

In a logistic regression model, we use a weighted sum of these two predictors to calculate a joint score S :

$$S = \beta_0 + \beta_1 x_1 + \beta_2 x_2. \quad (1)$$

The logistic regression model gives us appropriate values for the parameters $\beta_0, \beta_1, \beta_2$ using two sets of example genes:

- OP: Adjacent genes, on the same strand of DNA, known to belong to the same operon;
- NOP: Adjacent genes, on the same strand of DNA, known to belong to different operons.

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```

>>> xs = [[-53, -200.78],
           [117, -267.14],
           [57, -163.47],
           [16, -190.30],
           [11, -220.94],
           [85, -193.94],
           [16, -182.71],
           [15, -180.41],
           [-26, -181.73],
           [58, -259.87],
           [126, -414.53],
           [191, -249.57],
           [113, -265.28],
           [145, -312.99],
           [154, -213.83],
           [147, -380.85],
           [93, -291.13]]

>>> ys = [1,
           1,
           1,
           1,
           1,
           1,
           1,
           1,
           1,
           0,
           0,
           0,
           0,
           0,
           0,
           0,
           0]

>>> model = LogisticRegression().train(xs, ys)

```

Here, `xs` and `ys` are the training data: `xs` contains the predictor variables for each gene pair, and `ys` specifies if the gene pair belongs to the same operon (1, class OP) or different operons (0, class NOP). The resulting logistic regression model is stored in `model`, which contains the weights β_0 , β_1 , and β_2 :

```

>>> model.beta
[8.9830290157144681, -0.035968960444850887, 0.02181395662983519]

```

Note that β_1 is negative, as gene pairs with a shorter intergene distance have a higher probability of belonging to the same operon (class OP). On the other hand, β_2 is positive, as gene pairs belonging to the same operon typically have a higher similarity score of their gene expression profiles. The parameter β_0 is

```
    print "Iteration: ", iteration, "Log-likelihood function: ", log_likelihood
>>>
>>> model = LogisticRegression().train(xs, ys, update_fn=show_progress)
Iteration: 0 Log-likelihood function: -11.7835020695
Iteration: 1 Log-likelihood function: -7.15886767672
Iteration: 2 Log-likelihood function: -5.76877209868
Iteration: 3 Log-likelihood function: -5.11362294338
Iteration: 4 Log-likelihood function: -4.74870642433
Iteration: 5 Log-likelihood function: -4.50026077146
Iteration: 6 Log-likelihood function: -4.31127773737
Iteration: 7 Log-likelihood function: -4.16015043396
Iteration: 8 Log-likelihood function: -4.03561719785
Iteration: 9 Log-likelihood function: -3.93073282192
Iteration: 10 Log-likelihood function: -3.84087660929
Iteration: 11 Log-likelihood function: -3.76282560605
Iteration: 12 Log-likelihood function: -3.69425027154
Iteration: 13 Log-likelihood function: -3.6334178602
Iteration: 14 Log-likelihood function: -3.57900855837
Iteration: 15 Log-likelihood function: -3.52999671386
Iteration: 16 Log-likelihood function: -3.48557145163
Iteration: 17 Log-likelihood function: -3.44508206139
Iteration: 18 Log-likelihood function: -3.40799948447
Iteration: 19 Log-likelihood function: -3.3738885624
Iteration: 20 Log-likelihood function: -3.3423876581
Iteration: 21 Log-likelihood function: -3.31319343769
Iteration: 22 Log-likelihood function: -3.2860493346
Iteration: 23 Log-likelihood function: -3.2607366863
Iteration: 24 Log-likelihood function: -3.230e84091
Iteration: 25 Log-likelihood function: -3.21488073614
Iteration: 26 Log-likelihood function: -3.19403459259
Iteration: 27 Log-likelihood function: -3.17440646052
Iteration: 28 Log-likelihood function: -3.15588842703
Iteration: 29 Log-likelihood function: -3.13838533947
Iteration: 30 Log-likelihood function: -3.12181293595
Iteration: 31 Log-likelihood function: -3.10609629966
Iteration: 32 Log-likelihood function: -3.09116857282
Iteration: 33 Log-likelihood function: -3.07696988017
Iteration: 34 Log-likelihood function: -3.06344642288
Iteration: 35 Log-likelihood function: -3.05054971191
```

3 Using the logistic regression model for classification

Classification is performed by calling the `classify`

