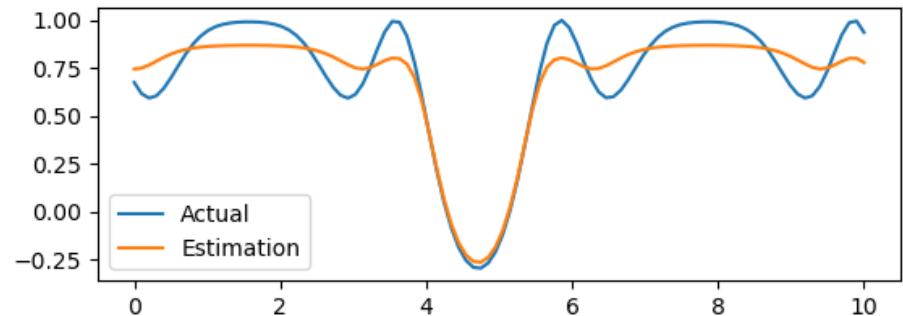
Contents

- 1 | Objectives
- 2 | Datasets
- 3 | Tasks
- 4 | Conclusions

■ Datasets

• 1 | Function approximation

- Simple problem – regression
- P4_function.csv



• 2 | Cardiac Risk

- **Classification**
- P4_cardiacRiskClassification.csv
- Output = {0,1} – {event, NO event}

• 3 | Cardiac Risk

- **Regression**
- P4_cardiacRiskRegression.csv
- Output = [0..1] – **probability to have an event**

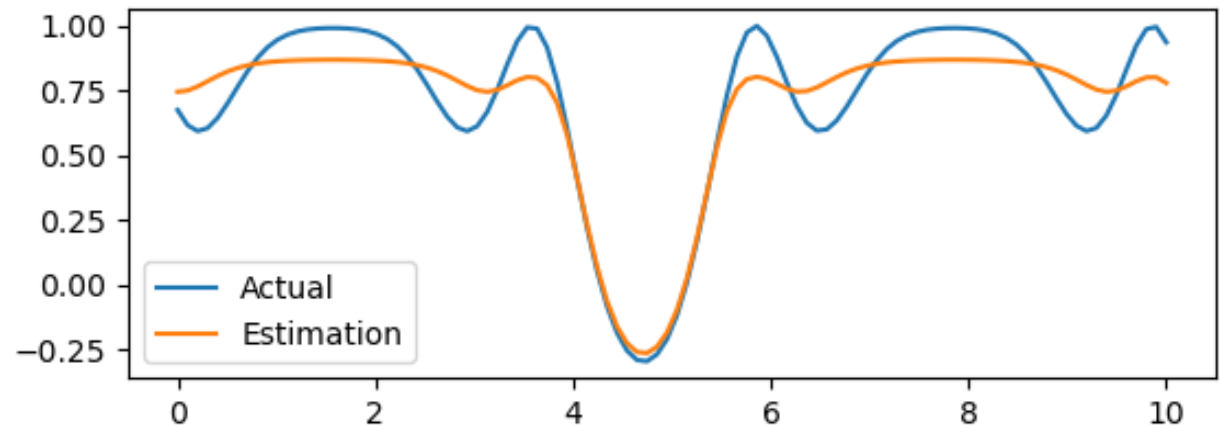
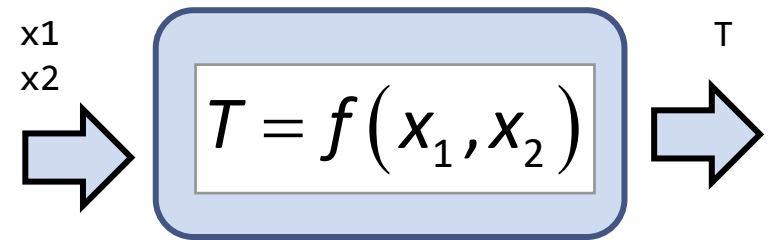


■ 1 | Datasets – function approximation

- Non-linear function

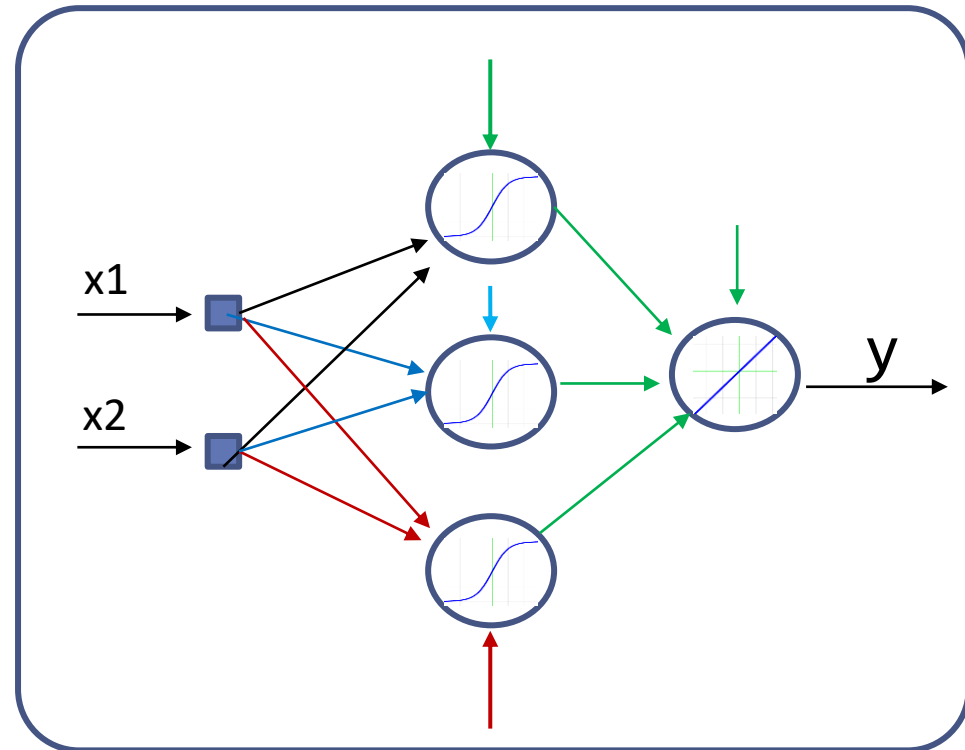
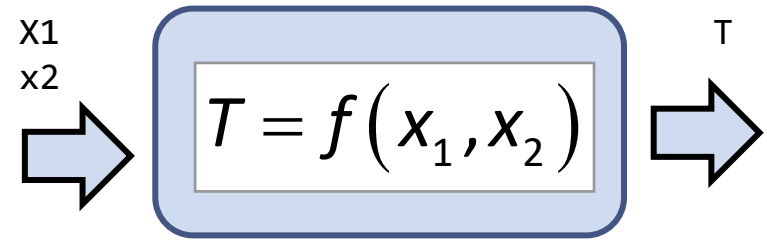
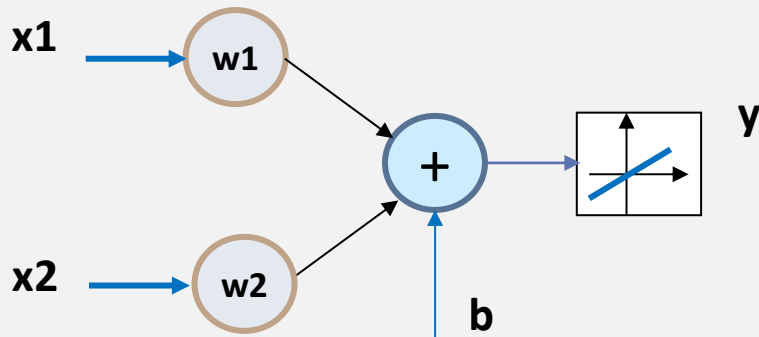
```
t1= 0:0.1:10  
x1= sin(t1);  
x2= sin(t).cos(t)^22;  
T = x1 + 2.4*x2;
```

P4_function.csv



■ 1| Function approximation

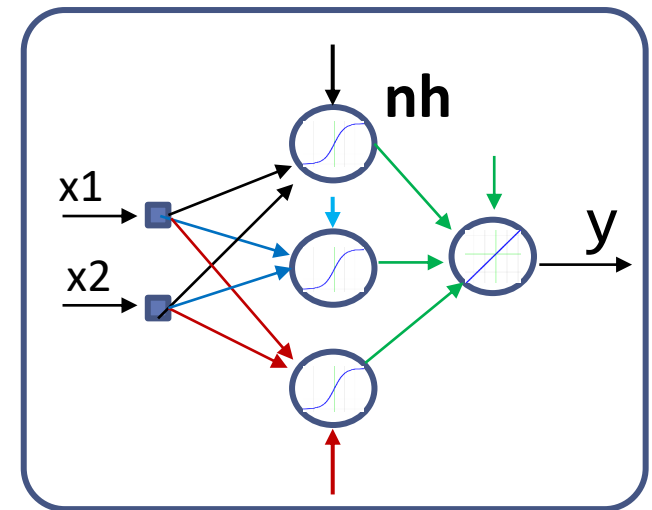
- Two possible structures
 - 1. Adaline : one layer
 - 2. MLNN : hidden layer
- Learning:
 - 1. RMLSE
 - 2. Backpropagation



■ 2 | Datasets - Cardiac Risk

■ Use of scikit learning functionalities

- Structure: MLNN – one hidden layer
- Activation functions
 - Hidden layer – sigmoidal (nh neurons)
 - Output layer – linear (one neuron)
- Classification $T=\{0,1\}$
- Regression $T=[0..1]$

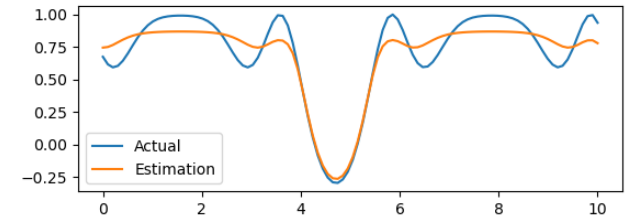


Contents

- 1 | Objectives
- 2 | Datasets
- 3 | Tasks
- 4 | Conclusions

■ 1 | Function approximation

P4_function.csv



1.1 | Structure

Activation function

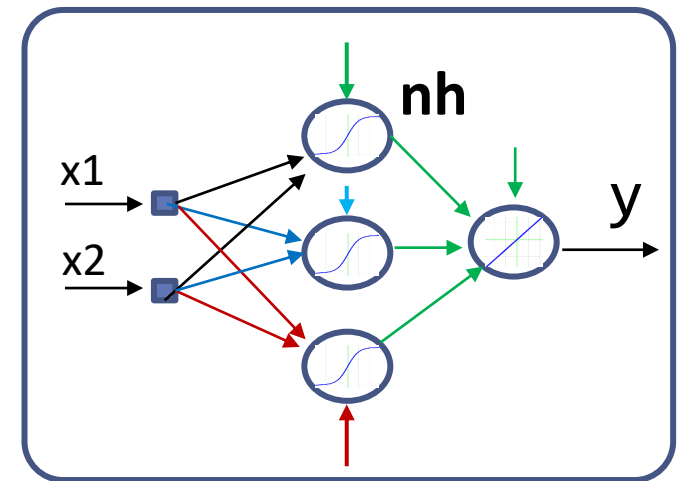
- Sigmoidal / Linear

Number of hidden layers

- $nh = ?$

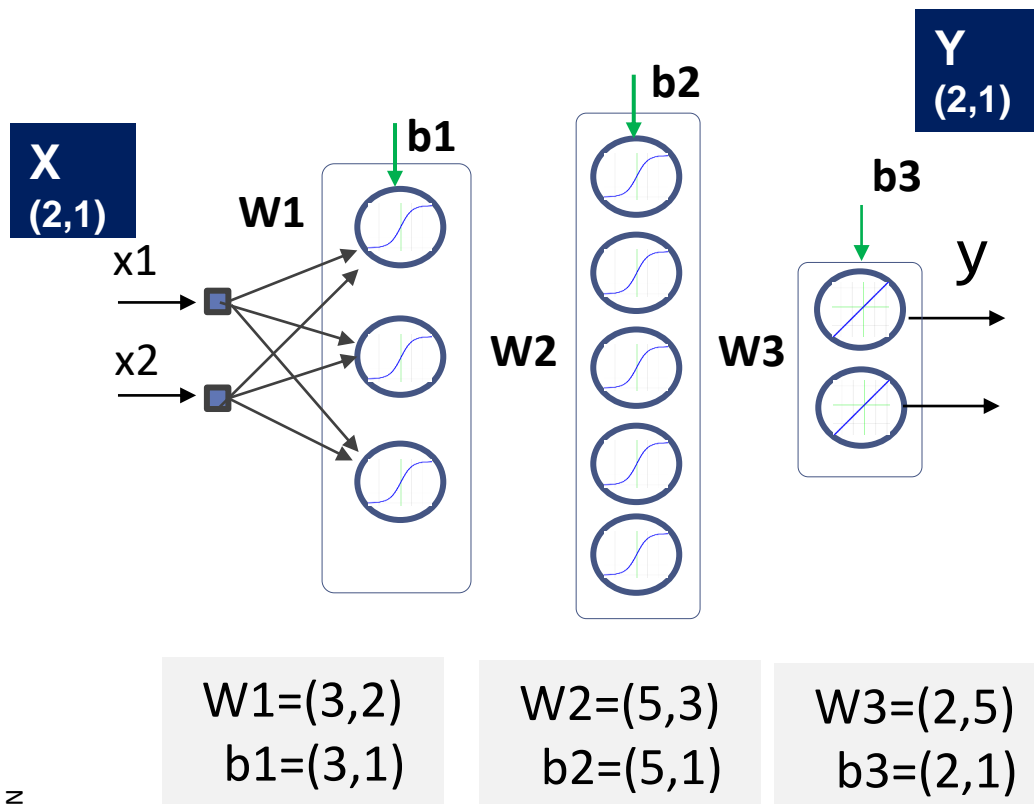
1.2 | Learning: Back Propagation

- Perform the computations by hand !



1 | BP: example

FORWARD



$$a^0 = x$$

$$n^1 = W^1 a^0 + b^1 \quad (3,2)(2,1) + (3,1)$$

$$a^1 = f^1(n^1) \quad (3,1)$$

$$n^2 = W^2 a^1 + b^2 \quad (5,3)(3,1) + (5,1)$$

$$a^2 = f^2(n^2) \quad (5,1)$$

$$n^3 = W^3 a^2 + b^3 \quad (2,5)(5,1) + (2,1)$$

$$a^3 = f^3(n^3) \quad (2,1)$$

$$y = a^3$$

■ Last layer ($m=3$)

$$n^3 = W^3 a^2 + b^3$$

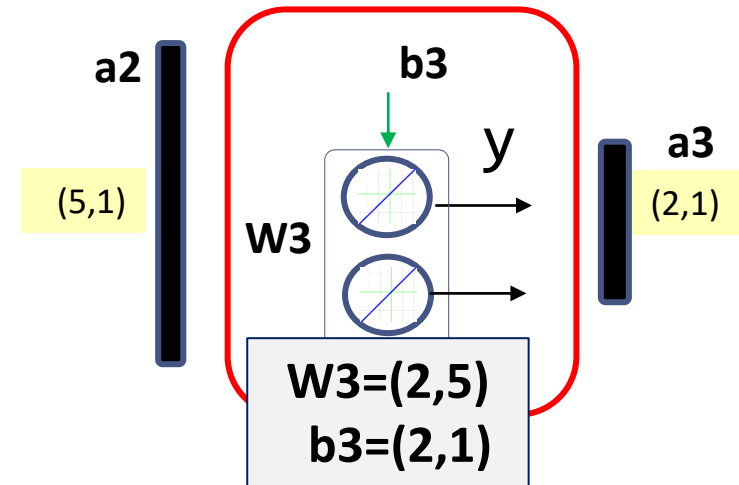
$$a^3 = f^3(n^3)$$

$$e = e^3 = t - a^3 = t - y$$

$$\dot{f}(n^3) = df^3(\cdot)$$

$$\frac{dE}{dw^3} = \frac{d(e^3)^2}{dw^3} = -2 e^3 f^3(\cdot) (a^2)^T$$

$$\frac{dE}{db^3} = \frac{d(e^3)^2}{db^3} = -2 f^3(\cdot) e^3$$



$$s^3 = e^3 \otimes df^3(\cdot) \quad (2,1) \otimes (2,1)$$

$$dW^3 = s^3 (a^2)^T \quad (2,1) (1,5)$$

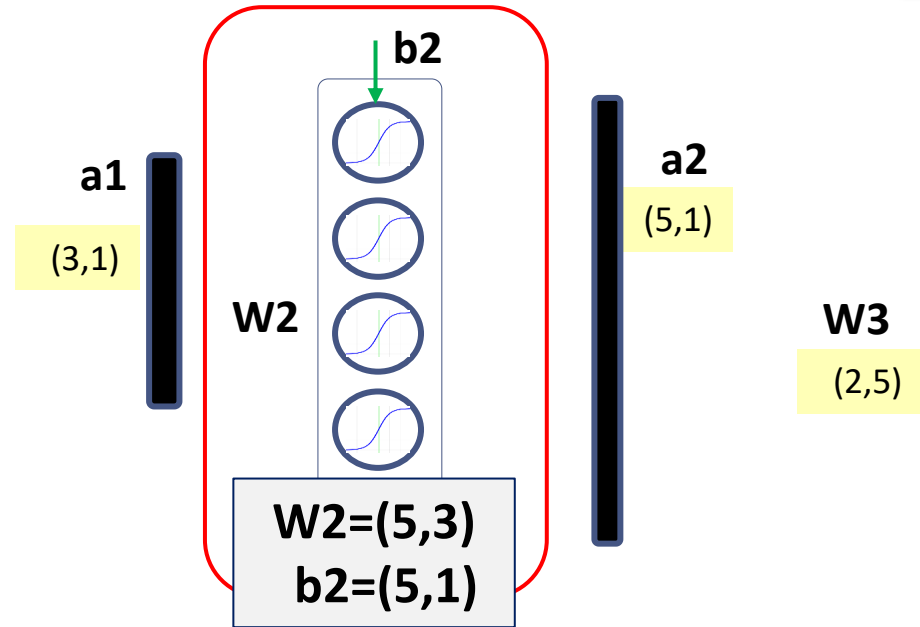
$$db^3 = s^3 \quad (2,1)$$

■ Hidden layer ($m=2$)

$$n^2 = W^2 a^1 + b^2$$

$$a^2 = f^2(n^2)$$

$$s^{m-1} \equiv (w^m)^T s^m \dot{f}(n^{m-1})$$



$$\dot{f}(n^2) = df^2(\cdot) \quad (5,1)$$

$$dW^2 = s^2 (a^1)^T \quad (5,1) (1,3) = (5,3)$$

$$db^2 = s^2 \quad (5,1)$$

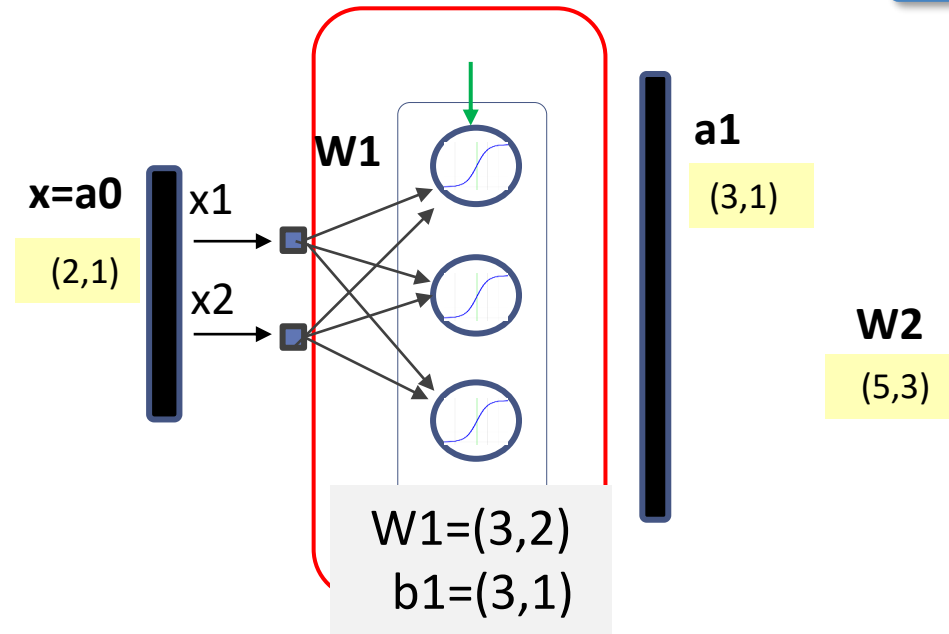
$$s^2 = (w^3)^T s^3 \otimes df^2(\cdot) \quad ((5,2)(2,1)) \otimes (5,1)$$

- First layer ($m=1$)

$$n^1 = W^1 a^0 + b^1$$

$$a^1 = f^1(n^1)$$

$$s^{m-1} \equiv (w^m)^T s^m \dot{f}(n^{m-1})$$



$$\dot{f}(n^1) = df^1(\cdot)$$

(3,1)

$$dW^1 = s^1 (a^0)^T$$

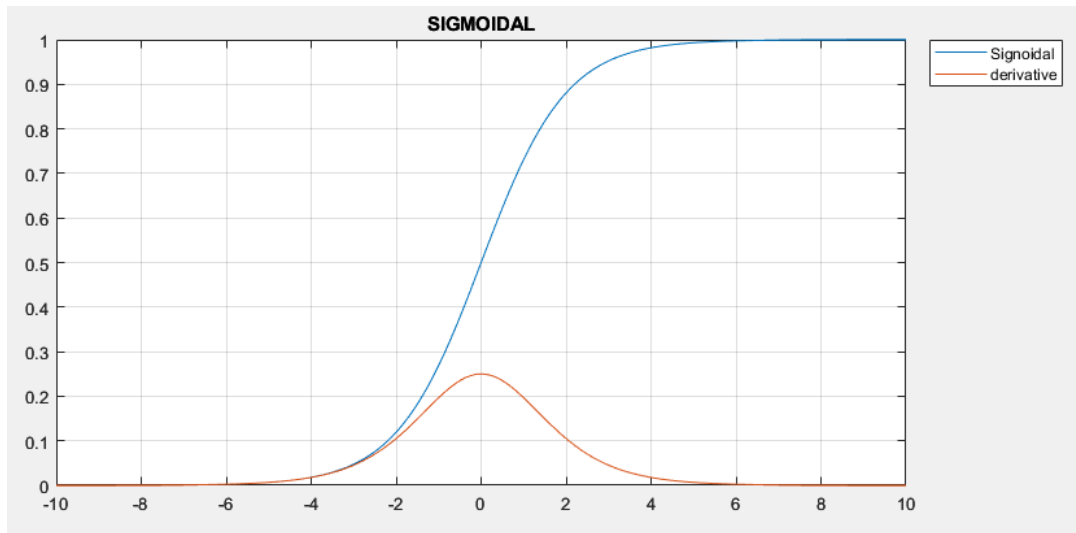
(3,1) (1,2) = (3,2)

$$s^1 = (w^2)^T s^2 \otimes df^1(\cdot)$$

((3,5)(5,1)) \otimes (3,1)

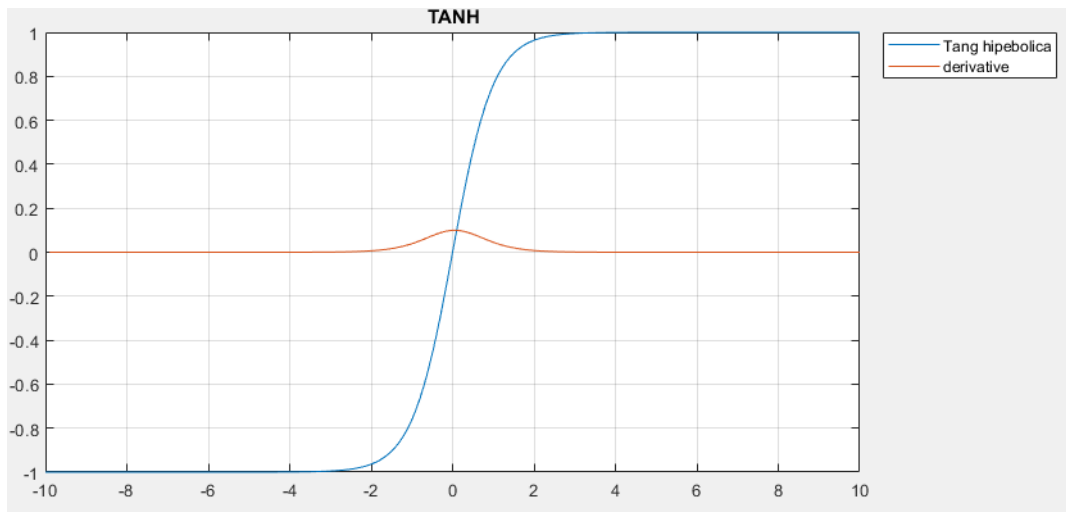
$$db^1 = s^1$$

(3,1)



$$f(x) = \frac{1}{1 + \exp(-x)}$$

$$df(x) = f(x)(1 - f(x))$$



$$f(x) = \frac{\exp(x) - \exp(-x)}{\exp(x) + \exp(-x)}$$

1 | Function approximation

P4_function.csv

```
alfa=0.001
```

```
W2=np.random.random((ny, nh1))
```

```
B2=np.random.random((ny, 1))
```

```
W1=np.random.random((nh1, nx))
```

```
B1=np.random.random((nh1, 1))
```

```
for numEP in range(EPOCHS):
```

```
#----- FORWARD
```

```
a0 = X
```

```
a1 = np.dot(W1,a0) + B1
```

```
y1 = sigmoid(a1); # sigmoidal
```

```
a2 = np.dot(W2,a1) + B2
```

```
y2= a2 # linear
```

```
#----- ERROR
```

```
e2 = T - y2
```

```
erro = np.dot( e2, e2.transpose()) # for each EPOCH
```

```
#----- BACKPROPAGATION
```

```
dy2 = 1
```

```
S2 = e2*dy2
```

```
dW2 = alfa*np.dot(S2, a1.transpose())
```

```
dB2 = alfa*np.dot(S2, MI)
```

```
W2 = W2+ dW2
```

```
B2 = B2+ dB2
```

```
# . . . . .
```

```
dy1 = y1*(1-y1)
```

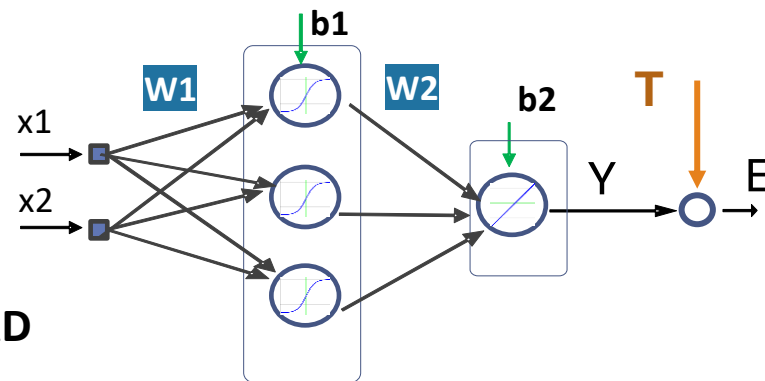
```
S1 = np.dot( W2.transpose(), S2)*dy1
```

```
dW1 = alfa* np.dot(S1, a0.transpose())
```

```
dB1 = alfa*np.dot(S1, MI)
```

```
W1 = W1+ dW1
```

```
B1 = B1 + dB1
```



FORWARD

$$f_1 = \text{sigmoidal}(x)$$

$$\dot{f}_1 = f_1(1 - f_1) = 1 - f_1^2$$

 $\otimes = *$
 $x = \text{np.dot}(.)$

element by element
multiplication

BACKPROPAGATION

- **2 | Dataset: Cardiac risk / P4_cardiacRiskClassification.csv**
- **2.1 | Implement MLNN - classification**
 - Structure
 - Number of neurons = ?
 - Activation functions : Hidden layer / Output layer (**sigmoidal/linear**)
 - Training / test data
 - Learning algorithm (**adam**)

```
from sklearn.neural_network import MLPClassifier

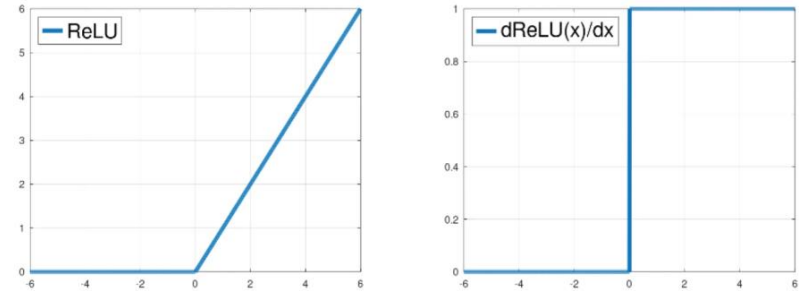
Xtrain, Xtest, Ttrain, Ttest = train_test_split(X,T,test_size = 0.3,
random_state = 42)

mlp = MLPClassifier(hidden_layer_sizes=(3,2), activation='relu',
                    solver='adam', max_iter=500)

mlp.fit(Xtrain,Ttrain)
```

■ RLU

$$f(x) = \max(0, x) = \begin{cases} x & \text{if } x > 0, \\ 0 & \text{otherwise.} \end{cases} \quad f'(x) = \begin{cases} 1 & \text{if } x > 0, \\ 0 & \text{if } x < 0. \end{cases}$$



■ Properties

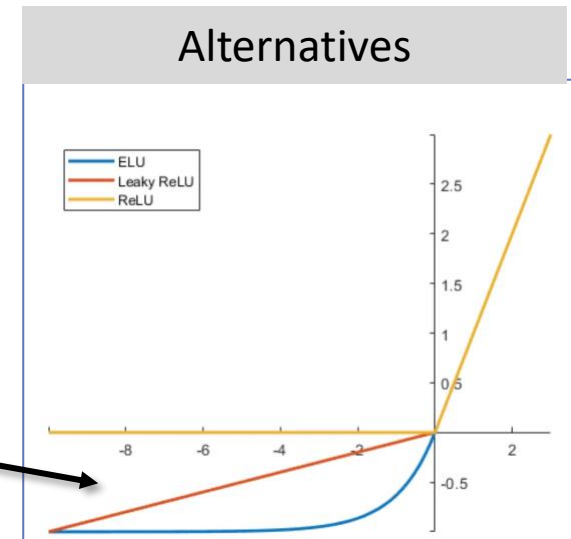
- Linear for positive values
- Non-linear for negative values

■ Advantages

- Avoids vanish gradient: unlike sigmodal does not saturate for positive values
- Simple, fast, efficient

■ Disadvantages

- Non-zero centered output
- Dead neurons: always zero (consistently negative values)



- **2 | Dataset: Cardiac risk / P4_cardiacRiskClassification.csv**
- **2.2 | Evaluation**
 - SE, SP, F1score, ?

```
from sklearn.metrics import confusion_matrix

cm = confusion_matrix(Ttest, Ytest)
TN, FP, FN, TP = cm.ravel()
SE = TP/(TP+FN)
SP = TN/(TN+FP)
F1 = ...
```

- **3 | Cardiac risk / P4_cardiacRiskRegression.csv**
- **3.1 | Implement MLNN – Regression**
 - Structure
 - Number of neurons = ?
 - Activation functions : Hidden layer / Output layer (++)**sigmoidal/linear**)
 - Training / test data
 - Learning algorithm (**adam**)

```
from sklearn.neural_network import MLPRegressor

mlp = MLPRegressor(hidden_layer_sizes=(13,6), activation='relu',
                    solver='adam', max_iter=500)

mlp.fit(Xtrain,Ttrain)
```

- **3 | Dataset: Cardiac risk / P4_cardiacRiskRegression.csv**
- **3.2 | Evaluation**
 - Mean squared Error

```
from sklearn.metrics import mean_squared_error

Etrain = mean_squared_error(Ttrain, Ytrain)
Etest  = mean_squared_error(Ttest, Ytest)
```

- **3 | Cardiac risk / P4_cardiacRiskRegression.csv**
- **Output – activation function**

```
# Output for regression

#if not is_classifier
    activation_ = 'identity'    ?
    activation_ = 'linear'

# Output for multi class
    activation_ = 'softmax'

# Output for binary class
    activation_ = 'logistic'    ?
    activation_ = 'sigmoid'
```

Contents

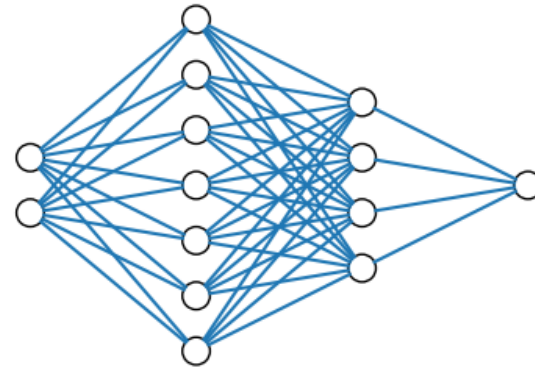
- 1 | Objectives
- 2 | Dataset
- 3 | Tasks
- 4 | Conclusions

- Multi Layer Neural Networks
 - Implement by hand the BP algorithm
 - Function approximation
 - Implement a MLNN
 - Classification
 - Regression problem
 - Evaluation
 - SE, SP - classification
 - Error - regression

■ Improvements

■ Function approximation

- 2 hidden layers ?



■ MLNN - Cardiac Risk

- Activation functions : Hidden layer / Output layer ??
- Learning algorithm ?

■ Other ideas ?

- Use of keras / tensorflow (more general tool)

■ Scikit / Keras

KERAS=0

```
from sklearn.neural_network import MLPClassifier
from sklearn.neural_network import MLPRegressor
```

KERAS=1

```
from keras.models import Sequential
from keras.layers import Dense
```

```
if KERAS==0:
    if regression:
        model = MLPRegressor( hidden_layer_sizes=(12,8),
                               activation='relu',
                               solver='adam', max_iter=3500)
    else:
        model = MLPClassifier(hidden_layer_sizes=(11,4),
                              activation='tanh',
                              solver='adam', max_iter=4500)

    model.fit(Xtrain,Ttrain)
```

■ Scikit / Keras

```
if KERAS==1:
    model = Sequential()
    if regression:
        model.add(Dense(32,activation='sigmoid'))
        model.add(Dense(18,activation='sigmoid'))
        model.add(Dense( 1,activation='linear'))

        model.compile(    loss='mse', optimizer='adam',
                        metrics=['mse'])
    else:
        model.add(Dense(33,activation='sigmoid')    )
        model.add(Dense(12,activation='sigmoid')    )
        model.add(Dense( 4,activation='sigmoid')    )
        model.add(Dense( 1,activation='sigmoid'))

        model.compile(    loss='binary_crossentropy', optimizer='adam',
                        metrics=['accuracy'])

model.fit(Xtrain,Ttrain, batch_size=40, epochs=125, verbose=1)
```

■ Scikit / Keras

```
history=model.fit(Xtrain,Ttrain, validation_split=0.3,
                  batch_size=40, epochs=125,verbose=1)

if KERAS==1:
    #df = pd.DataFrame.from_dict(history.history)

    print(history.history.keys())
    plt.plot(history.history['loss'])
    plt.plot(history.history['val_loss'])

    plt.title(' Training error')
    plt.ylabel('Error')
    plt.xlabel('epoch')
    plt.legend(['train', 'test'], loc='upper left')
    plt.show()
```