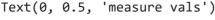
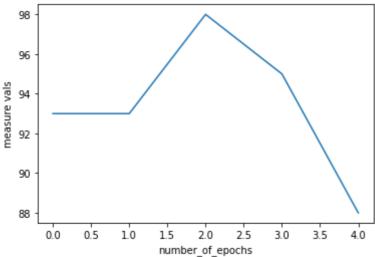
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# Backprop on the Seeds Dataset
from random import seed
from math import exp
import matplotlib.pyplot as plt
from random import randrange
from random import random
import numpy as np
from csv import reader
# how to Load a CSV file with a sample dataset(seeds Data Set)
def load csv func(filename):
 dataset = list()
 with open(filename, 'r') as file:
   csv_reader = reader(file)
   for i in csv_reader:
      if not i:
        continue
      dataset.append(i)
  return dataset
# next we are Converting the string column to float value
def convert_string_column_to_float(dataset, column):
 for i in dataset:
    i[column] = float(i[column].strip())
#then we Convert string column to integer value
def convert_string_column_to_integer(dataset, k):
 class_values = [i[k] for i in dataset]
 unique = set(class_values)
 vp = dict()
 for i, value in enumerate(unique):
   vp[value] = i
 for j in dataset:
    j[k] = vp[j[k]]
  return vp
#function to identify the maximum and minimum values within each column
def dataset_minmax_func(dataset):
 minmax = list()
  stats = [[min(column), max(column)] for column in zip(*dataset)]
  return stats
# changing the dataset range within 0-1
def normalizing dataset func(dataset, minmax):
 for j in dataset:
   for i in range(len(j)-1):
      j[i] = (j[i] - minmax[i][0]) / (minmax[i][1] - minmax[i][0])
# using k fold function to split our dataset
def func_for_cross_validation_split(dataset, number_of_folds):
  value = list()
  dataset conv = list(dataset)
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fold size = int(len(dataset) / number of folds)
  for i in range(number_of_folds):
   fold = list()
   while len(fold) < fold size:
      index = randrange(len(dataset_copy))
      fold.append(dataset_copy.pop(index))
   value.append(fold)
  return value
# finding the accuracy %
def accuracy_metric(actual, predicted):
  correct = 0
 for i in range(len(actual)):
    if actual[i] == predicted[i]:
      correct += 1
  return correct / float(len(actual)) * 100.0
# function for evaluating the algorithm using a cross validation split
def evaluate_algorithm(dataset, algorithm, n_folds, *args):
 folds = func_for_cross_validation_split(dataset, n_folds)
 vals = list()
 for i in folds:
   train_set = list(folds)
   train set.remove(i)
   train_set = sum(train_set, [])
   test_set = list()
   for row in i:
      row copy = list(row)
     test_set.append(row_copy)
      row_copy[-1] = None
   predicted = algorithm(train_set, test_set, *args)
    actual = [row[-1] for row in i]
   accuracy = accuracy_metric(actual, predicted)
   vals.append(accuracy)
  return vals
# function for finding neuron activation within any given input
def activate(weights, inputs):
  activation_val = weights[-1]
 for i in range(len(weights)-1):
    activation_val += weights[i] * inputs[i]
  return activation_val
# function for neuron activation transfer
def transfer(activation):
  return 1.0 / (1.0 + exp(-activation))
# function to forward propagate the given input into network output
def forward_propagate_func(network, row):
  input val = row
 for i in network:
   new inputs = []
   for j in i:
      activation = activate(j['weights'], input_val)
      i['output'] = transfer(activation)
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new inputs.append(j['output'])
    input_val = new_inputs
  return input_val
# finding the derivative of an neuron output
def transfer_derivative_func(output):
  return output * (1.0 - output)
# Backpropagate error function
def backward_propagate_error_func(network, expected):
 for i in reversed(range(len(network))):
    each_layer_val = network[i]
   error_val = list()
   if i != len(network)-1:
      for j in range(len(each_layer_val)):
        error = 0.0
        for k in network[i + 1]:
          error += (k['weights'][j] * k['delta'])
        error_val.append(error)
      for j in range(len(each_layer_val)):
        k = each_layer_val[j]
        error_val.append(k['output'] - expected[j])
   for j in range(len(each_layer_val)):
      k = each_layer_val[j]
      k['delta'] = error_val[j] * transfer_derivative_func(k['output'])
# function for updating network weights with error
def update_weights_func(network, row, l_rate):
 for i in range(len(network)):
    inputs = row[:-1]
   if i != 0:
      inputs = [neuron['output'] for neuron in network[i - 1]]
   for neuron in network[i]:
     for j in range(len(inputs)):
        neuron['weights'][j] -= l_rate * neuron['delta'] * inputs[j]
      neuron['weights'][-1] -= 1 rate * neuron['delta']
# next we train a network for a certain number of epochs
def train network(network, train, 1 rate, n epoch, n outputs):
 for epoch in range(n epoch):
    for row in train:
      outputs = forward_propagate_func(network, row)
      expected = [0 for i in range(n_outputs)]
      expected[row[-1]] = 1
      backward_propagate_error_func(network, expected)
      update_weights_func(network, row, l_rate)
def initialize_network(n_inputs, n_hidden, n_outputs):
  network = list()
  hidden_layer = [{'weights':[random() for i in range(n_inputs + 1)]} for i in range(n_hic
  network.append(hidden_layer)
  output_layer = [{'weights':[random() for i in range(n_hidden + 1)]} for i in range(n_out
  network.append(output layer)
  return network
```

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# predicting with a network
def predict(network, row):
  outputs = forward propagate func(network, row)
  return outputs.index(max(outputs))
# function for backpropagation Algorithm With Stochastic Gradient Descent
def back_propagation(train, test, l_rate, n_epoch, n_hidden):
  n inputs = len(train[0]) - 1
  n_outputs = len(set([row[-1] for row in train]))
  network = initialize_network(n_inputs, n_hidden, n_outputs)
  train_network(network, train, l_rate, n_epoch, n_outputs)
  predictions = list()
  for row in test:
    prediction = predict(network, row)
    predictions.append(prediction)
  return(predictions)
seed(1)
filename = '/content/drive/MyDrive/wheat-seeds.csv'
dataset = load csv func(filename)
for i in range(len(dataset[0])-1):
  convert_string_column_to_float(dataset, i)
convert_string_column_to_integer(dataset, len(dataset[0])-1)
minmax = dataset_minmax_func(dataset)
normalizing_dataset_func(dataset, minmax)
n folds = 5
1_{\text{rate}} = 0.3
n_{epoch} = 500
n_hidden = 5 #this value can be changed to represent both deep and shalow models and plot
scores = evaluate_algorithm(dataset, back_propagation, n_folds, l_rate, n_epoch, n_hidden)
measure=list(np.round(scores,0))
measure
epochs=[0,1,2,3,4]
plt.plot(epochs, measure)
plt.xlabel('number_of_epochs')
plt.ylabel('measure vals')
```





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