

Least squares method

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Breast cancer dataset

- 1) ID number
- 2) Diagnosis (M = malignant, B = benign)
- 3) radius (mean)
- 4) texture (mean)
- 5) perimeter (mean)
- 6) area (mean)
- 7) smoothness (mean)
- 8) compactness (mean)
- 9) concavity (mean)
- 10) concave points (mean)
- 11) symmetry (mean)
- 12) fractal dimension (mean)

Breast cancer dataset

- 13) radius (stderr)
- 14) texture (stderr)
- 15) perimeter (stderr)
- 16) area (stderr)
- 17) smoothness (stderr)
- 18) compactness (stderr)
- 19) concavity (stderr)
- 20) concave points (stderr)
- 21) symmetry (stderr)
- 22) fractal dimension (stderr)

Breast cancer dataset

- 23) radius (worst)
- 24) texture (worst)
- 25) perimeter (worst)
- 26) area (worst)
- 27) smoothness (worst)
- 28) compactness (worst)
- 29) concavity (worst)
- 30) concave points (worst)
- 31) symmetry (worst)
- 32) fractal dimension (worst)

Least squares method

Linear representation

$$A_{\text{lin}} = \begin{bmatrix} f_{1,1} & \dots & f_{1,m} \\ f_{2,1} & \dots & f_{2,m} \\ \vdots & \dots & \vdots \\ f_{n,1} & \dots & f_{n,m} \end{bmatrix} \quad (1)$$

Least squares method

$$Aw \cong b, \quad A \in \mathbb{R}^{n \times m}, \quad b \in \mathbb{R}^{1 \times n} \quad (2)$$

n - number of instances

m - number of features

$$\min_w ||Aw - b|| \quad (3)$$

Least squares method

Linear representation

$$A_{\text{lin}} = \begin{bmatrix} f_{1,1} & f_{1,2} & f_{1,3} & f_{1,4} \\ f_{2,1} & f_{2,2} & f_{2,3} & f_{2,4} \\ \vdots & \vdots & \vdots & \vdots \\ f_{n,1} & f_{n,2} & f_{n,3} & f_{n,4} \end{bmatrix} \quad (4)$$

Quadratic representation

$$A_{\text{quad}} = \begin{bmatrix} f_{1,1}, f_{1,2}, f_{1,3}, f_{1,4}, f_{1,1}^2, f_{1,2}^2, f_{1,3}^2, f_{1,4}^2, f_{1,1}f_{1,2}, f_{1,1}f_{1,3}, f_{1,1}f_{1,4}, f_{1,2}f_{1,3}, f_{1,2}f_{1,4}, f_{1,3}f_{1,4} \\ \vdots \\ f_{n,1}, f_{n,2}, f_{n,3}, f_{n,4}, f_{n,1}^2, f_{n,2}^2, f_{n,3}^2, f_{n,4}^2, f_{n,1}f_{n,2}, f_{n,1}f_{n,3}, f_{n,1}f_{n,4}, f_{n,2}f_{n,3}, f_{n,2}f_{n,4}, f_{n,3}f_{n,4} \end{bmatrix}$$

Normal equation

$$w = \underbrace{(A^T A)^{-1} A^T}_{\substack{\text{pseudoinverse} \\ \text{matrix}}} b \quad (5)$$

$$w = A^\dagger b \quad (6)$$

- $A^T A$ is not guaranteed to be non-singular
- The method is overly sensitive to the condition number of matrix
- QR and SVD are numerically more stable alternatives but computationally slower

$$\text{cond}(A) = \|A\| \cdot \|A^{-1}\| \quad (7)$$

Stability of normal equation:

$$\text{cond}(A^T A) = \text{cond}(A)^2 \quad (8)$$

References

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