

d5__breast__cancer__detector

October 21, 2022

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1 Breast Cancer Wisconsin (Diagnostic) Data Set

[https://archive.ics.uci.edu/ml/datasets/breast+cancer+wisconsin+\(diagnostic\)](https://archive.ics.uci.edu/ml/datasets/breast+cancer+wisconsin+(diagnostic))

1.1 Attribute Information:

1. ID number
2. Diagnosis (M = malignant:-1, B = benign:1) M: B:
3. 3-32

Ten real-valued features are computed for each cell nucleus:

- radius (mean of distances from center to points on the perimeter)
- texture (standard deviation of gray-scale values)
- perimeter
- area
- smoothness (local variation in radius lengths)
- compactness ($\text{perimeter}^2 / \text{area} - 1.0$)
- concavity (severity of concave portions of the contour)
- concave points (number of concave portions of the contour)
- symmetry
- fractal dimension (“coastline approximation” - 1)

mean, stderr, worst

<http://people.idsia.ch/~juergen/deeplearningwinsMICCAIgrandchallenge.html>

1.2

(training) \mathbf{A} \mathbf{b} (-1) (1)
 \mathbf{y} $C(\mathbf{y})$

1.3

() $\mathbb{R}^D \rightarrow \mathbb{R}$ $h(\cdot)$

$$C(\mathbf{y}) = \begin{cases} +1 & \text{when } h(\mathbf{y}) \geq 0 \\ -1 & \text{when } h(\mathbf{y}) < 0 \end{cases}$$

$h : \mathbb{R}^D \rightarrow \mathbb{R}$ D \mathbf{w}

$$h(\mathbf{y}) = \mathbf{w} \cdot \mathbf{y}$$

D \mathbf{w} \mathbf{w} h \mathbf{w}
 -1,1 \mathbf{w}

2

$$L(\mathbf{w}) = \sum_{i=1}^n (A_i \cdot \mathbf{w} - b_i)^2$$

\mathbf{w} j

$$\begin{aligned} \frac{\partial L}{\partial w_j} &= \sum_{i=1}^n \frac{\partial}{\partial w_j} (A_i \cdot \mathbf{w} - b_i)^2 \\ &= \sum_{i=1}^n 2(A_i \cdot \mathbf{w} - b_i) A_{ij} \end{aligned}$$

A_{ij} A_i j $\frac{\partial L}{\partial w_j}$ \mathbf{w}_j (slope) $L(\mathbf{w})$ (local minimum)
(steepest descent method)

2.1 print_w

\mathbf{w} j

```
[ ]: def print_w(w):  
    params = ["radius", "texture", "perimeter", "area",  
              "smoothness", "compactness", "concavity", "concave points",  
              "symmetry", "fractal dimension"]  
    print("      (params)      :      ", end="")  
    print("      (mean)      (stderr)      (worst)")  
    for i, param in enumerate(params):
```

```

print("%18s:" %param, end="")
for j in range(3):
    print("%13.9f" % w[i*3+j], end="")
print()

```

2.2

```

[ ]: import numpy as np
tmp = np.fromfile('./train_A.data', np.float64, -1, " ")
A = tmp.reshape(300,30)
tmp = np.fromfile('./train_b.data', np.float64, -1, " ")
b = tmp.reshape(300,1)
w = np.zeros(30).reshape(30,1)
for i in range(30):
    w[i] = 0

```

```
[ ]: A[0]
```

```

[ ]: array([1.799e+01, 1.038e+01, 1.228e+02, 1.001e+03, 1.184e-01, 2.776e-01,
          3.001e-01, 1.471e-01, 2.419e-01, 7.871e-02, 1.095e+00, 9.053e-01,
          8.589e+00, 1.534e+02, 6.399e-03, 4.904e-02, 5.373e-02, 1.587e-02,
          3.003e-02, 6.193e-03, 2.538e+01, 1.733e+01, 1.846e+02, 2.019e+03,
          1.622e-01, 6.656e-01, 7.119e-01, 2.654e-01, 4.601e-01, 1.189e-01])

```

```
[ ]: b[0]
```

```
[ ]: array([-1.])
```

2.3 w (steepest descent)

```

[ ]: loop, sigma = 300, 3.0*10**(-9)
for i in range(loop):
    dLw = A.dot(w)-b
    w = w - (dLw.transpose().dot(A)).transpose()*sigma

print_w(w)

```

(params)	:	(mean)	(stderr)	(worst)
radius:		0.000426997	0.000741817	0.002548876
texture:		0.001687946	0.000004707	0.000000127
perimeter:		-0.000003968	-0.000002078	0.000008954
area:		0.000003595	0.000002569	0.000070324
smoothness:		0.000001139	-0.000881778	0.000000430
compactness:		0.000000441	0.000000723	0.000000267
concavity:		0.000001200	0.000000191	0.000411499
concave points:		0.000921972	0.002395138	-0.001932789

```

        symmetry:  0.000005930 -0.000003750 -0.000008147
fractal dimension: -0.000002341  0.000011565  0.000003523

(params)      :      (mean)      (stderr)      (worst)
    radius:  0.000426997  0.000741817  0.002548876
    texture:  0.001687946  0.000004707  0.000000127
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        symmetry:  0.000005930 -0.000003750 -0.000008147
fractal dimension: -0.000002341  0.000011565  0.000003523

```

3

```

[ ]: def show_accuracy(mA, vb, vw):
    # M: (-1) B: (1)

    correct,safe_error,critical_error=0,0,0
    predict = mA.dot(vw)
    n = vb.size
    for i in range(n):
        if predict[i]*vb[i]>0:
            correct += 1
        elif (predict[i]<0 and vb[i]>0): #
            safe_error += 1
        elif (predict[i]>0 and vb[i]<0): #
            critical_error += 1
    print("      correct: %4d/%4d" % (correct,n))
    print("    safe error: %4d" % safe_error)
    print("critical error: %4d" % critical_error)

```

```

[ ]: show_accuracy(A, b, w)

```

```

        correct:  274/ 300
        safe error:    5
critical error:   21

```

```

[ ]: tmp = np.fromfile('./validate_A.data', np.float64, -1, " ")
A = tmp.reshape(260,30)
tmp = np.fromfile('./validate_b.data', np.float64, -1, " ")
b = tmp.reshape(260,1)

show_accuracy(A, b, w)

```

```

        correct:  240/ 260

```

```

    safe error:    10
critical error:    10

```

4 QR decomposition

QR A $\|A.w - b\|^2$ w

QR $n \times m$

$$A = QR$$

Q $n \times m$ R $m \times m$

$\|Aw - b\|$ QR

$$Q.R.w = b \Rightarrow R.w = Q^t.b \Rightarrow R^{-1}.R.w = R^{-1}.Q^t.b$$

```

[ ]: import numpy as np

tmp = np.fromfile('./train_A.data', np.float64, -1, " ")
A = tmp.reshape(300,30)
tmp = np.fromfile('./train_b.data', np.float64, -1, " ")
b = tmp.reshape(300,1)

q, r = np.linalg.qr(A)

```

```

[ ]: ww = np.linalg.inv(r).dot(np.transpose(q).dot(b))

```

```

[ ]: q.shape

```

```

[ ]: (300, 30)

```

```

[ ]: print(r[0,0:5])

```

```

[-2.57579883e+02 -3.32324268e+02 -1.68607899e+03 -1.29450676e+04
 -1.65446346e+00]

```

```

[ ]: show_accuracy(A, b, ww)

```

```

    correct: 286/ 300
    safe error:    1
critical error:    13

```

```

[ ]: print_w(ww)

```

```

(params)      :      (mean)      (stderr)      (worst)
    radius:  0.869921844 -0.024313948 -0.062679561
    texture: -0.003274619 -8.790300861  1.747147500
    perimeter: -0.202849407 -6.506451098  5.061760446
    area: 49.167541566 -0.956591421 -0.082052658

```

```
smoothness: -0.007943157 0.004976908-27.841944367
compactness: 3.301527110 4.985959134-16.318886295
concavity: 10.316289081-21.332232171 -0.408605816
concave points: -0.003345722 -0.000677873 0.002510735
symmetry: 4.531369718 0.590110016 -0.719368704
fractal dimension: -2.158965299 -3.803467225-12.298417038
```

```
[ ]: tmp = np.fromfile('./validate_A.data', np.float64, -1, " ")
A = tmp.reshape(260,30)
tmp = np.fromfile('./validate_b.data', np.float64, -1, " ")
b = tmp.reshape(260,1)

show_accuracy(A, b, ww)
```

```
correct: 252/ 260
safe error: 6
critical error: 2
```