

Introduction to Machine Learning

Evaluating Model Performance (Part 2)

Confusion matrix

| predicted→ real↓ | Class_pos | Class_neg |
|---------------------|-----------|-----------|
| Class_pos | TP | FN |
| Class_neg | FP | TN |

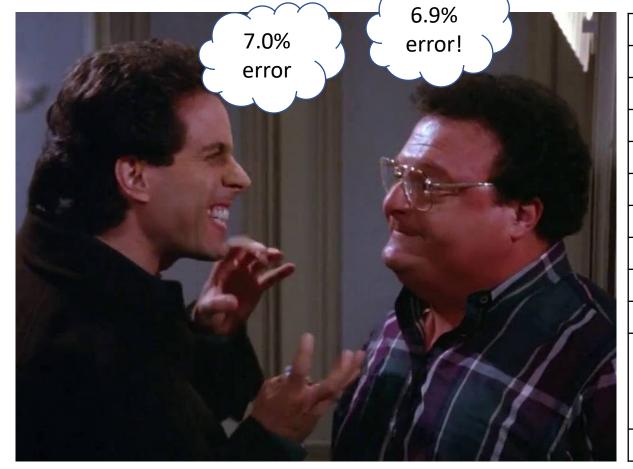
(Tra

$$P = TP + FN$$

$$N = TN+FP$$

 Accuracy is % correct (fraction correct)

| V |
|----------|
| 1 |
| √ |
| X |
| √ |
| X |
| X |
| X |
| √ |
| 1 |
| |
| |
| |
| V |
| |



| x1 | V |
|-------|----------|
| x2 | 1 |
| х3 | V |
| х4 | X |
| x5 | 1 |
| х6 | X |
| x8 | X |
| x9 | ***-*** |
| x10 | 1 |
| x11 | 1 |
| | |
| | |
| | |
| x1000 | √ |

| x1 | √ |
|-------|----------|
| x2 | 7 |
| х3 | V |
| x4 | X |
| x5 | V |
| х6 | X |
| x8 | X |
| x9 | X |
| x10 | 7 |
| x11 | 7 |
| | |
| | |
| | |
| x1000 | V |
| | • |



| √ |
|---|
| X |
| √ |
| √ |
| √ |
| 1 |
| X |
| √ |
| X |
| √ |
| |
| |
| |
| 1 |
| |

Hypothesis: Newman's algorithm is better than mine

• Null hypothesis: Newman's algorithm is not better than mine

Determine whether difference is statistically significant (not just due to random luck)

- Compute p-value
- Probability that observed difference was luck

"There is a 95% chance this difference was not by chance"

- Calculate sample mean (population mean)
- Degrees of freedom = n-1



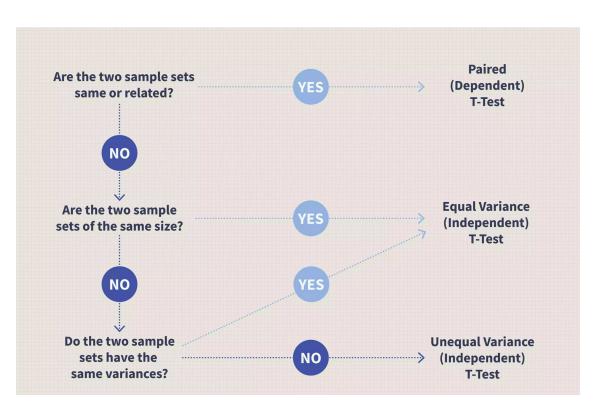
- Error of algorithm A is a₁, .., a_N
- Error of algorithm B is b₁, .., b_N
- Center data points around means μ_a and μ_b
 - Each a_i is now $\hat{a}_i = a \mu_a$

- Error of algorithm A is a₁, .., a_N
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- Center data points around means μ_a and μ_b
 - Each a_i is now $\hat{a}_i = a \mu_a$
 - Each b_i is now $\hat{b}_i = b \mu_b$

$$t = (\mu_a - \mu_b) \sqrt{\frac{N(N-1)}{\sum_n (\hat{a}_n - \hat{b}_n)^2}}$$

| t | significance |
|-------------|--------------|
| ≥ 1.28 | 90.0% |
| ≥ 1.64 | 95.0% |
| ≥ 1.96 | 97.5% |
| ≥ 2.58 | 99.5% |

One of many such tests



Calculating p-value from t statistic

- t distribution
- We care about values away from the mean
 - Mean +/- t
- Look up p value based on t statistic and degrees of freedom (= N-1)
- t table
 - Here for two tailed
 - https://www.medcalc.org/manual/t-distribution.php

Example

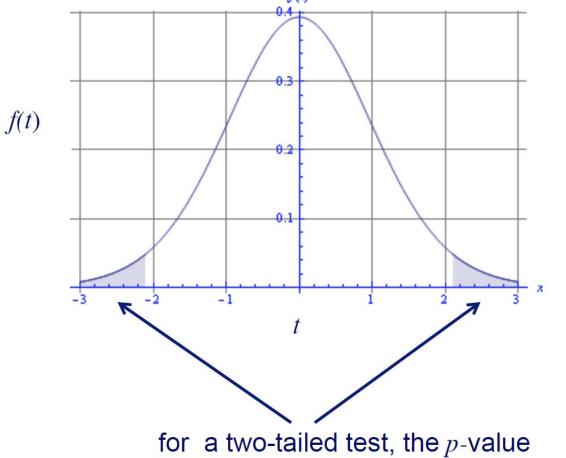
| Data | А | В | |
|------|-------|----|--|
| 1 | 3 | 20 | |
| 2 | 3 | 13 | |
| 3 | 3 | 13 | |
| 4 | 12 | 20 | |
| 5 | 15 | 29 | |
| 6 | 16 32 | 32 | |
| 7 | 17 | 23 | |
| 8 | 19 | 20 | |
| 9 | 23 | 25 | |
| 10 | 24 | 15 | |
| 11 | 32 | 30 | |

$$t = (\mu_a - \mu_b) \sqrt{\frac{N(N-1)}{\sum_n (\hat{a}_n - \hat{b}_n)^2}}$$

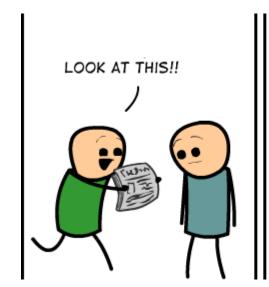
Generate p value

| Data | А | В |
|------|----|----|
| 1 | 3 | 20 |
| 2 | 3 | 13 |
| 3 | 3 | 13 |
| 4 | 12 | 20 |
| 5 | 15 | 29 |
| 6 | 16 | 32 |
| 7 | 17 | 23 |
| 8 | 19 | 20 |
| 9 | 23 | 25 |
| 10 | 24 | 15 |
| 11 | 32 | 30 |

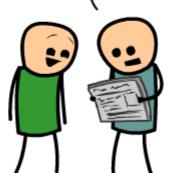
| DF | A = 0.2 | 0.10 | 0.05 | 0.02 | 0.01 | 0.002 | 0.001 |
|----|---------------|-------|--------|--------|--------|---------|---------|
| 00 | $t_a = 1.282$ | 1.645 | 1.960 | 2.326 | 2.576 | 3.091 | 3.291 |
| 1 | 3.078 | 6.314 | 12.706 | 31.821 | 63.656 | 318.289 | 636.578 |
| 2 | 1.886 | 2.920 | 4.303 | 6.965 | 9.925 | 22.328 | 31.600 |
| 3 | 1.638 | 2.353 | 3.182 | 4.541 | 5.841 | 10.214 | 12.924 |
| 4 | 1.533 | 2.132 | 2.776 | 3.747 | 4.604 | 7.173 | 8.610 |
| 5 | 1.476 | 2.015 | 2.571 | 3.365 | 4.032 | 5.894 | 6.869 |
| 6 | 1.440 | 1.943 | 2.447 | 3.143 | 3.707 | 5.208 | 5.959 |
| 7 | 1.415 | 1.895 | 2.365 | 2.998 | 3.499 | 4.785 | 5.408 |
| 8 | 1.397 | 1.860 | 2.306 | 2.896 | 3.355 | 4.501 | 5.041 |
| 9 | 1.383 | 1.833 | 2.262 | 2.821 | 3.250 | 4.297 | 4.781 |
| 10 | 1.372 | 1.812 | 2.228 | 2.764 | 3.169 | 4.144 | 4.587 |
| 11 | 1.363 | 1.796 | 2.201 | 2.718 | 3.106 | 4.025 | 4.437 |
| | | | | | | | |



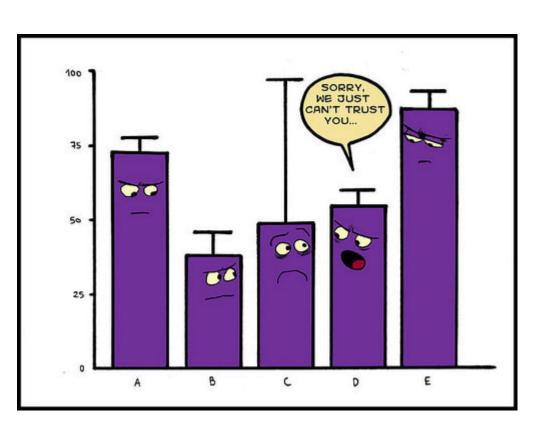
for a two-tailed test, the p-value represents the probability mass in these two regions











Confidence intervals

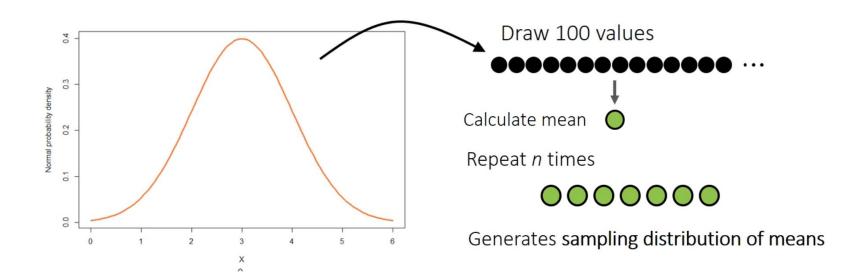
The **median**for the population
lies between

Confidence intervals



Confidence intervals

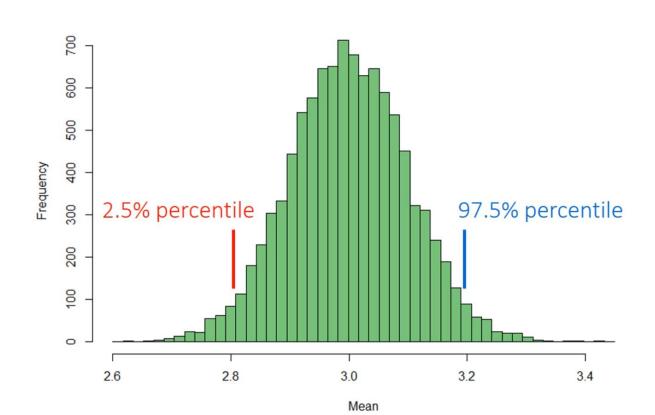
If you took many random samples from the population, 95% of the confidence intervals on those samples would include μ



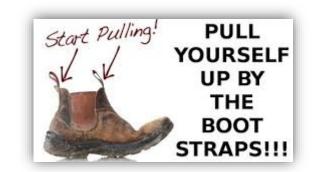
What if I do not have enough data?

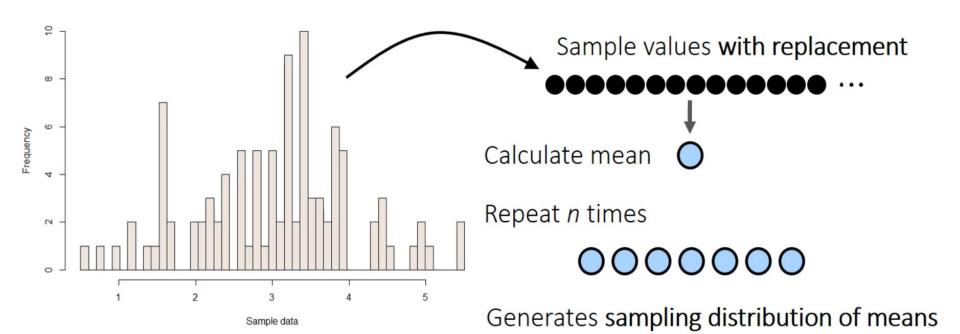


Resampling distribution



Bootstrapping





Algorithm 10 BOOTSTRAPEVALUATE $(y, \hat{y}, NumFolds)$

```
1: Scores ←
2: for k = 1 to NumFolds do
      truth \leftarrow []
                                                           // list of values we want to predict
      pred \leftarrow []
                                                        // list of values we actually predicted
      for n = \tau to N do
         m \leftarrow uniform random value from 1 to N
                                                                         // sample a test point
                                                                             // add on the truth
         truth \leftarrow truth \oplus y_m
         pred \leftarrow pred \oplus \hat{y}_m
                                                                       // add on our prediction
      end for
                                                                                      // evaluate
      scores \leftarrow scores \oplus F\text{-}score(truth, pred)
10:
··· end for
12: return (MEAN(scores), STDDEV(scores))
```

Why is the learning algorithm performing poorly?