ML Homework 4

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1

When w_0 increases, then the probability of predicting class 1 becomes higher, and vice versa.

$$h(x) = \frac{1}{1 + e^{-(w_0 + \dots)}}$$

This is because as w_0 gets bigger, the result of exponentiation gets smaller. As the denominator gets smaller, the quotient gets larger. h(x) is used as the probability we will predict a certain class. The higher h(x) is, the more likely we will predict class 1 (as opposed to class 0).

 $\mathbf{2}$

The logistic function is

$$\frac{1}{1 + e^{-\mathbf{w}^T \mathbf{x}}}$$

If we were to double \mathbf{w} , the result wouldn't actually change, as long as the threshold is 0.5.

The geometric interpretation is because $\mathbf{w}^T \mathbf{x}$ represents a (hyper)plane. Multiplying the equation of a plane doesn't affect any points that are on the plane (threshold = 0.5 \Longrightarrow on hyperplane), but it makes the points that are off it seem further away, increasing the certainty with which we pick a class. For example, if $\mathbf{w}^T \mathbf{x} = 0$, then there is no change $(0 \cdot 2 = 0)$. But if it's positive, it becomes a larger positive number, and vice versa.

(a)
$$\begin{bmatrix} 3 & 1 \\ 1 & 3 \end{bmatrix}$$

(b):

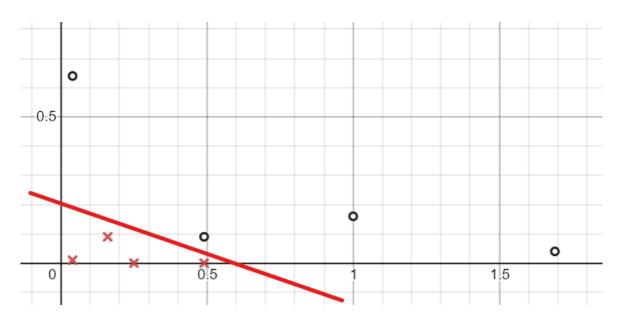


Figure 1: o: class 0, x: class 1

(c) TPR =
$$\frac{1}{3+1} = 1/4$$

(d)
$$FPR = \frac{3}{3+1} = 3/4$$

(e) accuracy =
$$\frac{3+3}{3+1+3+1} = 3/4$$

(f) recall =
$$\frac{3}{3+1} = 3/4$$

(g) precision =
$$\frac{3}{3+1} = 3/4$$

(h)

$$-[0 \cdot \ln{(0.389)} + (1 - 0) \ln{(1 - 0.389)}] - \dots - [1 \cdot \ln{(0.638)} + (1 - 1) \ln{(1 - 0.638)}] = 4.25205106\dots$$

(i)
$$-\left[0 \cdot \ln\left(\frac{1}{1 + \exp\left(-\mathbf{w'}^T\mathbf{X^{(1)}}\right)}\right)\right] + \dots - \left[0 \cdot \ln\left(\frac{1}{1 + \exp\left(-\mathbf{w'}^T\mathbf{X^{(8)}}\right)}\right)\right] = 3.0158793\dots$$

Since \mathbf{w}' has a lower cross-entropy, it is a better fit for this data.

(j) After one interation,

$$\mathbf{w} = \mathbf{w} + \frac{\alpha}{N} \mathbf{x}^{T} (y - \sigma(\mathbf{x}\mathbf{w}))$$

$$\begin{bmatrix} 0.66 \\ -2.24 \\ -0.18 \end{bmatrix}$$

$$\begin{bmatrix} 0 \\ 0 \end{bmatrix}$$

$$+\frac{0.01}{N}\begin{bmatrix}1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 \\ 0.49 & 1.69 & 0.04 & 1 & 0.16 & 0.25 & 0.49 & 0.04 \\ 0.09 & 0.04 & 0.64 & 0.16 & 0.09 & 0 & 0 & 0.01\end{bmatrix}\begin{bmatrix}0 & 1 & 0.49 & 0.09 \\ 1 & 1.69 & 0.04 \\ 1 & 0.04 & 0.64 \\ 1 & 0.16 & 0.09 \\ 1 & 0.25 & 0 \\ 1 & 0.04 & 0.01\end{bmatrix}\begin{bmatrix}0.66 \\ -2.24 \\ -0.18\end{bmatrix}$$

$$= \begin{bmatrix} 0.661 \dots \\ -2.239 \dots \\ -0.181 \dots \end{bmatrix}$$

- (k) These data points would contribute a medium amount to the new w.
- (1) These data points would contribute a low amount to the new \mathbf{w} .
- (m) These data points would contribute a high amount to the new w.

(n)
$$-\left[0 \cdot \ln\left(\frac{1}{1 + \exp\left(-\mathbf{w_g}^T \mathbf{X^{(1)}}\right)}\right)\right] + \dots - \left[0 \cdot \ln\left(\frac{1}{1 + \exp\left(-\mathbf{w_g}^T \mathbf{X^{(8)}}\right)}\right)\right] = 4.2519\dots$$

This is a very minor decrease from 4.2520. This makes sense because gradient ascent is supposed to make the model fit better and have less error.

lasso

$$-\lambda (|w_1| + \dots + |w_d|) + \sum_{i=1}^{N} y^{(i)} \ln (h(x)) + (1 - y^{(i)}) \ln (1 - h(x))$$

ridge

$$-\lambda \left(w_1^2 + \dots + w_d^2\right) + \sum_{i=1}^N y^{(i)} \ln \left(h(x)\right) + (1 - y^{(i)}) \ln \left(1 - h(x)\right)$$

• gradient

$$\mathbf{x}^{T}(y - \sigma(\mathbf{x}\mathbf{w})) - 2\lambda I'\mathbf{w}$$

• Optimal $\lambda = 0.6$. It did help, or else the best lambda would be 0. The training error would get worse (no more overfitting), but the test/validation data would get better.

Figure 2: Adding Ridge regression

```
from sklearn.model selection import KFold
def cross(L):
   nfold = 5
    kf = KFold(n_splits=nfold, shuffle=True)
   X unscaled = cancer.data
    X_unscaled = np.hstack((np.ones((X_unscaled.shape[0], 1)), X_unscaled))
    y orig = cancer.target
    likelihoods = []
    for train, test in kf.split(X_unscaled):
       X_tr = X_unscaled[train,:]
       y_tr = y_orig[train]
       X_ts = X_unscaled[test, :]
        y_ts = y_orig[test]
        scaler KFold = preprocessing.StandardScaler()
        X tr = scaler KFold.fit transform(X tr)
        X ts = scaler KFold.transform(X ts)
        y_2d_ts = y_ts.reshape((y_ts.shape[0], 1))
        y_2d_tr = y_tr.reshape((y_tr.shape[0], 1))
        w kfold, w likelihood = Gradient Ascent(X tr, y 2d tr, 0.001, 100000, L)
        likelihoods.append(likelihood(X_ts, y_2d_ts, w_kfold, X_ts.shape[0]))
    return max(likelihoods)
lambdas = []
div = 10
for lambda in range(0, 100):
    lambdas.append(cross(lambda /div))
    print(lambda_/div, lambdas[lambda_])
print(f"index: {lambdas.index(max(lambdas))}, value:{max(lambdas)}")
```

Figure 3: 5 fold

Basically the code iterates over a range of lambdas and does cross validation on each, and then we see which one has the best errors.

Interestingly, if the iterations for gradient ascent were too low, I would actually get $\lambda = 0$ as the best error. I got $\lambda = 0.6$ from increasing the iterations and letting the code run for like 30 minutes.

- (a) Predictors: average pitch; response: gender
- (b) Predictors: matrix of where the strokes were; response: letter or number written

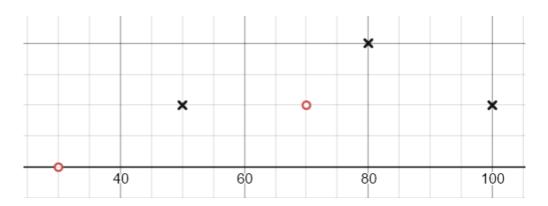


Figure 4: o: no donation, x: donation

(a)

(b) I will choose the horizontal line $x_2 = 0.5 \implies 0.5 + 0 \cdot x_1 + w_2$

$$\mathbf{w}^T = [-0.5, 0, 1]$$

- (c) The least likely would be the one that is incorrect. In this case, it's same 3, or the one that earns 70k, follows 1 site, and did not donate. No calculator was needed for this!
- (d) They would not change the values of \hat{y} , but they would change the likelihoods. If the prediction was correct, then the likelihood would increase. If the predictions were incorrect, then the likelihood would decrease.