



# Author Verification: Exploring a Large set of Parameters using a Genetic Algorithm

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#### Task

Given a set of documents written by author A and an unknown document, find whether the latter was written by A.

# Output: probability in [0, 1]

- Evaluation: product of
- ► Area under the ROC curve (**AUC**),
- c@1 (accuracy with "don't know" answer)

## Fine-grained strategy

#### Features

## Consistency

- how constant in known documents?
- at least two known documents
- ▶ std. dev., min-max range
- Divergence
- how specific to the author?
- against a reference corpus
- mean/median diff., Bhattacharrya

# Selected by the genetic algorithm

## Observation types

- Few features selected (from 3 to 11)
- ► POS *n*-grams, words *n*-grams
- word length, TTR
- stop-words *n*-grams

# Methods

Bhattacharrya (divergence measure)

- Find an optimal configuration:
- set of parameter/value pairs
- methods, features, thresholds...
- Regression model based on config
- SVM, decision trees (variants)
- optional: confidence estimation

#### Genetic learning

- vast space: about 10<sup>19</sup> configurations
- maximize performance
- risk of overfitting

### Uses a reference corpus

- assuming variability among authors
- using all documents in the dataset

#### Confidence

- how reliable?
- uses consistency and divergence

#### Distance

- compare known vs. unknown doc
- Cosine, Jaccard, normal distrib

# Genetic algorithm

- Basic genetic learning
- Selecting configs as "breeders":
- rank the configs by their performance
- better perf = higher probability
- pick any two breeders as parents

Consistency unused in most cases

- Simple distance metrics (e.g. cosine)
- Decision tree regression
- Confidence estimation unused

# Selection of the final models

Evaluation on the earlybird test set Hypothesis: robust strategy better if only one known document?



### Robust strategy

- A simple distance measure
- Words tetragrams only
- Divergence based on Jaccard sim.:

 $J_1 = \frac{(p+q)}{(p+q+r)}$   $J_2 = \frac{(p+r)}{(p+q+r)}$ 

with *p* words in both *X* and *Y*, *q* words in *X* but not in Y, and r words in Y but not in X

# Observation types

- ► *n*-grams
- tokens, characters, POS tags
- Combinations with skip-grams ► **e.g.** "<token> \_\_\_ <POS tag>"

crossover, mutation

#### variants: elitism, random



Avg. perf. by generation, main learning stage. Parameters: population, breeders prop.;

0.00-												
	DE	DR	ΕΈ	ΕN	ĠA	SA data	DE	DR	ΕΈ	ΕN	ĠA	SA

Dataset	Known docs	/case	Strategy	Perf. training	Perf. Earlybird	Perf. drop	Diff. average
Dutch occave	mean	1.79	robust	0.802	0.777	-0.025	+0.103
Duich essays	median	1	fine-g.	0.817	0.501	-0.316	-0.071
Dutch reviews	mean	1.02	robust	0.389	0.338	-0.051	+0.077
	median	1	fine-g.	0.608	0.253	-0.355	-0.111
English essays	mean	2.64	robust	0.292	0.265	-0.027	+0.101
	median	3	fine-g.	0.493	0.446	-0.047	+0.198
English novels	mean	1.00	robust	0.722	0.324	-0.398	-0.270
	median	1	fine-g.	0.860	0.370	-0.490	-0.245
Greek articles	mean	2.85	robust	0.359	0.246	-0.113	+0.015
	median	3	fine-g.	0.595	0.541	-0.054	+0.191
Spanich articlas	mean	5.00	robust	0.622	0.468	-0.154	-0.026
opanish articles	median	5	fine-g.	0.863	0.657	-0.206	+0.039
Correlation bo	oce by caso	robust	0.77				
COnciation De	UUS DY LASE	fine-g.	0.03				

# Results

Datasot	Training set CV		Ea	rlybird test s	set	Final test set				
Dalasel	robust	fine-grained	robust	fine-grained	mixed	robust	fine-grained	final	rank	
Dutch essays	0.802	0.817	0.777	0.501	0.777	0.755	0.563	0.777	4	
Dutch reviews	0.389	0.608	0.338	0.253	0.338	0.375	0.350	0.375	3	
English essays	0.292	0.493	0.265	0.446	0.446	0.325	0.372	0.372	3	
English novels	0.722	0.860	0.324	0.370	0.324	0.313	0.352	0.313	8	
Greek articles	0.359	0.595	0.246	0.541	0.541	0.436	0.565	0.565	3	
Spanish articles	0.622	0.863	0.468	0.657	0.657	0.335	0.634	0.634	2	
Macro-average	0.531	0.706	0.403	0.461	0.514	0.423	0.473	0.502	3	
Micro-average								0.451	4	

stop-words *n*-grams n-grams, only most frequent words ▶ **e.g.** "the \_\_\_\_ is \_\_\_"

Token length classes ▶ e.g. 2-3, 3-4, 5-6, 6-7, 8-9, 10+

Token-Type Ratio

Thresholds: min. frequency in a document, min. proportion of documents which contain the observation (known docs, ref corpus)

mutation probability; elitism prop.; random prop.

Quick convergence in every case

Small population sufficient

More stable with larger population

14 000 to 28 000 configs evaluated

main training: 3-fold cross-validation

final stage (best subset): 20-fold CV

Hypothesis does not hold

Selecting strategy by dataset better

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