

iris

April 30, 2022

```
[1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from sklearn import datasets
from sklearn.linear_model import LogisticRegression as LR
from sklearn.model_selection import train_test_split
import matplotlib.pyplot as plt
```

```
[2]: iris = datasets.load_iris()
data = iris.data
labels = iris.target
num_samples,num_features = data.shape[0],data.shape[1]
print(f'num_samples,num_features = {(num_samples,num_features)}')
indices = np.arange(num_samples)
X_train, X_test, y_train, y_test,train_indices,test_indices =
    ↪train_test_split(data, labels,indices, test_size=0.6, random_state=42)
train_indices = np.arange(1,num_samples,4)
test_indices = np.arange(0,num_samples,4)
print(f'train_indices ={train_indices.shape},test_indices ={test_indices.
    ↪shape}')
```

```
num_samples,num_features = (150, 4)
train_indices =(38,),test_indices =(38,)
```

```
[3]: data
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[3]: array([[5.1, 3.5, 1.4, 0.2],
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[6.5, 3. , 5.2, 2. ],
[6.2, 3.4, 5.4, 2.3],
[5.9, 3. , 5.1, 1.8]])

```

```
[4]: labels
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```
[4]: array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
           0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
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           2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2])
```

```
[5]: train_indices
```

```
[5]: array([ 1,  5,  9, 13, 17, 21, 25, 29, 33, 37, 41, 45, 49,
           53, 57, 61, 65, 69, 73, 77, 81, 85, 89, 93, 97, 101,
           105, 109, 113, 117, 121, 125, 129, 133, 137, 141, 145, 149])
```

```
[6]: test_indices
```

```
[6]: array([ 0,  4,  8, 12, 16, 20, 24, 28, 32, 36, 40, 44, 48,
           52, 56, 60, 64, 68, 72, 76, 80, 84, 88, 92, 96, 100,
           104, 108, 112, 116, 120, 124, 128, 132, 136, 140, 144, 148])
```

```
[7]: from numpy.random import randint
      from numpy.random import rand

      def select_features(elem, features):
          selected_elem = np.where(elem==1)[0]
          selected_features = features[:,selected_elem]
          return selected_features

      def classification_accuracy(labels, preds):
          correct = np.where(labels == preds)[0]
          accuracy = correct.shape[0]/labels.shape[0]
          return accuracy

      def objective(pop, data, labels, train_ind, test_ind):
          accuracies = np.zeros(pop.shape[0])
          idx= 0
          for elem in pop:
              selected_features = select_features(elem, data)
              train_data = selected_features[train_ind,:]
              test_data = selected_features[test_ind,:]
              if train_data.shape[0]==0 or train_data.shape[1]==0 or test_data.
→shape[0]==0 or test_data.shape[1]==0:
                  idx=idx+1
                  continue
              train_labels = labels[train_indices]
              test_labels = labels[test_indices]
```

```

        LR_classifier = LR(random_state=0)
        LR_classifier.fit(X=train_data, y=train_labels)
        predictions = LR_classifier.predict(test_data)
        accuracies[idx] = classification_accuracy(test_labels, predictions)
        idx = idx + 1
    return accuracies

def parent_selection(pop,n_pop,scores,k=3):
    selected = []
    for _ in range(n_pop):
        idx = randint(len(pop))
        for ix in randint(0, len(pop),k-1):
            # check if better (e.g. perform a tournament)
            if scores[ix] < scores[idx]:
                idx = ix
        selected.append(pop[idx])
    return selected

def crossover(p1,p2,r_cross):
    c1 = p1.copy()
    c2 = p2.copy()
    if rand() < r_cross:
        pt = randint(1, len(p1)-2)
        c1 = list(p1[:pt])+list(p2[pt:])
        c2 = list(p2[:pt])+list(p1[pt:])
    return [np.array(c1), np.array(c2)]

def mutation(bitstring, r_mut):
    for i in range(len(bitstring)):
        # check for a mutation
        if rand() < r_mut:
            # flip the bit
            bitstring[i] = 1 - bitstring[i]
    return bitstring

def get_children(selected_parents,n_pop,r_cross,r_mut):
    children = []
    for i in range(0, n_pop, 2):
        p1, p2 = selected_parents[i], selected_parents[i+1]
        for c in crossover(p1, p2, r_cross):
            mutation(c, r_mut)
            children.append(c)
    return np.array(children)

def genetic_algorithm(epochs,data,labels,train_indices,test_indices):
    pop_size = 10
    k = 4

```

```

r_cross = 0.9
r_mut = 1/pop_size
pop_shape = (pop_size, num_features)
#initial population
new_population = np.random.randint(low=0, high=2, size=pop_shape)
print(f"new_population: {new_population} ")

best_outputs = []
num_generations = epochs
for gen in range(num_generations):
    #measure fitness of each member in population
    scores = objective(new_population, data, labels, train_indices,
↳test_indices)

    #print current best in population
    best_outputs.append(np.max(scores))
    print(f"Gen: {gen} => Best result : {best_outputs[-1]}")

    #Select parent in current population to generate children for next
↳generation
    selected = parent_selection(new_population, pop_size, scores)

    #Get children of parents
    children = get_children(selected, pop_size, r_cross, r_mut)

    #replace old population
    new_population = children

#    best_outputs.append(np.max(scores))
#    print(f"Gen: {gen} => Best result : {best_outputs[-1]}")

# Getting the best solution after iterating finishing all generations.
# At first, the fitness is calculated for each solution in the final
↳generation.
    scores = objective(new_population, data, labels, train_indices,
↳test_indices)

    # Then return the index of that solution corresponding to the best fitness.
    best_match_idx = np.where(scores == np.max(scores))[0]
    best_match_idx = best_match_idx[0]
    print(f'np.max(scores) ={np.max(scores)}')
    best_solution = new_population[best_match_idx, :]
    best_solution_indices = np.where(best_solution == 1)[0]
    best_solution_num_elements = best_solution_indices.shape[0]
    best_solution_fitness = scores[best_match_idx]

    print("best_match_idx : ", best_match_idx)
    print("best_solution : ", best_solution)

```

```

print("Selected indices : ", best_solution_indices)
print("Number of selected elements : ", best_solution_num_elements)
print("Best solution fitness : ", best_solution_fitness)

plt.plot(best_outputs)
plt.xlabel("Iteration")
plt.ylabel("Fitness")
plt.show()

genetic_algorithm(100,data,labels,train_indices,test_indices)

```

```

new_population: [[0 1 0 0]
 [1 0 1 0]
 [1 1 1 0]
 [1 0 0 1]
 [1 1 0 0]
 [1 1 0 0]
 [1 1 1 1]
 [1 1 1 0]
 [1 1 1 1]
 [0 1 0 1]]
Gen: 0 => Best result : 0.9736842105263158
Gen: 1 => Best result : 0.9736842105263158
Gen: 2 => Best result : 0.9736842105263158
Gen: 3 => Best result : 0.9210526315789473
Gen: 4 => Best result : 0.9210526315789473
Gen: 5 => Best result : 0.9210526315789473
Gen: 6 => Best result : 0.9736842105263158
Gen: 7 => Best result : 0.9210526315789473
Gen: 8 => Best result : 0.9736842105263158
Gen: 9 => Best result : 0.9736842105263158
Gen: 10 => Best result : 0.9736842105263158
Gen: 11 => Best result : 0.9736842105263158
Gen: 12 => Best result : 0.8947368421052632
Gen: 13 => Best result : 0.9736842105263158
Gen: 14 => Best result : 0.9736842105263158
Gen: 15 => Best result : 0.9736842105263158
Gen: 16 => Best result : 0.5526315789473685
Gen: 17 => Best result : 0.7894736842105263
Gen: 18 => Best result : 0.9736842105263158
Gen: 19 => Best result : 0.9736842105263158
Gen: 20 => Best result : 0.9736842105263158
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Gen: 23 => Best result : 0.9210526315789473
Gen: 24 => Best result : 0.9473684210526315
Gen: 25 => Best result : 0.9736842105263158
Gen: 26 => Best result : 0.7894736842105263

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Gen: 27 => Best result : 0.9210526315789473
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Gen: 29 => Best result : 0.9736842105263158
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Gen: 95 => Best result : 0.9736842105263158
Gen: 96 => Best result : 0.8947368421052632
Gen: 97 => Best result : 0.9736842105263158
Gen: 98 => Best result : 0.9736842105263158
Gen: 99 => Best result : 0.9736842105263158
np.max(scores) =0.9210526315789473
best_match_idx : 9
best_solution : [0 0 1 0]
Selected indices : [2]
Number of selected elements : 1
Best solution fitness : 0.9210526315789473
```

