Conference Speaker Biography Analysis

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Q: Do the self-presentation of male and female researchers differ? In what ways?

Basic Methodology

- **Data:** Biographies of speakers in computer science conferences
 - A concise self-introduction to be read by fellow academics
 - Should contain information the speaker considers to be important about their career
 - No unified format across conferences

Metrics:

- Cosine similarity: simple way to quantify difference between texts
- Term frequency: what information gets emphasized over and over again?

Basic Methodology

- Most research into gender representation in computer science focuses on the male-to-female ratio, and the larger societal trends that causes less women entering the field
 - Many compiled statistical reports are also available from government organizations (Bureau of Labor etc...)
- Most research are also focused on the computer science workforce instead of academia
- Much less research on how women already in computer science view their roles
- <u>"Women in Technology"</u> report: highlights the unique barriers facing women working in the technology sector
 - O Do the same barriers exist in academia?
 - Does this affect how women researchers presents themselves and how they are presented by others?

Data Collection



▶ HOME

▶ PROGRAM

▶ GENERAL INFORMATION

RECOMB/ISCB Conference on Regulatory and Systems Genomics, with DREAM Challenges

KEYNOTE SPEAKER ABSTRACTS & BIOGRAPHIES

updated Oct 27, 2013

DREAM Challenges



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Turning Networks Into Ontologies: Towards A Data-driven Gene Ontology Print

Abstract: Ontologies have been very useful for capturing knowledge as a hierarchy of concepts and their interrelationships. In biology, a prime challenge has been to develop ontologies of gene function given only partial biological knowledge and inconsistency in how this knowledge is curated by experts. I will discuss how large networks of gene and protein interaction, as are being mapped systematically for many species, can be transformed to assemble an ontology with equivalent coverage and power to the manually-curated Gene Ontology (GO). Our network-extracted ontology contains 4,123 biological concepts and 5,768 relations, capturing the majority of known cellular components as well as many additional concepts, triggering subsequent updates to GO. Using genetic interaction profiling we provide further support for novel concepts related to protein trafficking, including a link between Nn/2 and YEL043W. This work enables a shift from using ontologies to evaluate data to using data to construct and evaluate ontologies.

Biography: Trey Ideker, Ph. D. Is Professor of Medicine at the University of California at San Diego. He serves as Division Chief of Medical Genetics and Director of the National Resource for Network Biology, as well as being Adjunct Professor of Bioengineering and Computer Science and Member of the Moores UCSD Cancer Center, Ideker received Bachelor's and Master's degrees from MIT in Electrical Engineering and Computer Science and his Ph.D. from the University of Washington in Molecular Biology under the supervision of Dr. Leroy Hood. He is a pioneer in assembling genome-scale measurements to construct network models of cellular processes and disease. His recent research activities include assembly of networks governing the response to DNA damage, development of the Cytoscape and NetworkBLAST software packages for biological network visualization and cross-species network comparison; and methods for Identifying network-based biomarkers in development and disease. Ideker serves on the Editorial Boards for Bioinformatics and PLoS Computational Biology, is on the Scientific Advisory Boards of the Sanford-Burnham Medical Research Institute and the Institute for Systems Biology, and is a regular consultant for companies such as Monastin and Mandel Biotechnology.

invited speakers: 5

Invited speakers: 19

Collects biography for each scientist from the conference websites.

Data Collection

 Extract all the significant information out of each conference such as biography,gender,publish year,Name of speaker,

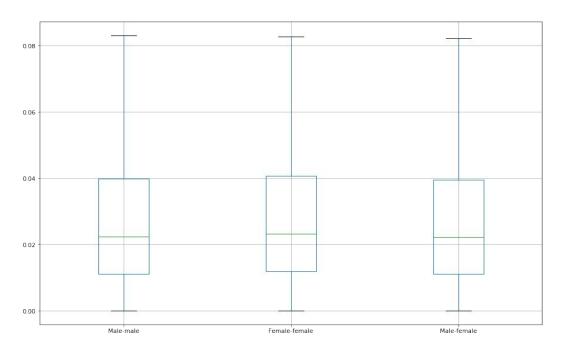
	A	В	С	D
1	Bio	Gender =	Year =	Name =
2	Kathryn S. McKinley is a Principal Research	F	2016	Kathryn S McKinley
3	Dr Larry Persons PhD is on the Faculty at S	M	2018	Larry Persons
4	Dr Arthur Shelley is an independent educate	M	2018	Arthur Shelley
5	Shane McCarthy Shane is the CEO of Blue	M	2018	Shane McCarthy
6	Stephen O'Leary is managing director at Oly	M	2018	Stephen O'Leary
7	Olav Lysne is a director of Simula Metropolit	M	2018	Olav Lysne
8	Wallace Chigona is a Professor in Information	М	2018	Wallace Chigona
9	Dr. Johannes Cronié is the Dean of Informati	M	2018	Dr. Johannes Cronié

Preliminary Results in June

- Calculated cosine similarity on a small data set of 94 biographies (74 male, 20 female), with primitive preprocessing (small list of stop words, no stemming)
- Average cosine similarity of male-male pairs: 0.0331
- Average cosine similarity of female-female pairs: 0.0588
 - High similarity due to small sample size?
- Average cosine similarity of male-female pairs: 0.0255
- There does seem to be a slight difference between male and female biographies

Result: Cosine Similarity

- Cosine similarity calculated over a set of 191 bios (153 male, 38 female)
- Improved preprocessing via NLTK



Result: Cosine Similarity

- Average cosine similarity of male-male pairs: 0.031
- Average cosine similarity of female-female pairs: 0.032
- Average cosine similarity of male-female pairs: 0.030
- Similar distribution
- No significant difference!

Result: Term Frequency

Frequency of word stems in male and female bios

Male

comput	353
research	297
univers	280
scienc	242
professor	130
system	114
award	110
algorithm	98
work	93
includ	89

research	84
comput	71
scienc	66
univers	64
award	44
professor	42
engin	27
receiv	25
learn	24
data	23

Result: Term Frequency

- Term frequency separated by part-of-speech tags (noun, adjective, verb, adverb) showed similar results: significant overlap of most common terms across genders.
- Calculated without removing stop words and stemming to ensure accuracy of tagger

Result: Document Frequency

Frequency of word stems by documents in male and female bios

Male

research	121
univers	115
scienc	112
comput	109
professor	93
includ	65
work	60
receiv	58
institut	55
award	54

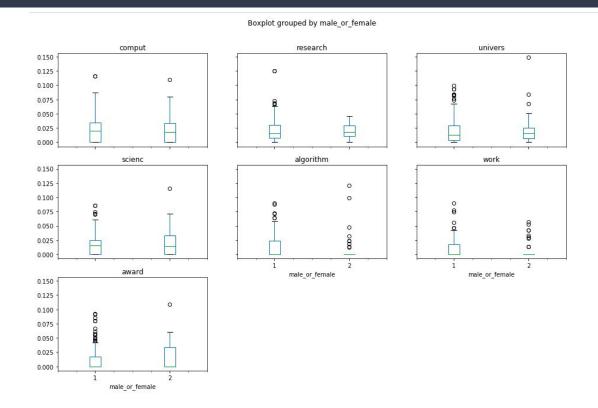
Female

research	33
univers	30
professor	28
comput	27
scienc	25
receiv	19
associ	16
engin	16
fellow	15
interest	14

Result: Document Frequency

- Document frequency separated by part-of-speech tags (noun, adjective, verb, adverb) showed similar results: significant overlap of most common terms across genders
- Again, calculated with minimal preprocessing

Result: TFIDF



Result: Mutual Information

> Top pairs of terms representing Male class

Term 1	Term 2
'aachen'	'cross'
'aachen'	'envoy'
'aberration'	'array'
'ababa'	'age'
'abstract'	'deeper'
'abstract'	'pure'
'academics'	'brothers'
'accenture'	'watch'
'adapted',	'serious'
address'	'bacterial'

> Top pairs of terms representing Female class

Term 1	Term 2
'ability',	'alzheimers'
'ability'	'anita',
'ability'	'disease'
'about'	'failure'
'about'	'smart'
'accounting'	'assist'
'accounting'	'algorithm'
'activities'	'it'
'activities',	'leadership'

Conclusion: No significant difference exists in the language used by male and female computer scientists in their conference biographies.

Final Thoughts

- Related research shows large gender differences in the tech industry
 - Academic advancement (based on blind peer reviews) more objective than metrics for climbing the "corporate ladder"?
- More significant differences across conferences than across gender?
 - Individual "conference traditions", since no unified format across conferences exist?
- Does the result hold for older conferences?
 - Would the presumably larger gender gap make this harder to calculate?
- The importance of having a large enough data set
- Format standards (or lack thereof) sometimes have a larger influence than the data they carry!
- Different preprocessing procedures for different calculations

