

Computational Literary Text Analysis (1): an Overview

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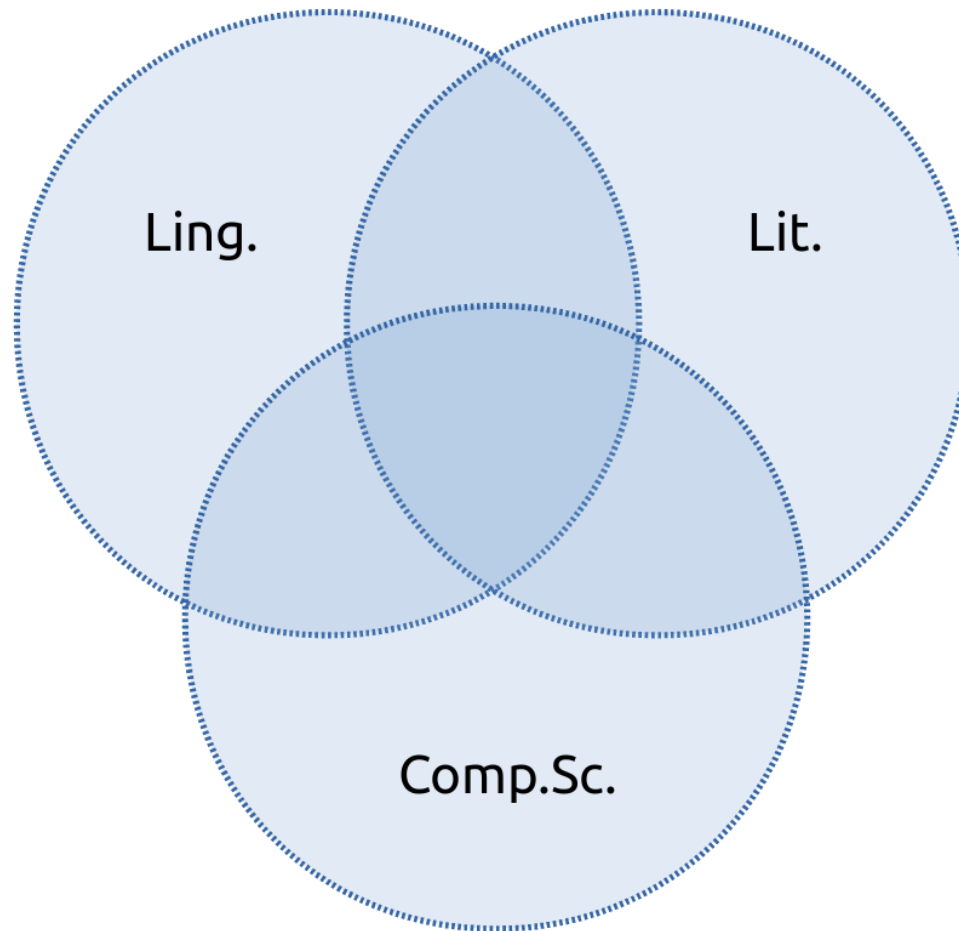


Overview

1. What is computational literary text analysis?
2. Stylometric Similarity Analysis
3. Machine Learning in Computational Narratology
4. Thematic Analysis with Topic Modeling
5. Conclusion

1. What is computational literary text analysis?

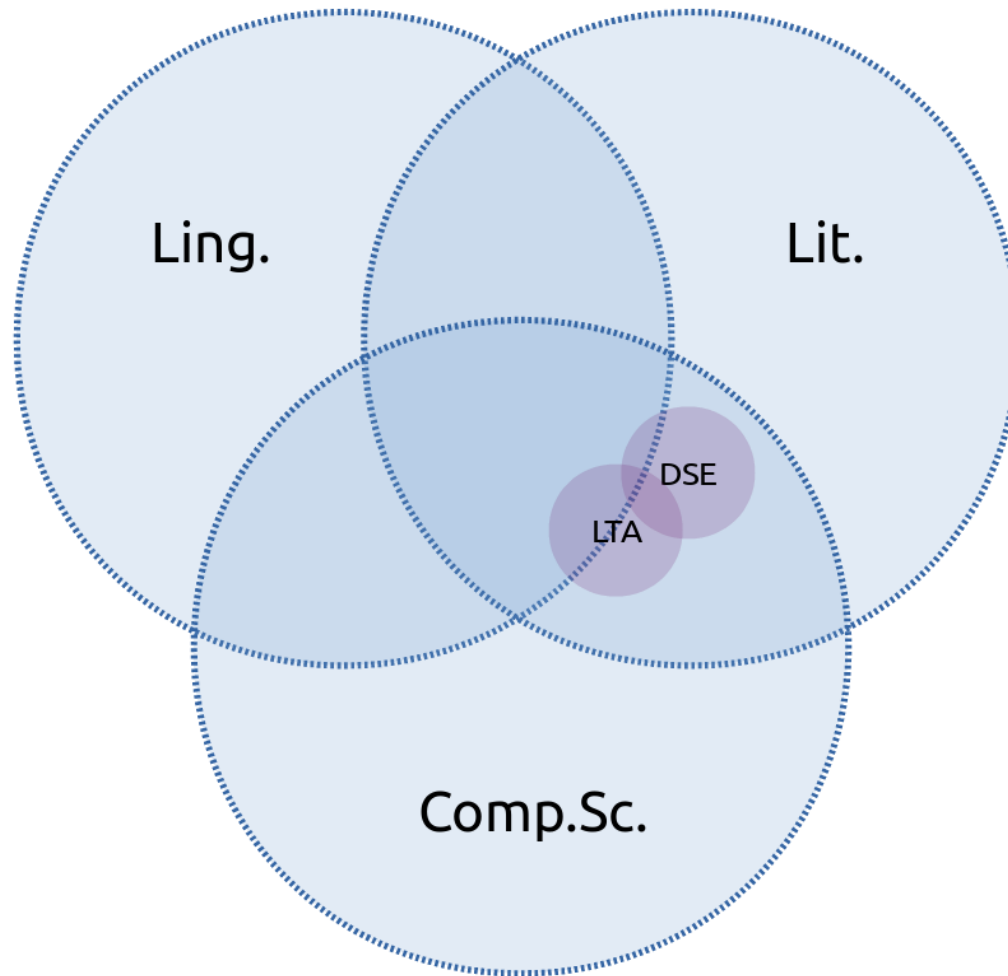
Literary Computing



Literary Computing

- Creating Texts: Digitisation, Digital Scholarly Editing (TEI), Text Collections
- Analysing Texts: Style, Content, Space, Time, Characters; Authors, Genres, Periods, etc.

Literary Text Analysis



(DSE = Digital Scholarly Editing, LTA = Computational Literary Text Analysis)

Some distinctions

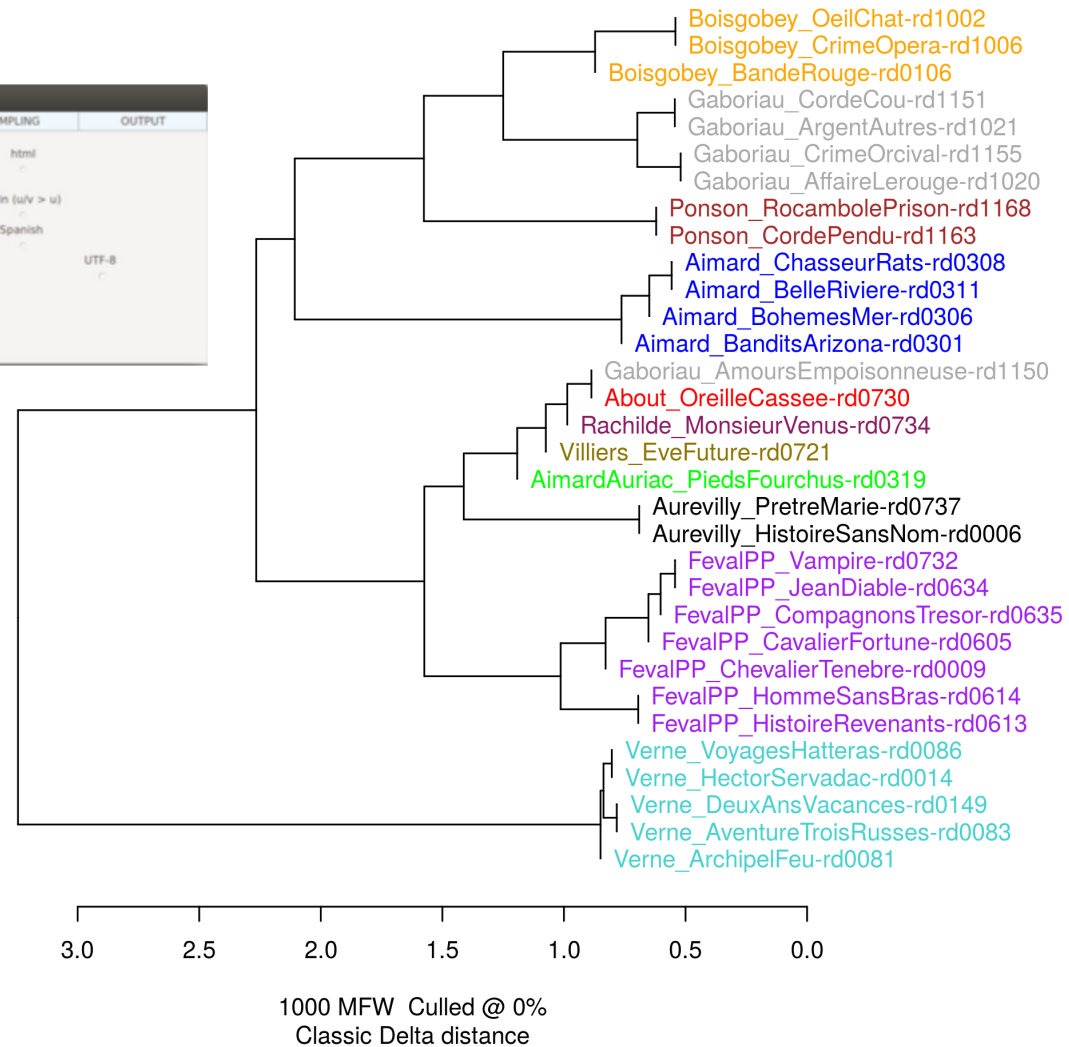
Some areas of LTA

Some techniques

2. Stylometric Similarity Analysis

(With thanks to Maciej Eder, Jan Rybicki, Stefan Evert, Fotis Jannidis, Thorsten Vitt, Steffen Pielström, ...)

stylo for R



Stylometric Similarity Analysis

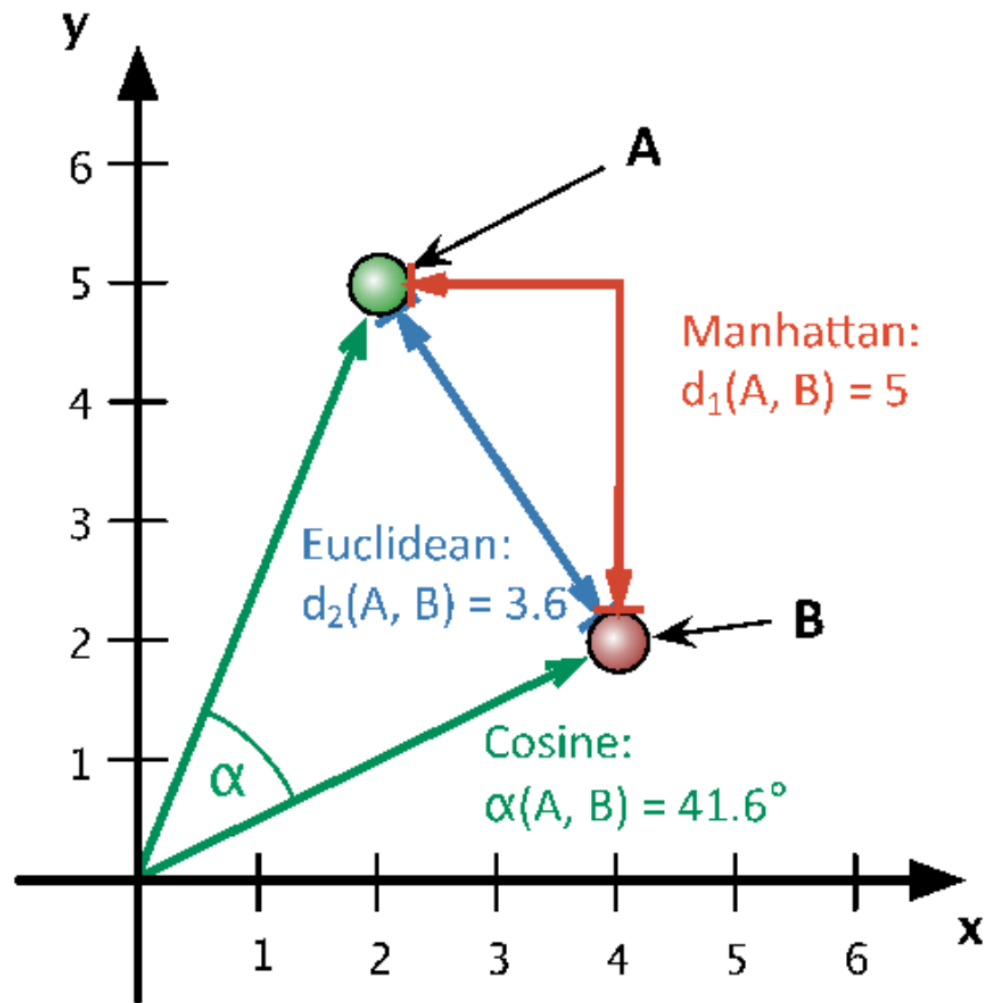
- Clearly quantitative method
- Usually using surface features
- Can be used for clustering or classification
- There is also a GUI ;-)

How does stylometric similarity work?

Texts as word frequency vectors

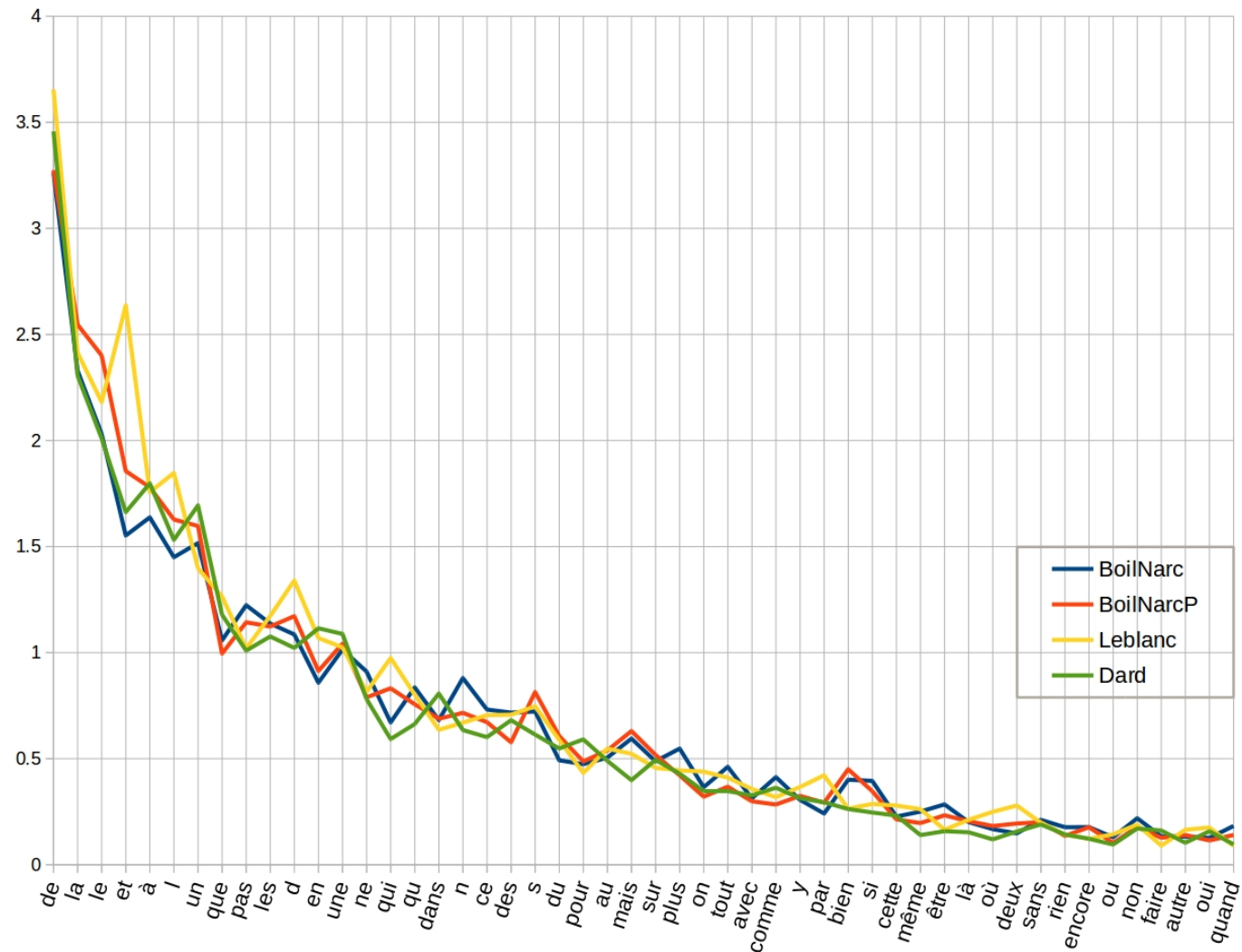
	A	B	C	D	E	F	G
1		Leblanc_1934=rp046.txt	Leblanc_1935=rp039.txt	BoilNarcP_1974=rp260.txt	BoilNarcP_1975=rp261.txt	BoilNarc_1955=rp253.txt	BoilNarcP_1979=rp263.txt
2	de	0.0364230093	0.0360980047	0.0324750152	0.0342983665	0.0345434543	0.0319564549
3	la	0.0262979924	0.0222841226	0.0270066417	0.0247315871	0.0215821582	0.0231113647
4	il	0.0224141596	0.023743202	0.0273628193	0.0256715545	0.0080008001	0.026798648
5	le	0.0210809035	0.0223030716	0.0204487838	0.025232903	0.0174817482	0.0233747421
6	et	0.0250420265	0.0272677316	0.0184583796	0.0173998412	0.0159615962	0.0181291427
7	à	0.0182791336	0.0168646846	0.0173479436	0.0184024732	0.0173617362	0.0173390106
8	l	0.0213707418	0.018986982	0.0173688952	0.0155825709	0.0123012301	0.016658619
9	je	0.0082893745	0.0103461998	0.0122357477	0.0144337219	0.0333433343	0.0113471753
10	un	0.0146464939	0.0137570348	0.0151899265	0.0168776371	0.0147414741	0.0146613406
11	d	0.0141634301	0.0144581509	0.0096586981	0.0118227013	0.0117211721	0.0125982178
12	pas	0.0111297896	0.0115020939	0.0115443441	0.0118644776	0.0114611461	0.0115886045
13	que	0.0118640465	0.0134349004	0.0103920049	0.0103396416	0.0118211821	0.0100302884
14	elle	0.0095260178	0.0068974665	0.0086530202	0.0044074028	0.0195619562	0.0104473026
15	les	0.0097192433	0.0087544767	0.0128223931	0.0109662865	0.0115811581	0.0100302884
16	une	0.0102216297	0.0092282038	0.0096796497	0.0109453983	0.0101210121	0.0101839252
17	est	0.0121538848	0.0108388759	0.0110624568	0.0099845428	0.0086408641	0.0122909442
18	vous	0.0107433386	0.0101946071	0.0126547801	0.0154990183	0.0083208321	0.0119178263
19	en	0.0103568875	0.0107441305	0.008883488	0.0090445753	0.0080208021	0.0097010667
20	ne	0.0083666647	0.0092471529	0.0079616166	0.0081046079	0.0097209721	0.0079671656
21	qui	0.010337565	0.0099293199	0.0074587777	0.00887747	0.0072007201	0.0086695053
22	n	0.0072846019	0.0079396661	0.0073121163	0.0068722062	0.0079807981	0.0075062552
23	qu	0.0080381814	0.0077880734	0.0079825682	0.0065171074	0.0064006401	0.00803301
24	s	0.0074198597	0.0069543138	0.008275891	0.0086477002	0.004760476	0.0075282033
25	était	0.0042896064	0.0053057435	0.0078568585	0.0064335547	0.0056605661	0.0081427505
26	se	0.0069367959	0.0068406193	0.0084016007	0.0080210553	0.00410041	0.0077696326
27	ce	0.0073618921	0.0075038372	0.006055019	0.0067886535	0.0065606561	0.0069795005
28	dans	0.0069174734	0.0075796335	0.0080873264	0.0060784559	0.0073807381	0.0061674202
29	des	0.0062798292	0.0058363178	0.0063902449	0.006976647	0.0064606461	0.0047188447
30	avait	0.0046760574	0.0047751691	0.0061178738	0.0048251661	0.004560456	0.0067380712
31	mais	0.004907928	0.0059689614	0.0064530998	0.0050967122	0.0056005601	0.0071989816

Vector similarity / distance



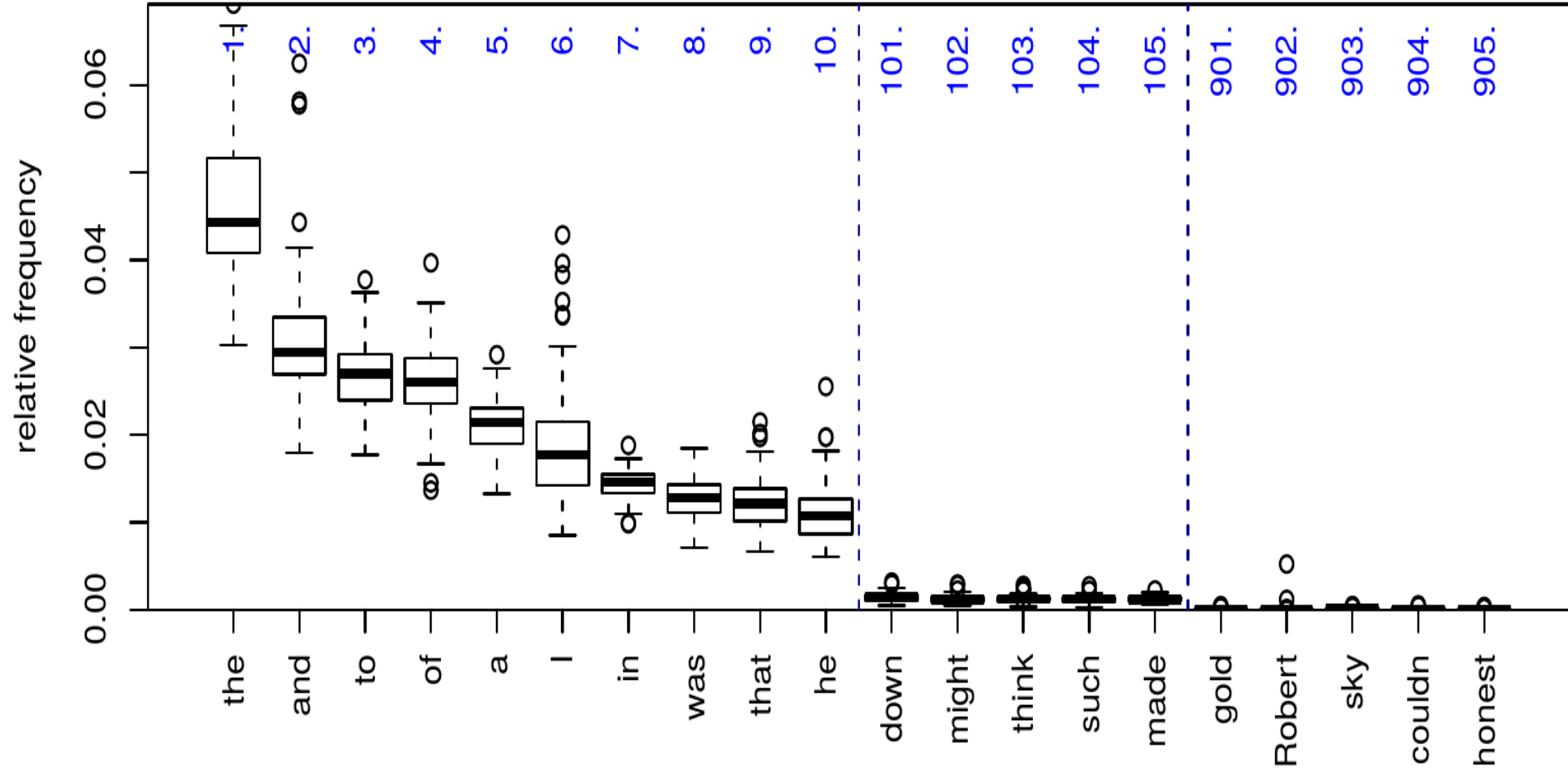
(Two texts, A and B; two words on x and y axes)

Frequency vectors plotted



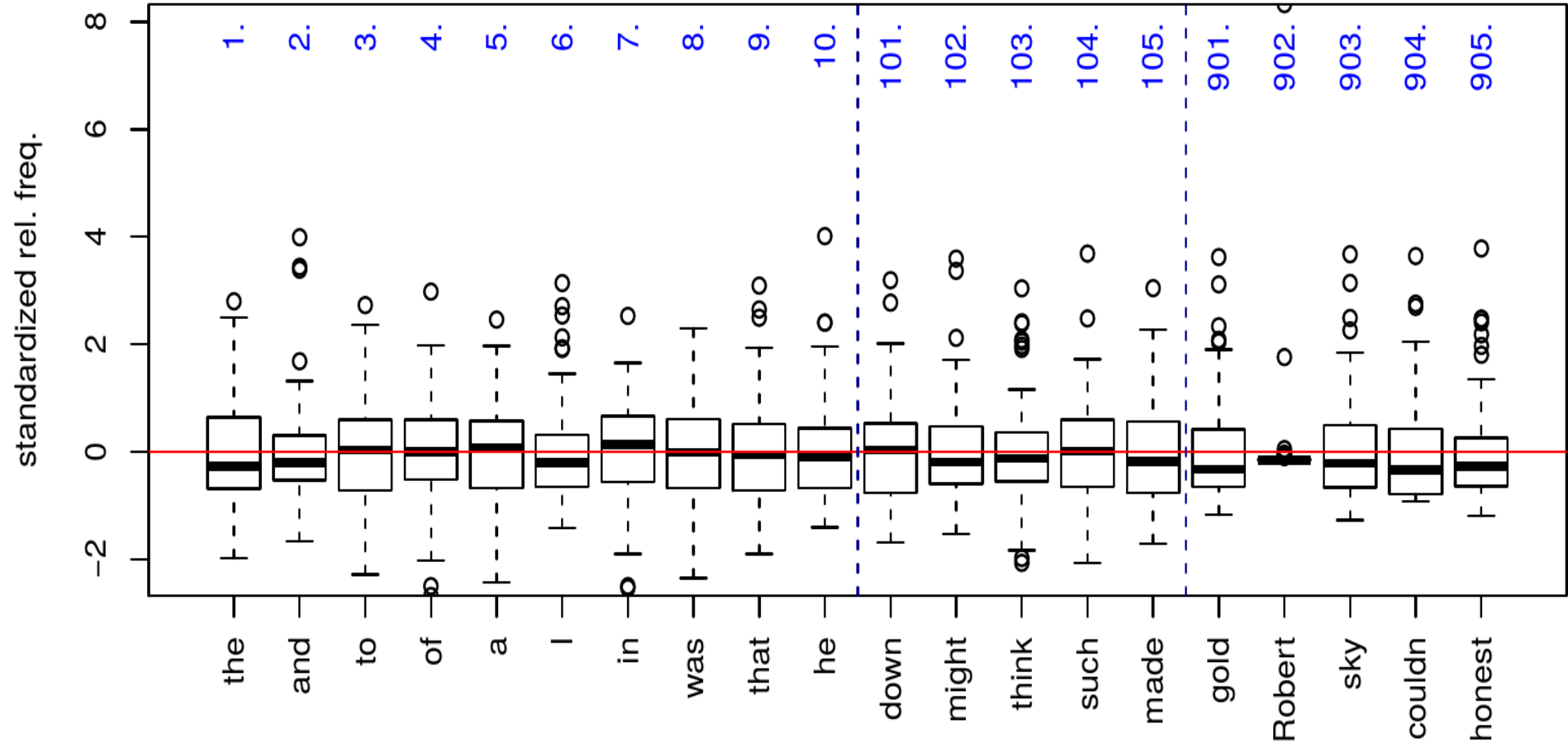
(One text for three/four different authors)

Distribution of relative frequencies



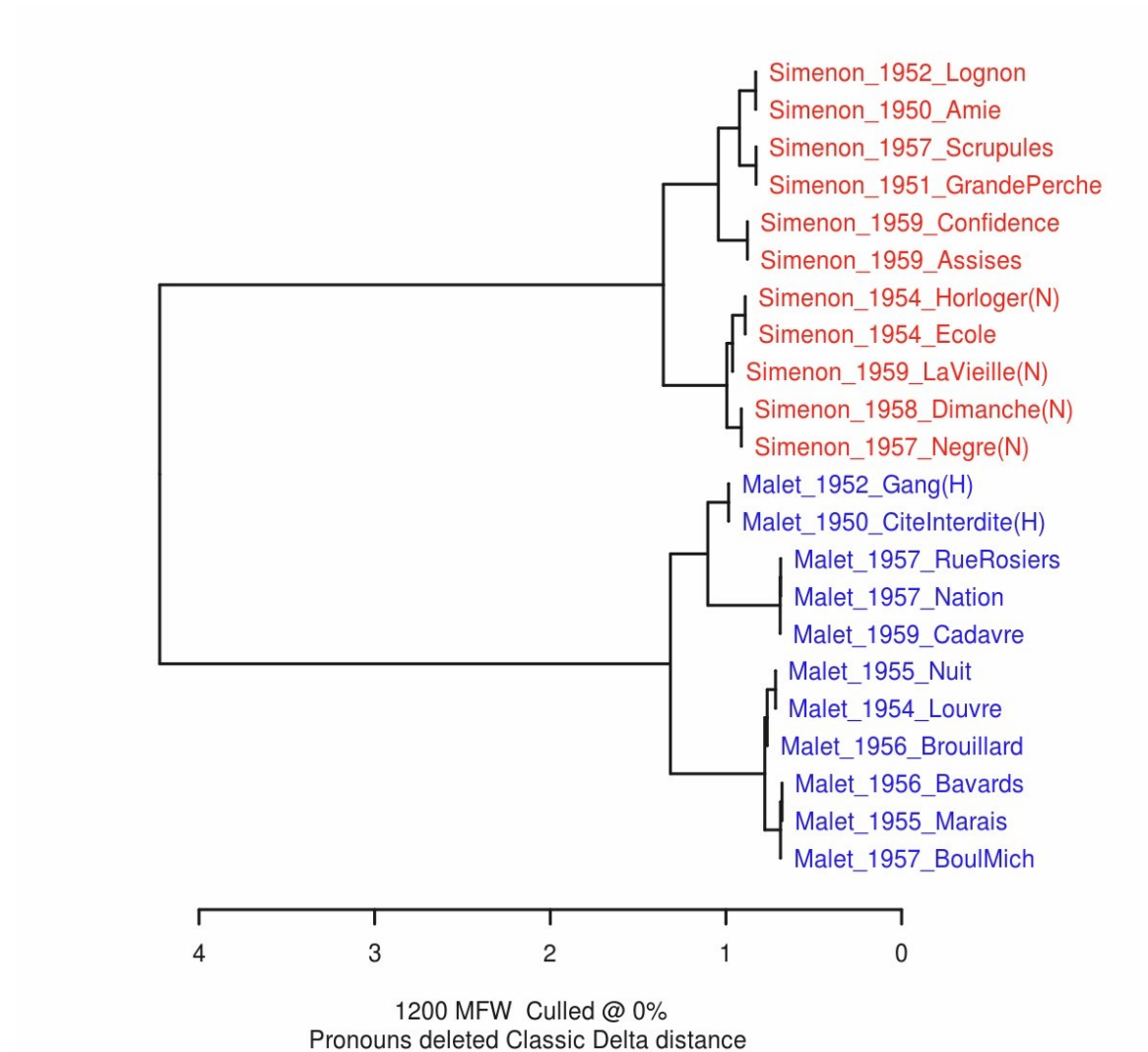
(High-frequency words dominate; graph by Stefan Evert)

Distribution of standardized frequencies



(This is what Burrows' Delta does; graph by Stefan Evert)

Dendrogram



(Based on the similarity matrix)

What does such similarity tell us?

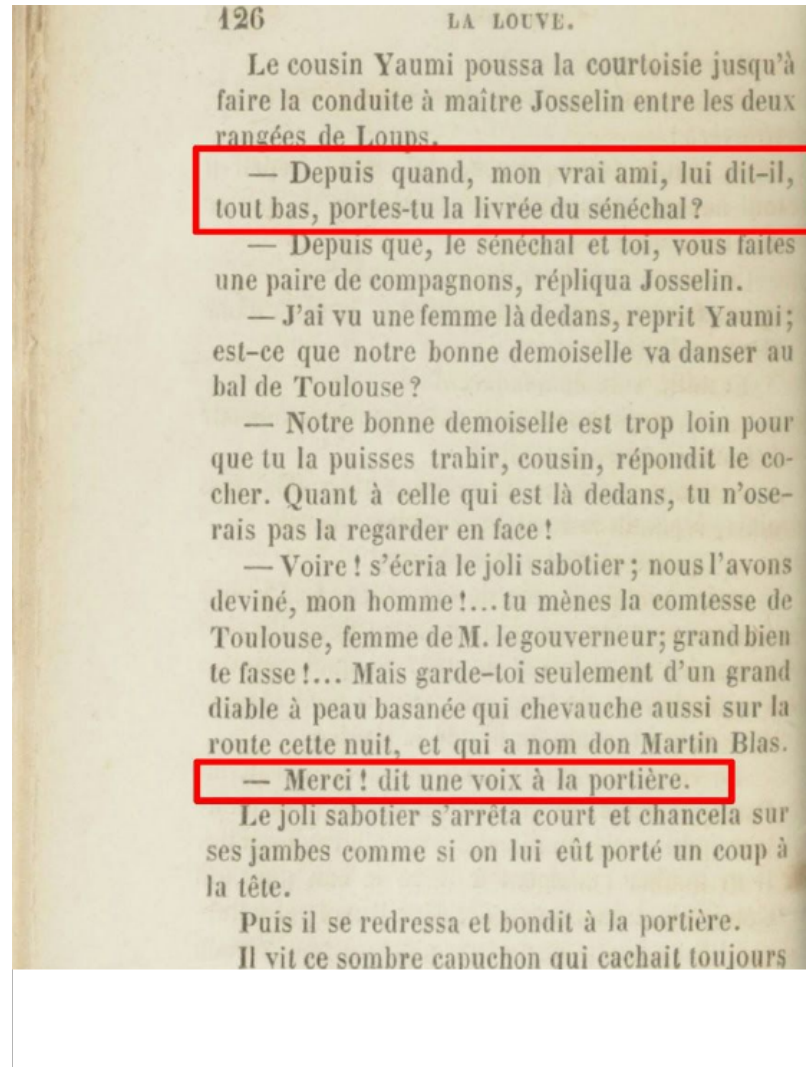
3. Machine Learning in Computational Narratology

(Example: Automatic Recognition of Direct Speech, with Daniel Schlör and
Stefanie Popp)

Automatic Recognition of Direct Speech

- Computational Narratology
- Relies on annotation and Machine Learning
- Necessitates feature generation
- Derives a higher-order feature from surface / lower-order features

What is the problem?



The corpus used

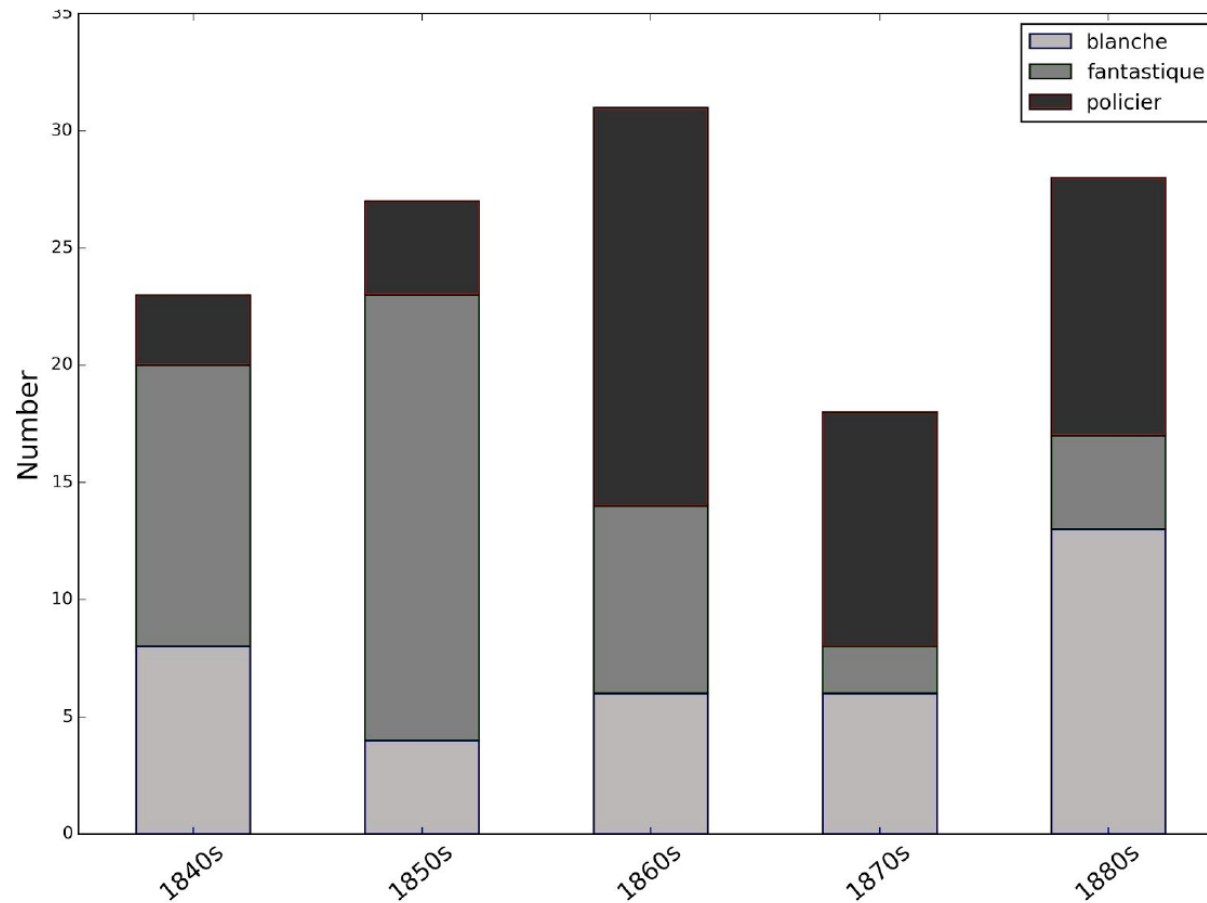


Figure 2: Distribution of novels per subgenre and decade.

127 French Novels, 1840-1890

The method

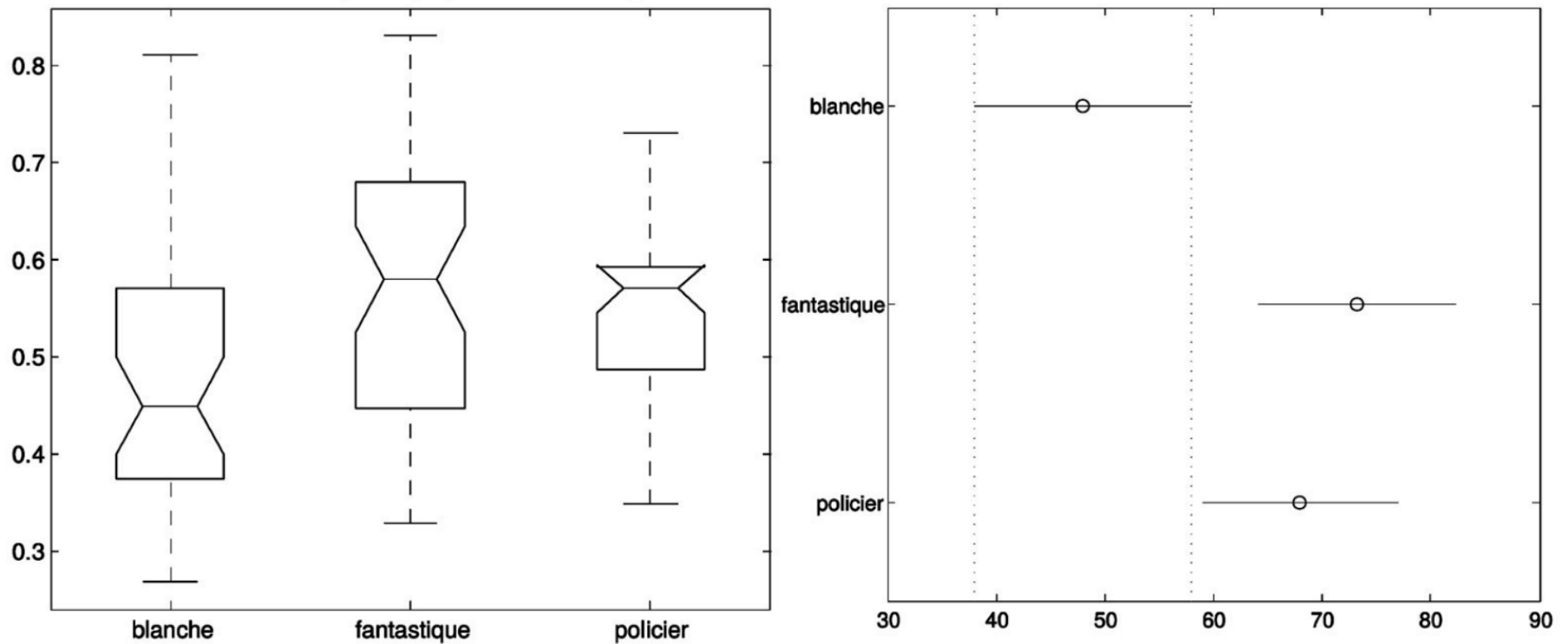
Feature generation (81 features)

Performance

	Direct speech (3222 Instances)			Non-direct speech (2512 Instances)			Weighted average (5734 instances)			Without Speechsign
	Precision	Recall	F1 Score	Precision	Recall	F1 Score	Precision	Recall	F1 Score	F1 Score
Baseline Speechsign	0.948	0.569	0.711	0.634	0.96	0.764	0.810	0.740	0.734	
N.Bayes	0.863	0.906	0.884	0.834	0.884	0.859	0.850	0.896	0.873	0.831
MaxEnt	0.894	0.887	0.89	0.856	0.865	0.861	0.877	0.877	0.877	0.847
JRip	0.881	0.912	0.896	0.882	0.842	0.861	0.881	0.881	0.881	0.849

Random Forests: F1-score 0.938

Results: Subgenres



***Figure 4: Distribution (left) and significance (right)
of direct to non-direct speech ratios across three subgenres***

Popular vs. literary subgenres

What is it useful for?

- Investigate subgenre preferences and long-term evolution
- Investigate differences between different character's styles
- Separate narrator and character speech for stylometry
- Identify narrator speech and further classify it (narrative, descriptive)

Top Features

average merit	average rank	attribute
74.028 +- 0.168	1 +- 0	79 SPEECHSIGN
71.743 +- 0.16	2 +- 0	57 VER:impf
65.847 +- 0.234	3 +- 0	54 VER:pres
63.893 +- 0.155	4 +- 0	55 VER:simp
63.248 +- 0.136	5 +- 0	6 PUNCTMARKDOT
59.48 +- 0.12	6 +- 0	29 MATCHINGPPER_SON
58.835 +- 0.094	7.7 +- 0.64	30 MATCHINGPPER_SES
58.695 +- 0.208	8.1 +- 0.94	24 MATCHINGPPER_IL
58.713 +- 0.104	8.4 +- 0.92	35 VERB_MOTION
58.364 +- 0.083	10.6 +- 0.49	28 MATCHINGPPER_SA
58.344 +- 0.417	10.8 +- 1.78	7 SENTENCELENGTH
58.172 +- 0.078	11.7 +- 0.46	61 VER:subi
57.492 +- 0.091	14 +- 1.41	25 MATCHINGPPER_ELLE
57.422 +- 0.103	14.5 +- 1.36	44 VERB_PERCEPTION
57.387 +- 0.248	14.9 +- 1.51	50 INNERSUBCLAUSE
57.356 +- 0.4	15.8 +- 2.09	48 UNKNOWNLEMMA
57.213 +- 0.07	16.5 +- 1.02	31 MATCHINGPPER_LEUR
57.143 +- 0.162	17.3 +- 1.1	60 VER:ppre
56.672 +- 0.042	20.2 +- 0.98	36 VERB_BODY
56.672 +- 0.115	21 +- 1.84	52 VER:cond
56.62 +- 0.136	21.7 +- 2.1	40 VERB_EMOTION
56.567 +- 0.072	22.3 +- 1.19	26 MATCHINGPPER_ILS
56.497 +- 0.033	23.9 +- 1.3	41 VERB_COGNITION
56.428 +- 0.044	25 +- 1	46 VERB_CONSUMPTION

long hyphen, verb tense, personal pronouns

4. Thematic Analysis with Topic Modeling

Basic Idea

Words, Topics, Documents

Topics

gene	0.04
dna	0.02
genetic	0.01
...	

life	0.02
evolve	0.01
organism	0.01
...	

brain	0.04
neuron	0.02
nerve	0.01
...	

data	0.02
number	0.02
computer	0.01
...	

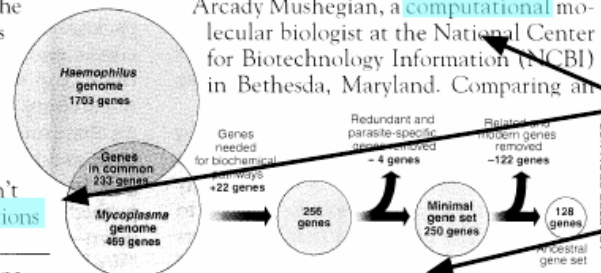
Documents

Seeking Life's Bare (Genetic) Necessities

COLD SPRING HARBOR, NEW YORK—How many **genes** does an **organism** need to **survive**? Last week at the genome meeting here,* two genome researchers with radically different approaches presented complementary views of the basic genes needed for **life**. One research team, using **computer** analyses to compare known **genomes**, concluded that today's **organisms** can be sustained with just 250 genes, and that the earliest life forms required a mere 128 **genes**. The other researcher mapped genes in a simple parasite and estimated that for this organism, 800 genes are plenty to do the job—but that anything short of 100 wouldn't be enough.

Although the numbers don't match precisely, those **predictions**

"are not all that far apart," especially in comparison to the 75,000 **genes** in the human genome, notes Siv Andersson of Uppsala University in Sweden, who arrived at the 800 number. But coming up with a consensus answer may be more than just a **genetic** numbers game, particularly as more and more **genomes** are completely mapped and sequenced. "It may be a way of organizing any newly **sequenced genome**," explains Arcady Mushegian, a **computational** molecular biologist at the National Center for Biotechnology Information (NCBI) in Bethesda, Maryland. Comparing an

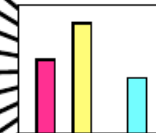


* Genome Mapping and Sequencing, Cold Spring Harbor, New York, May 8 to 12.

Stripping down. Computer analysis yields an estimate of the minimum modern and ancient genomes.

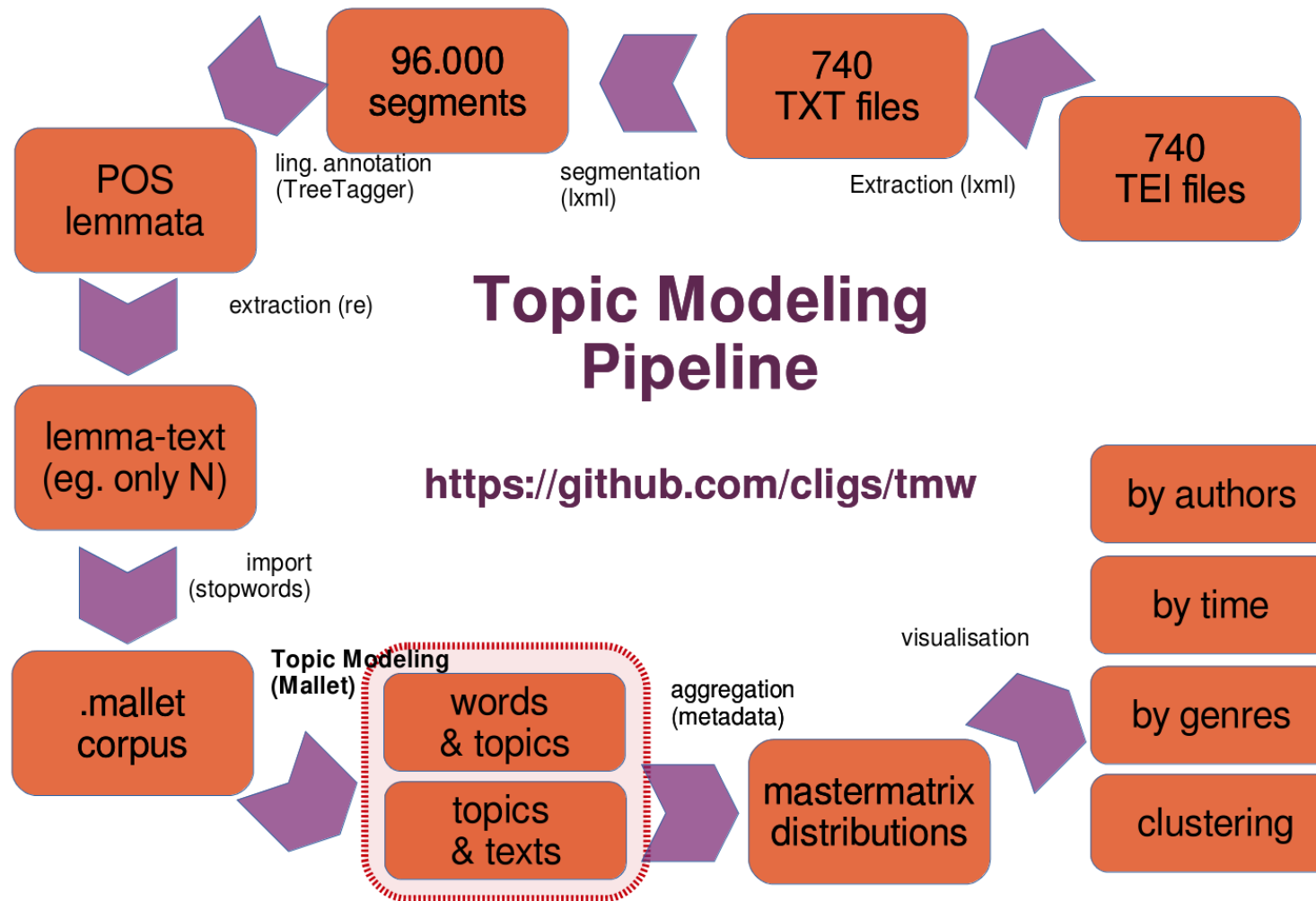
SCIENCE • VOL. 272 • 24 MAY 1996

Topic proportions and assignments



(David Blei, "Probabilistic Topic Models", 2012)

Preprocessing, Modeling, Postprocessing



Topics: Crime

topic 79 (73/240)

trace cadavre
déTECTIVE mort police découverte

Topics: Music

topic 45 (102/240)

œuvre ●

clavier

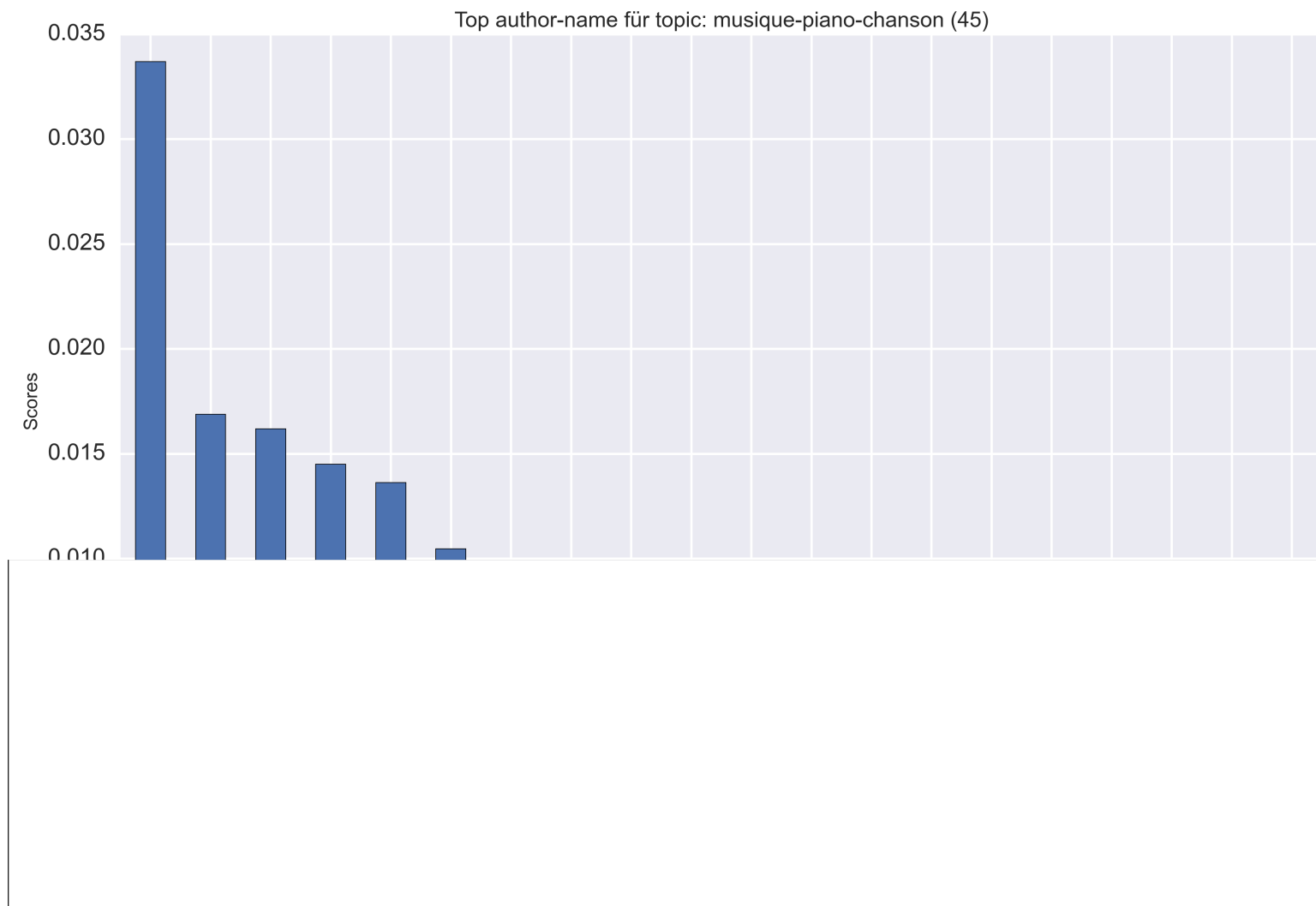
artiste

mesure ●

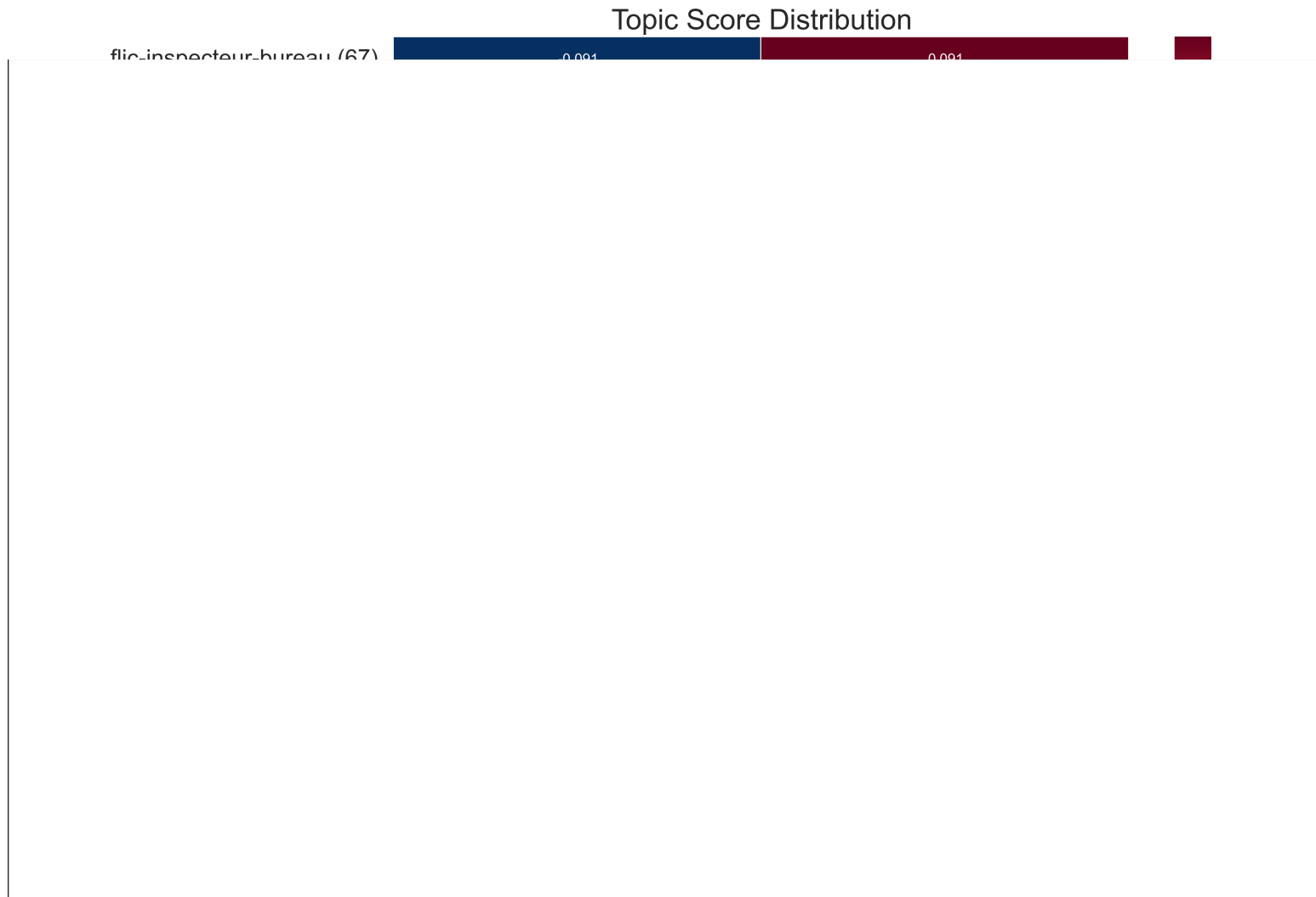
● mélodie



Top-Authors for Music



Crime fiction vs. non-crime fiction



Topics over time

What is it good for?

Conclusion

Conclusion

Further Reading

- Shillingsburg, P. (2006). *From Gutenberg to Google. Electronic Representations of Literary Texts*. Cambridge: Cambridge Univ. Press.
- Alpaydin, E. (2010). *Introduction to Machine Learning*. 2nd ed. Cambridge, Mass: MIT Press.
- Ramsay, S. (2011). *Reading Machines: Toward an Algorithmic Criticism*. Urbana Ill.: University of Illinois Press.
- *Doing Digital Humanities Bibliography*:
https://www.zotero.org/groups/doing_digital_humanities_-_a_dariah_bibliography

Thank you!

Christof Schöch, 2016

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