

Text Mining Techniques

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Overview

- Gentle introduction to TM
- Applications/case studies
 - To help understand what TM could do for you





NaCTeM

• First publicly-funded (JISC) national text mining centre in the world

Provide services to research community

- Initial focus on biology, then social sciences, medicine, chemistry, ...
- Processing on a large scale, e.g. for UKPMC (Wellcome T.+17 other funders)
- www.nactem.ac.uk





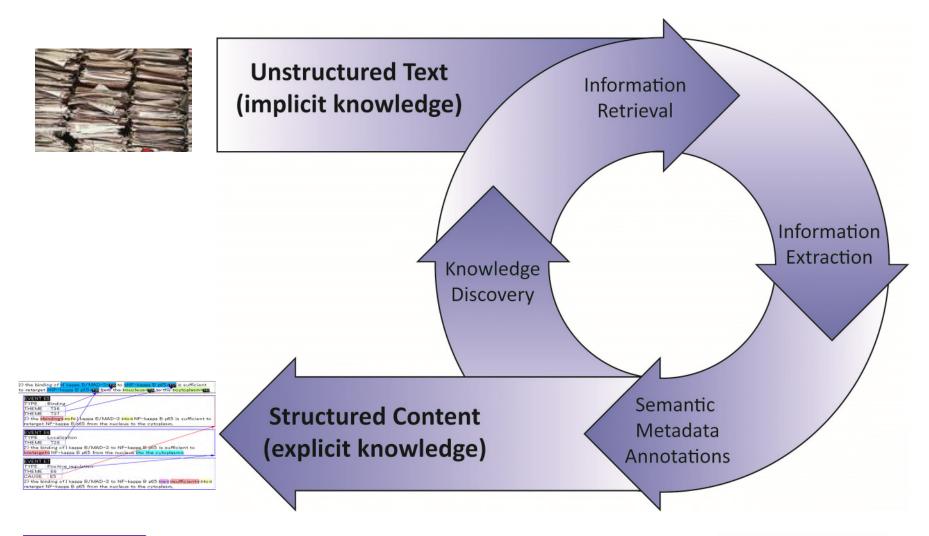
What is text mining?

- Goal: Discover new knowledge from old
- How:
 - Process (typically) very large amounts of text
 - Millions of documents not unusual
 - Identify and extract information
 - (Link extracted information to already curated knowledge)
 - Mine to discover implicit significant associations
 - Flag (unknown) associations for researcher to investigate further
 - Spin-off on the way: render information explicit





From text to new knowledge







What does TM offer?

- Finds unsuspected knowledge
 - E.g. Disease-gene associations
- Enables discoveries human effort could not achieve (information overload/overlook)
- Enables better search/navigation of literature
 - Semantic search via extracted semantic metadata
- Reduces time spent searching
 - 15-48% of researcher time spent on classic search, 20-50% of classic searches unsatisfied

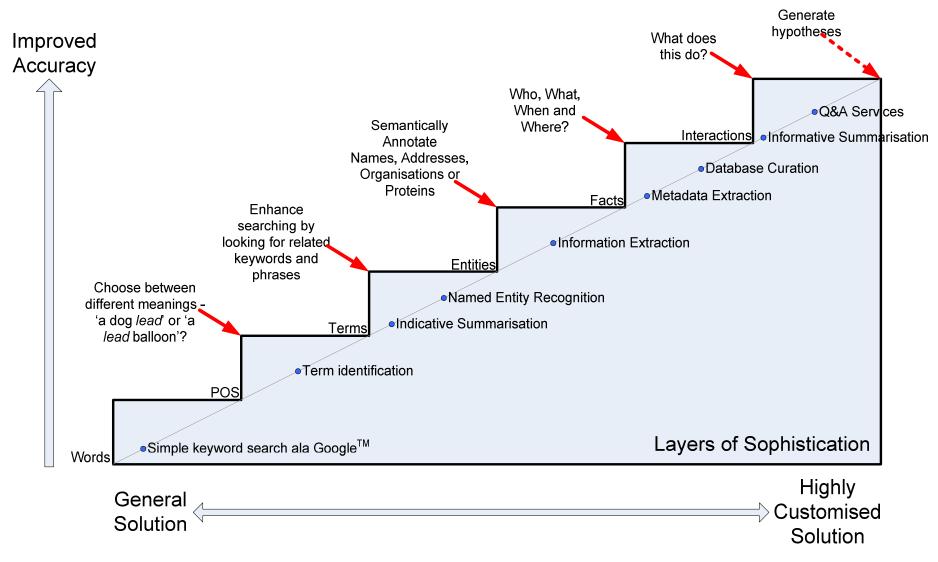
A complex problem

- TM involves
 - Many components (converters, analysers, miners, visualisers, ...)
 - Many resources (grammars, ontologies, lexicons, terminologies, thesauri, CVs)
 - Many combinations of components and resources for different applications
 - Many different user requirements and scenarios, training needs





Accurate mining of sophisticated knowledge demands customisation



Customisation

- Rule writing
 - Expert expresses knowledge in formalism
 - Issues of exhaustivity, specificity
- Machine learning (supervised)
 - Expert annotates large amounts of text for desired phenomena
 - Accelerated annotation editor (e.g. NaCTeM's Argo)
 - System 'completes' annotation as it learns from your decisions
 - Train machine learning algorithm
 - Apply trained model to find similar patterns of phenomena in new texts
- Hybrid approaches
- Knowledge poor vs. knowledge rich approaches





Brief aside

 Many of the systems and applications that follow are based on biomedicine

– A small intellectual leap required...

- Techniques are portable to other domains and text types
 - Modulo customisation...
 - Modulo understanding user requirements
 - What's a frame (for you, for your colleague)?





Producing multilevel annotated data: NaCTeM's Argo editor

Specialized secretion systems of pathogenic bacteria commonly transport multiple effectors that act in concert to control and exploit the host cell as a replication-permissive niche. Both the Mycobacterium marinum and the Mycobacterium tuberculosis genomes contain an extended region of difference 1 (extRD1) locus that encodes one such pathway, the early secretory antigenic target 6 (ESAT-6) system 1 (ESX-1) secretion apparatus. ESX-1) is required for virulence and for secretion of the proteins ESAT-6, culture filtrate protein 10 (CFP-10), and EspA. Here, we show that both Rv3881c and its M. marinum homolog, Mh3881c, are secreted proteins, and disruption of RD1 in either organism blocks secretion. We have renamed the Rv3881c/Mh3881c gene espB for ESX-1 substrate protein B. Secretion of M. marinum EspB (EspBM Remove Gene/Protein genes within RD1, while CFP-10 secretion is not affected by disru ruption of Change to Biological process Mh3866 or Mh3867 within the extRD1 locus prevents CFP-10 sec lutants that fail to secrete only EspBM or only CFP-10 are less attenuated in n Change to Cellular component to secrete both substrates. EspBM physically interacts with Mh3879c; the M. ysically Change to Metabolite interacts with Rv3879c; and mutants of EspBM that fail to bind Mh o found Change to Molecular Function interaction between Rv3879c and Rv3871, a component of the ES anism for the Change to Organism secretion of EspB. The results establish EspB as a substrate of E and growth Change to Unknown in macrophages and suggests that the contribution of ESX-1 to vir on of multiple independent substrates.

Extract from McLaughlin et al. (2007) in PLoS Pathogens

Flexible annotation

Mechanisms of Toxicity of 3-Alkylpyridinium Polymers from Marine Sponge Reniera sarai

Polymeric 3-alkylpyridinium salts (boly-APS) present in the marine sponge Reniera sarai show a broad spectrum of biological activities. They are lytic to erythrocytes and various other mammalian cells, enabling the transfection of the latter with alien DNA. Furthermore, they show inhibitory effects to marine bacteria and can inhibit fouling of micro- and macroorganisms to submerged surfaces. Finally, poly-APS act as potent cholinesterase inhibitors. The kinetics of acetylcholinesterase inhibition by poly-APS in vitro is complex and comprises several successive phases ending in irreversible inhibition of the enzyme. The latter is accounted for by aggregation and precipitation of the enzyme-inhibitor complexes. Poly-APS are lather to arts in concentrations above 2.7 mg/kg. Monitoring of the basic vital functions and histor MDG lysis showed that the effects directly ascribable to acetylcholinesterase inhibition are o Remove er application of lower concentrations of poly-APS. At higher concentrations, such e Change to ٠ ked by other, more pronounced and faster developing lethal effects of the toxin, such as Add AGENT aggregation. Properties COMMENT ENZ

V	AGENT
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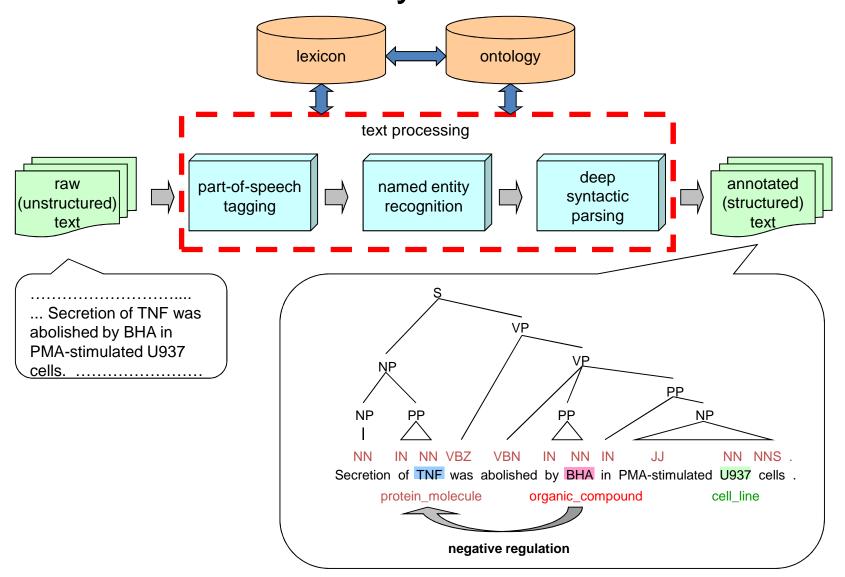
1. Introduction

Monomeric, oligomeric and polymeric <u>3-alkylpyridiniums</u> and <u>3-alkylpyridines</u> comprise a group of active compounds found in several sponges of the order Haplosclerida [1–4]. In general, the var potency of the biological effects of these compounds increase with their degree of polymerisation, recurring increase and unprecedented mechanisms of action and/or toxicity. <u>3-Alkylpyridinium polymers</u> isolated so far from haplosclerid marine sponges are shown inFigure 1. Among them, polymeric <u>3-alkylpyridinium salts</u> (poly-APS), isolated from crude extracts of the Mediterranean marine sponge Reniera sarai, show the highest degree of polymerisation.

Extract from Turk et al. (2007) in Marine Drugs

INTERACTION

Information extraction: many techniques at many levels



Semantic search using named entity recognition

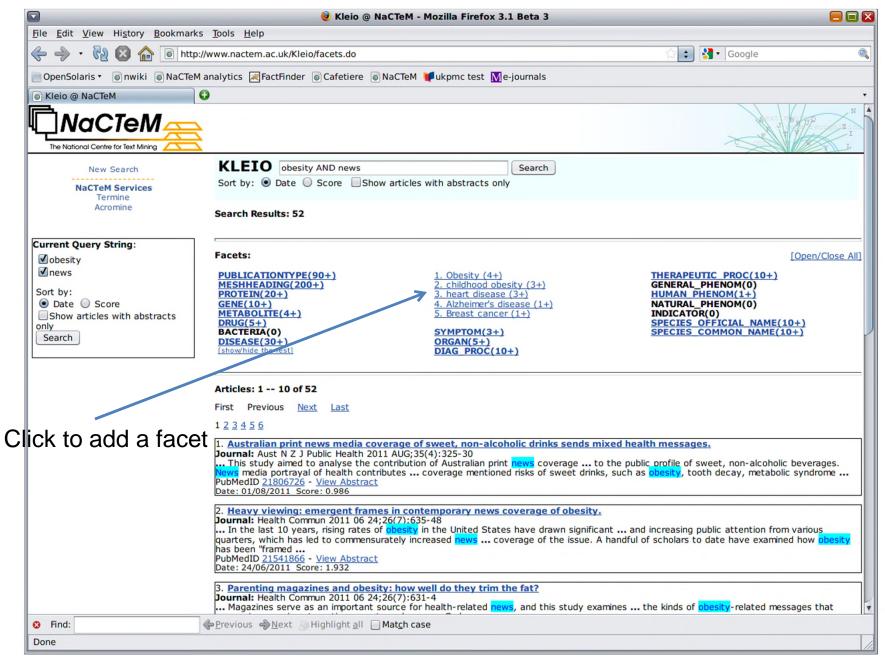
- Identify named entities in texts
 - People, organisations, locations, genes, drugs, ...
- Use entity types in searching

 Reduce hits to precise, relevant ones
- Make life easy by offering facetted search using the NEs
- NaCTeM's Kleio
- www.nactem.ac.uk/software/Kleio/





General search



Add a mined NE facet

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for stories regarding childhood obesity. Of the 201 stories appearing the framing of news regarding childhood obesity are discussed. PubMedID <u>21153987</u> - <u>View Abstract</u> Date: 01/12/2010 Score: 1.652						
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Noisy data

- Full-scale analysis techniques often not suitable for noisy data
 - OCR, digitised texts
 - Emails, unprofessional blogs, tweets, ...
- May have to use partial analysis techniques, often based on machine learning
- Even apparently 'clean' data can be hard to process
 - "Hamburger PDFs" (P. Murray-Rust)





Full-scale analysis

- For formally-written text, full-scale analysis possible
- Deep parsing with MEDIE
 - Allows semantic search
 - Query consists of partial or fully specified fact/event template
 - www.nactem.ac.uk/medie/
 - Developed by Univ. Tokyo (close collaboration)





MEDIE: semantic search of pre-analysed collection

Sentences from MEDLINE where news is the object, with obesity specified in (here unseen) advanced options

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What users want...

- Many users just give a one-word query when searching
 - Very few explore 'advanced search'
- So NaCTeM uses deep parsing results to generate questions relevant to a query that are known to have answers
 - UKPMC EvidenceFinder
 - http://labs.ukpmc.ac.uk
 - UKPMC: archive of full texts in life sciences
 - EBI, British Library, Univ of Manchester (Mimas and NaCTeM)





UKPMC EvidenceFinder

Semantic search based on analysis of full text articles

But currently, EF oriented more towards Biology... and can only text mine OA subset

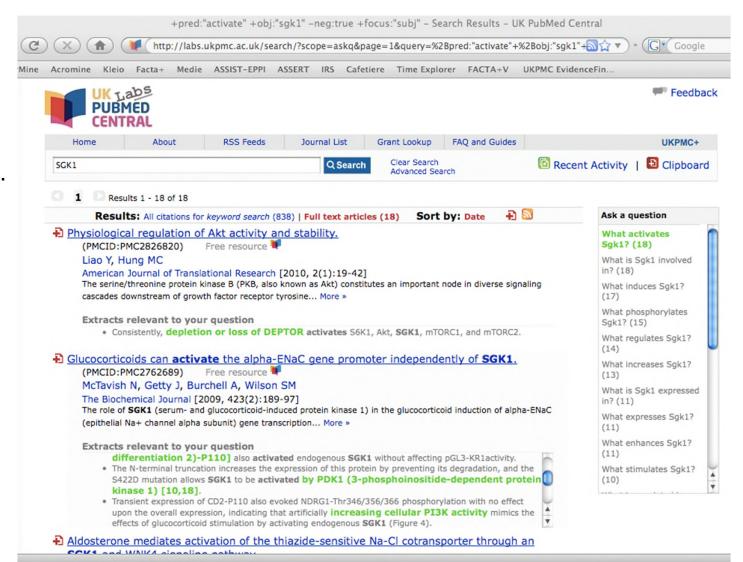
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	 Heavy viewing: emergent frames in contemporary news coverage of obesity. (PMID:21541866) Shugart HA Health Communication [2011, 26(7):635-48] In the last 10 years, rising rates of obesity in the United States have drawn significant and increasing public attention from various quarters, which has led to commensurately More * The Emergence and Portrayal of Obesity in The Irish Times: Content Analysis of Obesity 	What is Obesity associated with? (3010) What causes Obesity? (1615) What reduces Obesity? (554)			
	<u>Coverage, 1997-2009.</u> (PMID:21978090)	Popular content sets P			
	De Brún A, McKenzie K, McCarthy M, McGloin A Health Communication [2011] Both global obesity prevalence rates and media attention to obesity have increased significantly in recent years. The current study examined the representation of obesity More »	All results (92) Reviews (4)			
	Framing obesity: the framing contest between industry and public health at the New Zealand inquiry into obesity. (PMID:21831231) Jenkin GL, Signal L, Thomson G Obesity Reviews : an Official Journal of the International Association for the Study of Obesity [2011] Drawing on submissions to the 2006-2007 New Zealand Inquiry into Obesity and Type 2 Diabetes, this article outlines how the food and marketing industries (industry) and More »	More »			
	Obesity in the New Media: A Content Analysis of Obesity Videos on YouTube. (PMID:21809934) Yoo JH, Kim J Health Communication [2011] This study examines (1) how the topics of obesity are framed and (2) how obese persons are portrayed on YouTube video clips. The analysis of 417 obesity videos revealed More *				
🕴 Find:	⊕Previous → Next → Highlight all → Match case				
Done					

UKPMC EvidenceFinder

But asking about e.g. a protein gives better idea of functionality...

330 thousand OA full text articles analysed.

Generated questions known to have answers.



Finding associations

- Known associations (but not to you)
 Unknown knowns (the one DR missed out)
- Unknown associations (to anyone)
 - Statistics of surprise
 - A related to B, B to C, so infer A related to C
- www.nactem.ac.uk/services.php
- Check out FACTA+ and FACTA+ visualizer





Marrying IR with TM

- Classifying
- Clustering
- Summarisation
- Term extraction
- Related documents based on analysis of returned results
- More detail: talk by James Thomas
- Recommendation systems





Opinion, sentiment mining

- Opinion polling
 - Ratios for/against a proposition
- What ± passions are aroused about
 - A topic?
 - A social group?
- What are the terms in which the debate is conducted?
- How are media used in the formation and dissemination of opinion?





Case study: Alternative vote debate

- Named entity recognition
 - Finding mentions of political parties, politicians
 - Including alternative designations
- Term and topic discovery
 - Most significant terms used in a (sub) collection of text
- Subjectivity analysis
 - Positive or negative orientation, whether subjective or objective style
- Speech act and rhetorical analysis
 - Whether initiating, reacting, supporting, arguing, citing evidence, etc.
- Semantic search engines based on all the above





NaCTeM analysis of tweets on AV: overall (document-level) sentiment

Analysis Tools	
 ♀ T Ⅰ 1-15 of 315 	
	Sentiment
(DocID:34) File name: 64117217322741760 @GrumpyOldYouth @NickClegg_UK He reminded me slightly of Jim Trott from Vicar of Dibley with the No No No NoYes To AV!	
(DocID:35) File name: 64126672559022080 Yes to AV. Nothing changes if you keep voting the same people in~! say Yes to AV http://www.globalresearch.ca/index.php?context=va&aid=24538	•
(DocID:36) File name: 64131138117050368 Vote Yes to AV. Politicos think we can't do 1,2,3 (and maybe 4). Prove the patronising ******** wrong.	1
(DocID:37) File name: 64213752714760192 @Gaijinsan21 I agree. And am "Yes to AV" (and Labour). FPP is outdated nonsense. DC's stance has been disingenuous throughoutcontd	
(DocID:38) File name: 64215670069530624 @Gaijinsan21 let's hope yes to AV wins and is just a stepping stone to a better future. You have to have hope else you become bitter :)	-

Sentiment analysis of 1 document

File	Annotations	Terms	Rules	Gazetteer	Search	My Account	
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Θ	subjectivity						
	negative						
	strong						
	weak						

Sentiment: challenges

- Sentiment, subjectivity, emotion
 - Polarity of subjective expression can be confused with being *pro* or *anti* a given proposition ("I support AV" is not subjective)
- Irony
- Rhetoric (last element often overturns predecessors)
- Quality and cultural specificity of resources (e.g. sentiment lexicon)





Opinion/sentiment mining tutorial

http://www.cs.uic.edu/~liub/FBS/Sentiment-Analysis-tutorial-AAAI-2011.pdf

• Recent comprehensive tutorial by Bing Liu at AAAI 2011, August.





Conclusion

- TM has many aspects, components, levels
 - Different combinations for different tasks
 - Can be applied in any domain (customisation)
- No "one size fits all"
- Domain experts and developers must engage to ensure appropriate application for needs
- Copyright & licensing: support adoption of Hargreaves recommendations!





Acknowledgements

- Text mining team at NaCTeM (~19)
- 東京大学
- Funders and sponsors





EPSRC Engineering and Physical Sciences Research Council

Working together for a healthier world"



Wellcome Trust + 17 other sponsors (UKPMC)



Elsevier



The University of Manchester