### Design & Analysis of Data Science Experiments Model Inter-comparison

Nagiza F. Samatova, <a href="mailto:samatova@csc.ncsu.edu">samatova@csc.ncsu.edu</a>
Professor, Department of Computer Science
North Carolina State University



### Assessing Model Performance TESTING ERROR RATES

#### **Testing Error Rates: Application**

- Classification error
- Squared error in regression
- Log likelihood in unsupervised learning
- Expected reward in reinforcement learning

#### **Book Chapter Sections**

- 19.10.1. Binomial Test for Single Classifier, Single Validation Set
- 19.10.2. Approximate Normal Test for Single Classifier, Single Validation Set
- 19.10.3. t-Test for Single Classifier with K-fold Cross-Validation
- 19.11.1. McNemar's Test to Compare Two Classifiers w/ Single Validation Set
- 19.11.2. K-fold Cross-Validated Paired t-test
- 19.11.3. 5x2 cv Paired t-test

- 19.12 Comparing Multiple Algorithms: Analysis of Variance
- 19.13.1 Nonparametric tests for comparison over multiple data sets: Wilcoxon Signed Rank Test for Two Algorithm Comparison
- 19.13.2. Nonparametric tests for comparison over multiple data sets: Kruskal-Wallis Test for Multiple Algorithm Comparison
- 19.14.1 Multi-criterion Assessment: Multivariate Test for Two Algorithms
- 19.14.2 Multi-criterion Assessment: Multivariate Test for Multiple Algorithms (MANOVA)

#### Single Classifier Assessment BINOMIAL TEST FOR SINGLE CLASSIFIER USING SINGLE VALIDATION SET (19.10.1)

help (binom.test)

http://www.instantr.com/2012/11/06/performing-a-binomial-test/

#### When and how to use?

- The binomial test is used to assess the error rate of a single dichotomous (binary) classification algorithm.
- Assumptions
  - The training is done on the single training set
  - The single validation set (non-overlapping with the training set) is used for the assessment.
- Hypothesis Testing using binom.test():
  - Let's assume that p is the probability that the classifier makes a misclassification error
  - Estimate or test a hypothesis about the probability of a success in a Bernoulli experiment

#### **Binomial Test**

- Let x<sup>t</sup> denote the correctness of the classifier decision:
  - it is a 0/1 Bernoulli random variable
  - it is 1 when the classifier commits the error; 0, otherwise
- Let  $X = \sum_{t=1}^{N} x^t$  denote the binomial random variable for the total number of errors
- The Null Hypothesis (H<sub>0</sub>):
  - The error probability  $p: p \le p_0$  (e.g.,  $p_0 = 0.5$ )
- The Alternative Hypothesis  $(H_1)$ :  $p > p_0$
- If the probability of error is p then

• 
$$P(X=j) = \binom{N}{j} p^j (1-p)^{N-j}$$

• Binomial test reject  $H_0$  if for the significance value of  $\alpha$  (e.g.,  $\alpha = 0.05$ ):

• 
$$P(X \ge e) = \sum_{j=e}^{N} {N \choose j} p_0^j (1 - p_0)^{N-j} < \alpha$$

#### binom.test()

binom.test (x, n, p = 0.5, alternative = c("two.sided", "less", "greater"), conf.level = 0.95)

- x The parameter can be a single value indicating number of successes or a vector with 2 values indicating number of successes and failures
- n The parameter is used to specify the total number of trials. It is unused
  if the first parameter is a vector.
- p The hypothesis for the probability of success. It has a default value of 0.5.
- alternative This parameter should be one of "two.sided", "less" or "greater"
  - two.sided implies the true probability of success should not be equal to the value of parameter 'p'
  - less implies the true probability of success should be lesser than the value of parameter 'p'
  - greater implies the true probability of success should be greater than the value of parameter 'p'
- conf.level The confidence level which has a default value of 0.95

# Single Classifier Assessment APPROXIMATE NORMAL TEST FOR SINGLE CLASSIFIER USING SINGLE VALIDATION SET (19.10.2)

#### **Assumptions & Problem Statement**

- Assumptions:
  - The training is done on the single training set
  - The single validation set (non-overlapping with the training set) is used for the assessment.
  - For large N,  $\frac{X}{N} \sim N\left(p_0, \frac{p_0(1-p_0)}{N}\right)$  is approximately normal distribution with the mean  $p_0$  and the variance  $\frac{p_0(1-p_0)}{N}$ 
    - where X is the total number of errors
- Hypothesis Testing using qqnorm():
  - Let's assume that p is the probability that the classifier makes a misclassification error
  - Estimate or test a hypothesis about the probability p

#### **Approximately Normal Test**

Step 1.  $H_0$ :  $P <= P_0 \text{ vs } H_1$ :  $P > P_0$ 

Step 2.  $H_0$ : One-tail test. The critical value is 1.64 for  $\alpha$ =0.05

Step 3. The test value is

$$z_{\text{stat}} = \frac{X / N - p_0}{\sqrt{p_0 (1 - p_0) / N}}$$

#### Step 4. Reject the null hypothesis if Z<sub>stat</sub>>1.64, otherwise fail to reject

- Assumptions:
  - N is not too small and p is not very close to 0 or 1
  - as a rule of thumb: Np>=5 and N(1-p)>=5.

# Single Classifier Assessment T-TEST FOR SINGLE CLASSIFIER USING K-FOLD CROSSVALIDATION SETS (19.10.3)

help (qqnorm) help(t.test)

#### When and how to use?

- The t-test is used to assess the error rate of a single classification algorithm using K-fold cross-validation results.
- Assumptions
  - The training is done on the K-fold training set
  - The K-fold cross-validation result sets (non-overlapping with the corresponding training set for each fold) are used for the assessment.
- Hypothesis Testing using t.test():
  - classification algorithm has  $p_0$  or less error percentage at significance level  $\alpha$  using the t-statistic with parameters  $\alpha$  and (K-1):  $t_{\alpha,K-1}$

#### t-Test using *t*-statistic: $t_{\alpha,(K-1)}$

- Let the algorithm run K times, on K training/ validation set pairs:
  - then  $p_i$ , i = 1, 2, ..., K are error percentages on K validation sets
- Let  $x_i^t$  denote the correctness of the classifier on the  $i^{th}$  training set:
  - it is 1 when the classifier commits the error on the corresponding  $i^{th}$  validation set of the pair; 0, otherwise
  - Then  $p_i = \frac{1}{N} \sum_{t=1}^{N} x_i^t$  is the error percentage for fold i
- The mean and the variance across all the K folds:
  - $m = \frac{1}{K} \sum_{i=1}^{K} p_i$
  - $S^2 = \frac{1}{K-1} \sum_{i=1}^{K} (p_i m)^2$
- The *t*-statistic with (K-1) degrees of freedom:  $t_{K-1} \sim \frac{\sqrt{K}(m-p_0)}{S}$
- The Null Hypothesis  $(H_0)$ :  $p < p_0$
- The t-test rejects  $H_0$  at significance level  $\alpha$  (e.g.,  $\alpha = 0.05$ ):
  - if  $t_{K-1} > t_{\alpha,(K-1)}$  (e.g.,  $t_{0.05,9} = 1.83$  for K=10)

#### TWO Classifier Assessment

### McNEMAR'S TEST FOR TWO CLASSIFIERS USING SINGLE VALIDATION SET (19.11.1)

help (mcnemar.test)

#### When and how to use?

- The McNemar's test is used to assess whether the two classification algorithms have the same error rate using a single validation set.
- Assumptions
  - Two different classification algorithms
  - The training is done on the single training set
  - The single validation set (non-overlapping with the training set) is used for the assessment.
- Hypothesis Testing using mcnemar.test():
  - McNemar's test rejects the hypothesis that the two classification algorithms have the same error rate at significance level  $\alpha$  if the proper  $\chi^2$ -statistic is greater than  $\chi^2_{\alpha,1}$

#### McNemar's Test using $\chi_1^2$ -statistic

$e_{00}$ : number of examples misclassified by both	$e_{01}$ : number of examples misclassified by 1 but not 2
$e_{10}$ : number of examples misclassified by 2 but not 1	$e_{11}$ : number of examples correctly classified by both

- Under the null hypothesis (H<sub>0</sub>):
  - both classification algorithms have the same error rate

• 
$$e_{01} = e_{10} = (e_{01} + e_{10})/2$$

The chi-squared statistic with one degree of freedom:

$$\left| \chi_1^2 \sim \frac{(|e_{01} - e_{10}| - 1)^2}{e_{01} + e_{10}} \right|$$

- McNemar's test rejects the hypothesis that the two classification algorithms have the same error rate at significance level  $\alpha$  if the proper  $\chi_1^2$ -statistic is greater than  $\chi_{\alpha,1}^2$ :
  - for  $\alpha = 0.05$ ,  $\chi^2_{0.05,1} = 3.84$

#### **TWO Classifiers Assessment**

## PAIRED 7-TEST FOR TWO CLASSIFIERS USING K-FOLD CROSS-VALIDATION SETS (19.11.2)

t.test (..., paired=TRUE)

#### When and how to use?

- The paired t. test is used to assess whether the two classification algorithms have the same error rate using K-fold cross-validation results.
- Assumptions
  - Two different classification algorithms
  - The training is done on the K-fold training set
  - The K-fold cross-validation result sets (non-overlapping with the corresponding training set for each fold) are used for the assessment.
- Hypothesis Testing using t.test(paired=TRUE):
  - Use paired test; that is, for each i, both algorithms see the same training and validation sets

#### Paired t-Test using t-statistic: $t_{K-1}$

- Let two algorithms run K times, on K training/ validation set pairs:
  - then  $p_i^1$  and  $p_i^2$ , i = 1, 2, ..., K are error percentages on K validation sets for algorithm 1 and 2, resp.
- If the two classification algorithms have the same error rate, then
  we expect them to have the same mean, or equivalently, that the
  difference of their means is 0.
  - The difference in error rates on fold i:  $p_i = p_i^1 p_i^2$
  - Assuming that each  $p_i^1$  and  $p_i^2$  is normal, so is  $p_i$
- The Null Hypothesis  $(H_0)$ :  $p_i$  has the mean  $\mu = 0$ 
  - $H_1$ :  $\mu \neq 0$
- The mean and the variance across all the K folds:
  - $\bullet m = \frac{1}{K} \sum_{i=1}^{K} p_i$
  - $S^2 = \frac{1}{K-1} \sum_{i=1}^{K} (p_i m)^2$
- The *t*-statistic with (K-1) degrees of freedom:  $t_{K-1} \sim \frac{\sqrt{K}(m-\mu)}{S} = \frac{\sqrt{K}m}{S}$

#### K-fold CV Paired t-test: $t_{\alpha,(K-1)}$

- Two classification algorithms have the same error rate at significance level  $\alpha$  if  $t_{K-1}$  value is outside the interval:
  - $\bullet \quad (-t_{\frac{\alpha}{2},K-1};\ t_{\frac{\alpha}{2},K-1})$
  - Example: K = 10 and  $\alpha = 0.05$ ,  $t_{0.025,9} = 2.26$

21

# TWO Classifiers Assessment 5X2 CV PAIRED *T*-TEST FOR TWO CLASSIFIERS (19.11.3)

t.test (..., paired=TRUE)

#### When and how to use?

- The paired t. test is used to assess whether the two classification algorithms have the same error rate using 5x2 cross-validation results.
- Assumptions
  - Two different classification algorithms
  - The 5x2 cross-validation result sets are used for assessment
- Hypothesis Testing using t.test(paired=TRUE):
  - Use paired test; that is, for each i, both algorithms see the same training and validation sets

#### 5x2 CV Paired t Test

- 5x2 CV Paired t-Test is a simplified variation of the k-fold crossvalidation t-test.
- To perform the test, the following procedure is followed:
  - Perform 2-fold cross validation 5 times:
    - Generating 10 (training-testing) data set pairs
  - 2. For each pair:
    - Train both algorithms on the training data
    - Test on the test data
    - Compute the difference between the accuracies of the two algorithms

#### 5x2 CV Paired t Test

 Once you have these differences, compute the tstatistic of the first training-test pair (the difference of accuracies between the two classifiers on the first run's first fold)

$$\frac{\Delta_{1,1}}{\left(\frac{1}{5}\sum_{i=1}^{5} \left(\Delta_{1,i} + \frac{\Delta_{1,i} + \Delta_{2,i}}{2}\right)^{2} + \left(\Delta_{2,i} + \frac{\Delta_{1,i} + \Delta_{2,i}}{2}\right)^{2}\right)^{.5}}$$

#### Paired t-Test using t-statistic: $t_{K-1}$

- Let two algorithms run K times, on K training/ validation set pairs:
  - then  $p_i^1$  and  $p_i^2$ , i = 1, 2, ..., K are error percentages on K validation sets for algorithm 1 and 2, resp.
- If the two classification algorithms have the same error rate, then
  we expect them to have the same mean, or equivalently, that the
  difference of their means is 0.
  - The difference in error rates on fold i:  $p_i = p_i^1 p_i^2$
  - Assuming that each  $p_i^1$  and  $p_i^2$  is normal, so is  $p_i$
- The Null Hypothesis  $(H_0)$ :  $p_i$  has the mean  $\mu = 0$ 
  - $H_1$ :  $\mu \neq 0$
- The mean and the variance across all the K folds:
  - $\bullet m = \frac{1}{K} \sum_{i=1}^{K} p_i$
  - $S^2 = \frac{1}{K-1} \sum_{i=1}^{K} (p_i m)^2$
- The *t*-statistic with (K-1) degrees of freedom:  $t_{K-1} \sim \frac{\sqrt{K}(m-\mu)}{S} = \frac{\sqrt{K}m}{S}$