MOLIERE

Automatic Biomedical Hypothesis Generation System

<u>Justin Sybrandt</u>

Clemson University
School of Computing

Michael Shtutman

University of South Carolina
College of Pharmacy

Ilya Safro

Clemson University

School of Computing

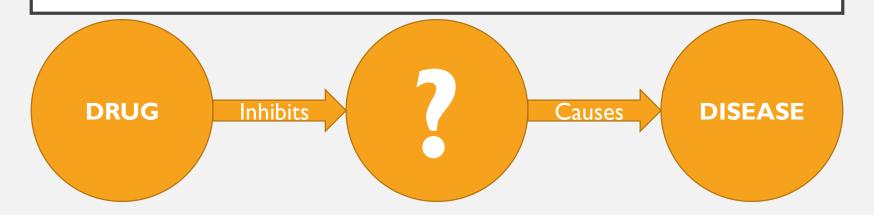
UNDISCOVERED PUBLIC KNOWLEDGE

- Proposed by Swanson in 1986
- The set of public knowledge is too large
- Contains implicit connections

HUMAN LIMITS

- No person can read everything
 - 2,000 4,000 new papers *daily*
- People Specialize
 - Limits knowledge sharing across disciplines
- Bias and Inconsistent Assumptions

WHAT IS A HYPOTHESIS?

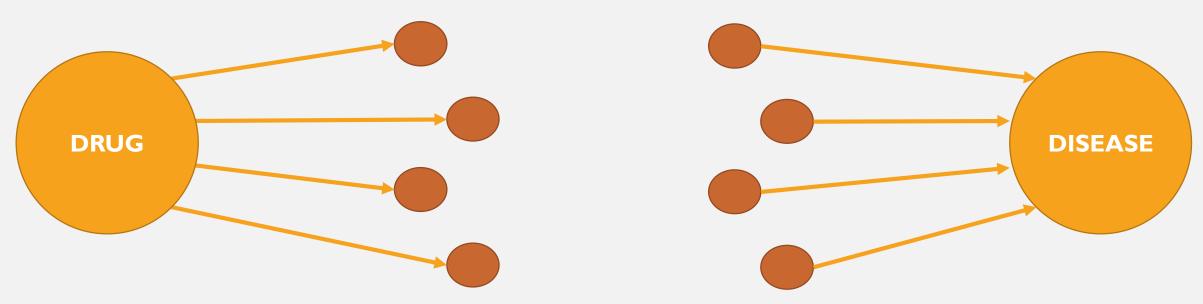


- An idea or explanation for something that is based on known facts but has not yet been proven
 - (Definition of "hypothesis" from the Cambridge Academic Content Dictionary © Cambridge University Press)

AUTOMATIC HYPOTHESIS GENERATION

Network of Medical Objects

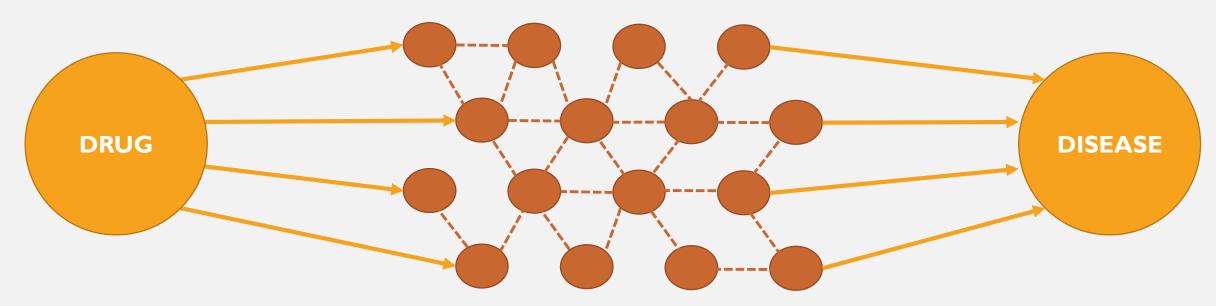
Connections Not Well Studied



AUTOMATIC HYPOTHESIS GENERATION

Network of Medical Objects

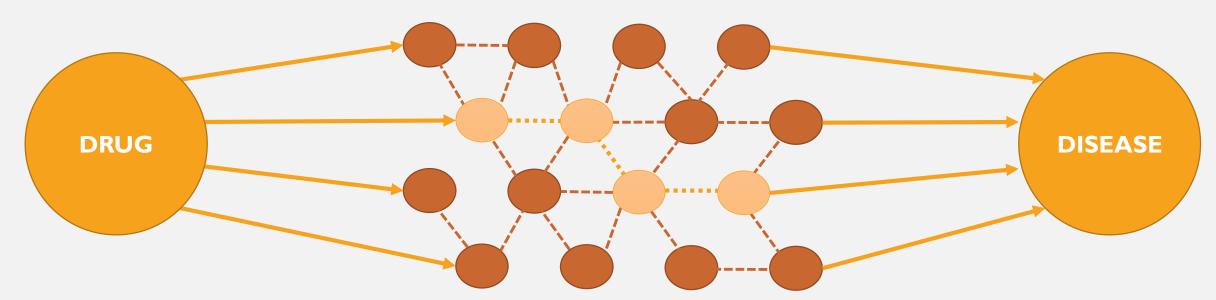
Connections Not Well Studied



AUTOMATIC HYPOTHESIS GENERATION

Network of Medical Objects

Connections Not Well Studied



RELATED APPROACHES

Methodology	Highlights		
ARROWSMITH	 One of the first hypothesis generation systems. Found link between Fish Oil and Raynaud's Disease. 		
DiseaseConnect	 Finds connections between genes and diseases. Displays information in an interactive prompt. 		
BioLDA	 Constructs high quality topic models aided by domain-specific information. Identified link between Venlafaxine and HTRIA. 		

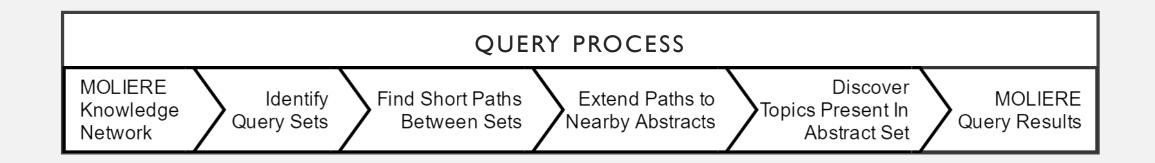
RELATED APPROACHES

Methodology	Limitation	Reference
ARROWSMITH	Limited Document Set	Neil R Smalheiser and Don R Swanson. 1998.
DiseaseConnect	Limited Document Set	Chun-Chi Liu et al. 2014.
BioLDA	Limited Vocabulary	Huijun Wang et al. 2011.

NEW APPROACH

- Construct a large network
- Identify meaningful paths
- Extend paths to neighboring nodes
- Mine neighborhoods for important topics







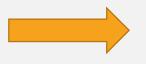
- National Library of Medicine (NLM)
- National Center for Biotechnology Information (NCBI)
- MEDLINE
 - 25 Million Documents
 - Titles and Abstracts

NETWORK CONSTRUCTION MOLIERE Raw Project Construct Text Discover Integrate Fit Centroids Phrases Data Network of Knowledge Cleaning Phrases Other Data to Abstracts Files to Vectors Abstracts Network

- SPECALIST NLP TOOLSET
- Natural Language ToolKit (NLTK)

Raw Data

This is some example text that is hopefully more understandable than the typical medical abstract.



Cleaned Text

some example text that hopefully more understandable than typical medical abstract

NETWORK CONSTRUCTION MOLIERE Raw Project Construct Text Discover Integrate Fit Centroids Phrases Knowledge Data Network of Cleaning Phrases Other Data to Abstracts to Vectors Files Abstracts Network

- Topical Pattern Mining
 - Groups together common phrases
 - Creates 2,3,...,n-grams

Cleaned Text

some example text that hopefully more understandable than typical medical abstract

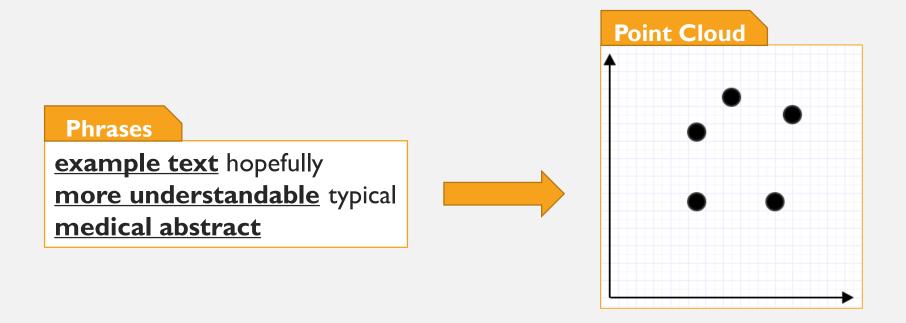


Phrases

example text hopefully
more understandable typical
medical abstract

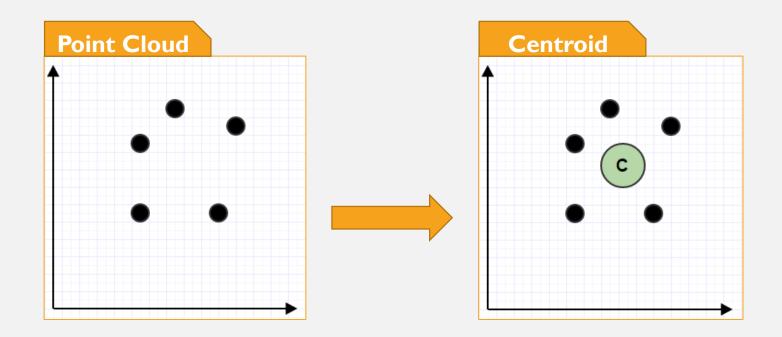
NETWORK CONSTRUCTION Project MOLIERE Raw Construct Text Discover Integrate Fit Centroids Phrases Knowledge Data Network of Cleaning Phrases Other Data to Abstracts Files to Vectors Abstracts Network

- FastText: Projects phrases into real valued vector space
- Long word embedding composed from subwords



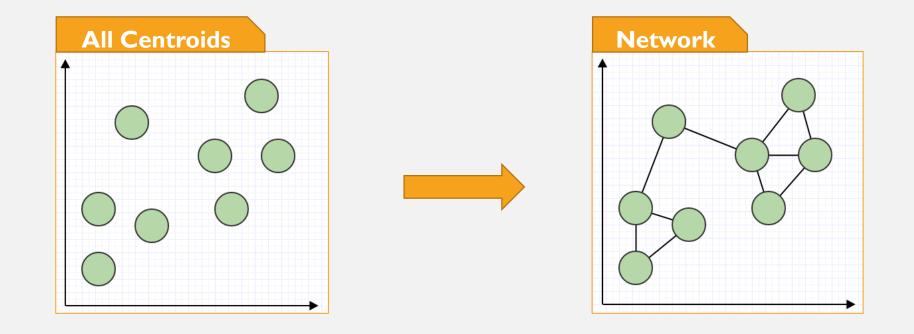


Embed documents by averaging over point clouds



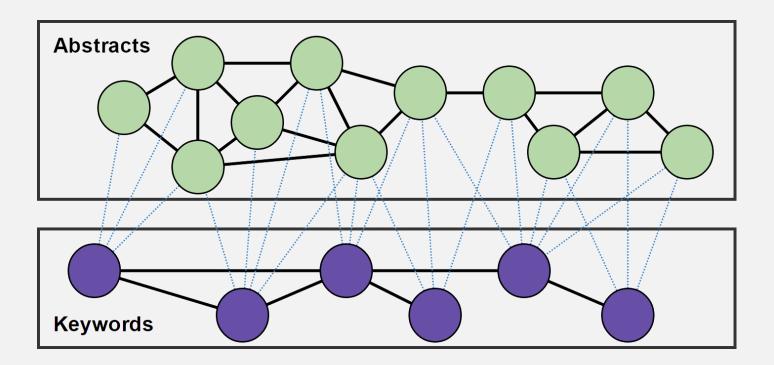


- Construct KNN
 - Fast Library for Approximate Nearest Neighbors

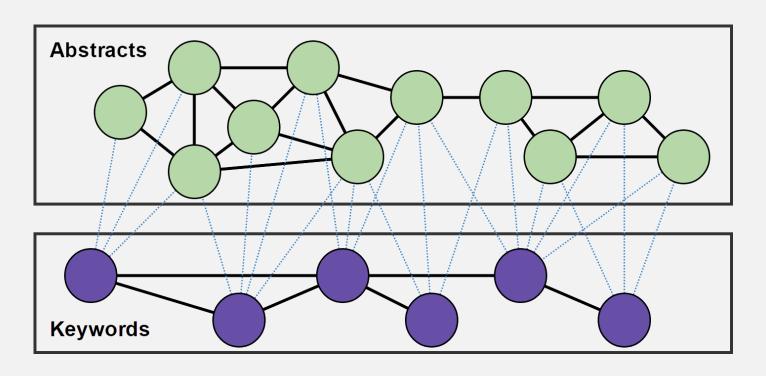




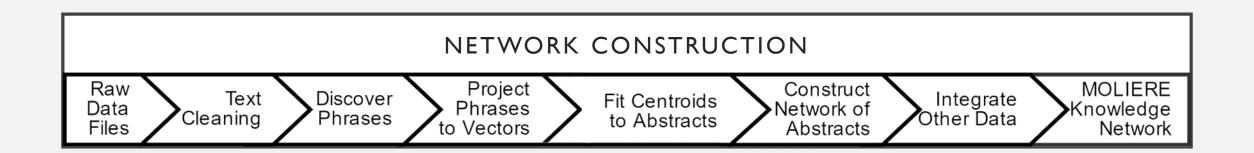
- UMLS Metathesarus
 - Curated keyword network
 - 2 Million Nodes
 - Superset of MESH

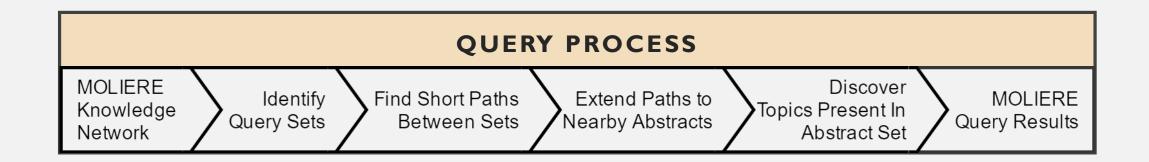






- Edge weight ~ Distance
- Inv.TF-IDF cross-layer edges
- Edges normalized [0,1]





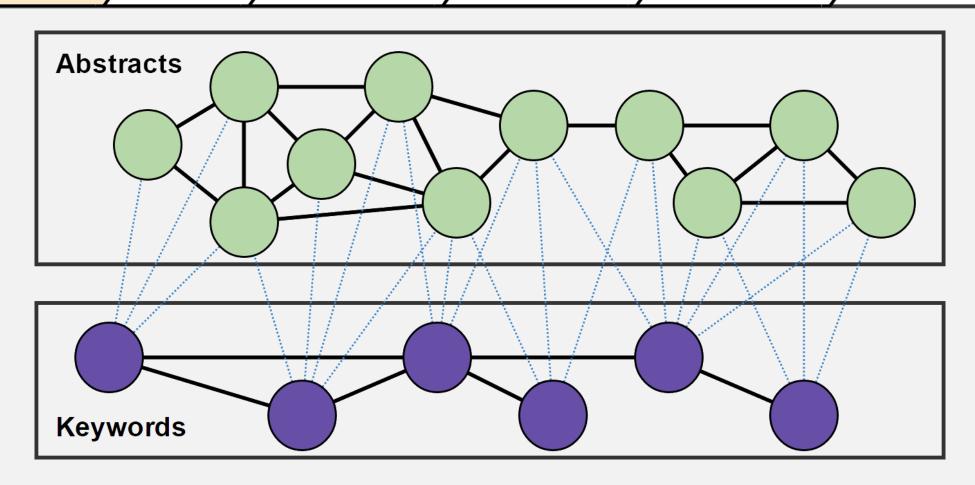
QUERY PROCESS

MOLIERE Knowledge Network

Identify Query Sets Find Short Paths Between Sets Extend Paths to Nearby Abstracts

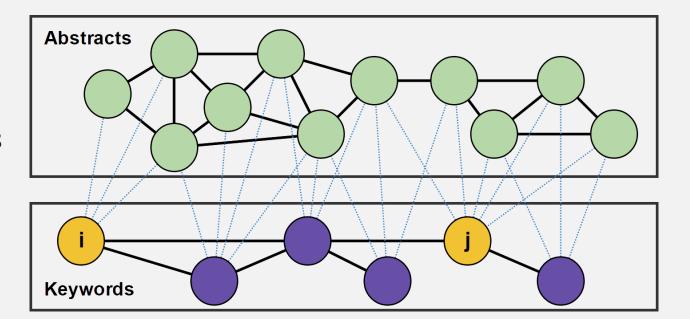
Discover Topics Present In Abstract Set

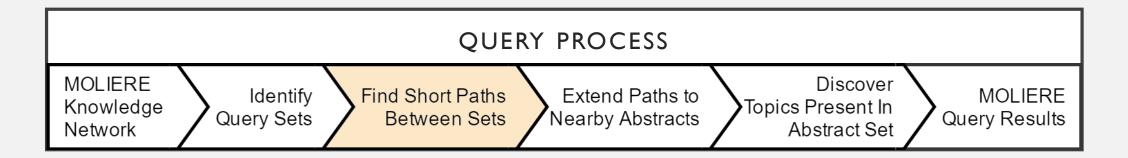
MOLIERE Query Results



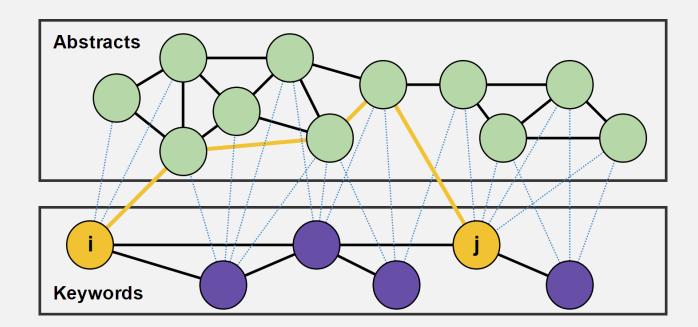
MOLIERE Knowledge Network MOLIERE Knowledge Network MOLIERE Knowledge Network MOLIERE Nearby Abstracts Extend Paths to Nearby Abstracts Discover Topics Present In Abstract Set MOLIERE Query Results

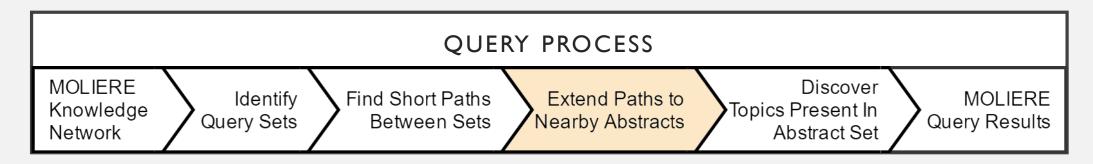
- User selects two nodes
 - Restrained to keywords
- Can generalize to two sets



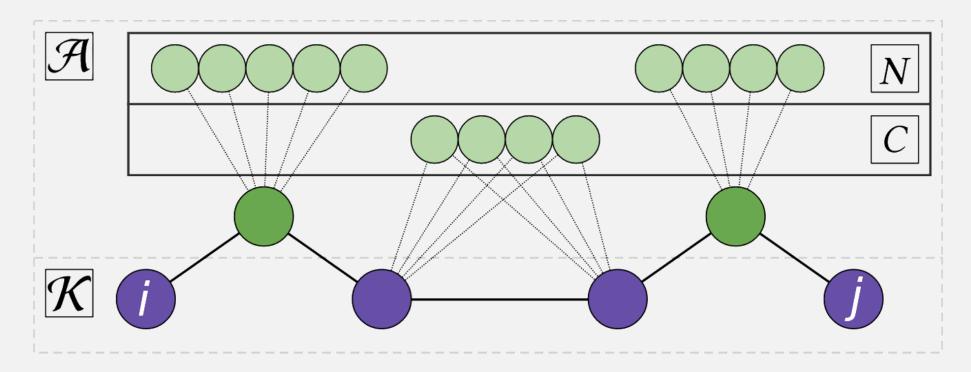


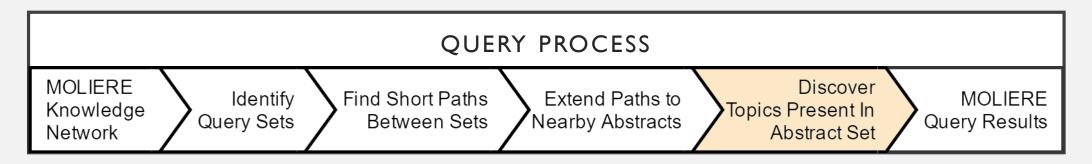
 Identify shortest path between query sets



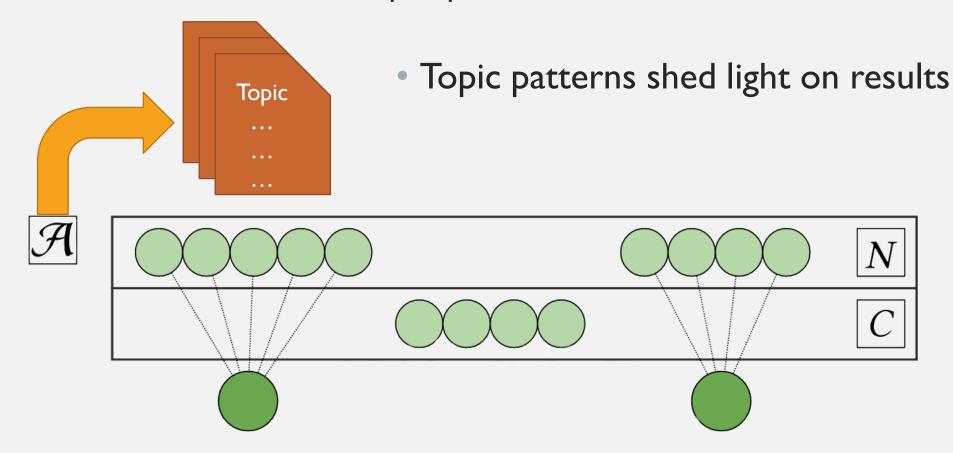


- N: Abstracts close to those in original path
- C: Abstracts which share path-adjacent keywords





• PLDA+: Identifies topics present in a set of text.



MOLIERE Knowledge Network MOLIERE Knowledge Network MOLIERE Knowledge Network MOLIERE Network Extend Paths to Nearby Abstracts Topics Present In Abstract Set MOLIERE Query Results

- Hypothesis represented as a topic model
- Shown: Venlaflaxine vs. HTRIA

TOPIC: 0 antidepressant_drugs milnacipran org selected ht

TOPIC: I increase reduced treatment dorsal_raphe_nucleus effect

rats sert ht_receptor escitalopram potency

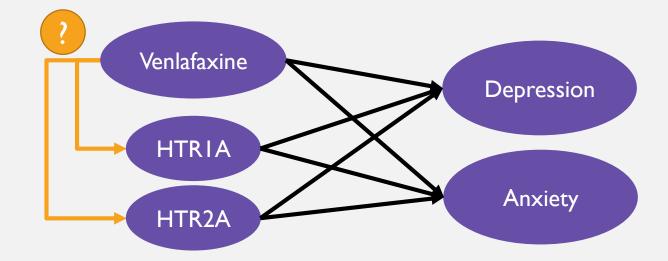
RESULTS

VENLAFAXINE

DRUG REPURPOSING

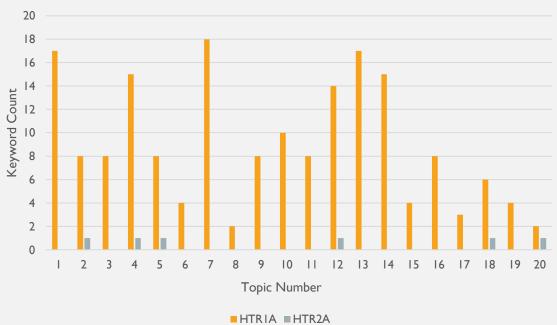
VENLAFAXINE EXAMPLE

- <u>Venlafaxine</u>:
 - Treats depression / anxiety
- <u>HTR[12]A</u>:
 - Linked to depression / anxiety
- No paper linked these concepts

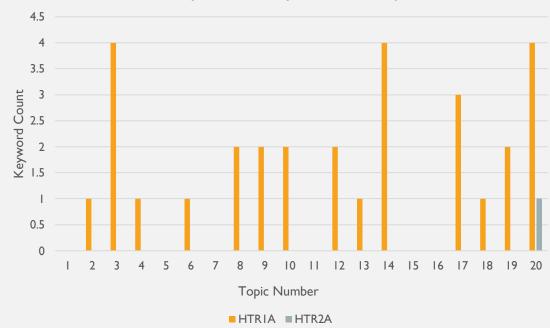


VENLAFAXINE RESULTS



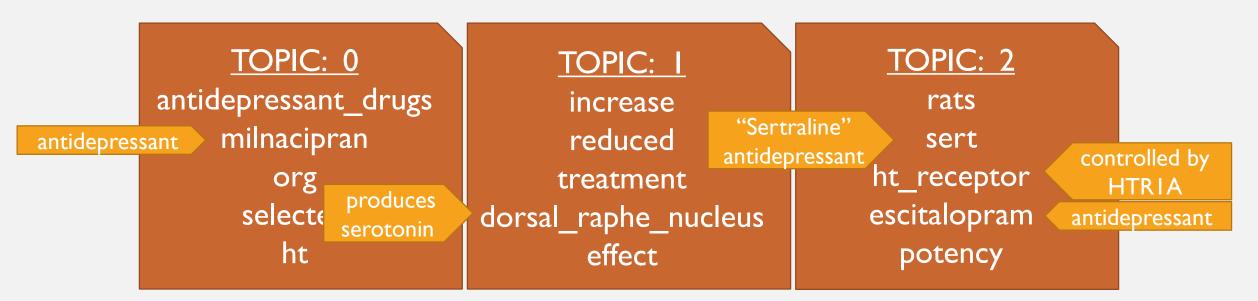


Anxiety Related Keywords Per Topic



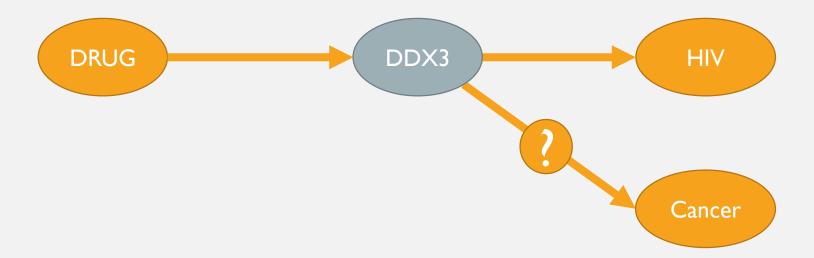
VENLAFAXINE RESULTS

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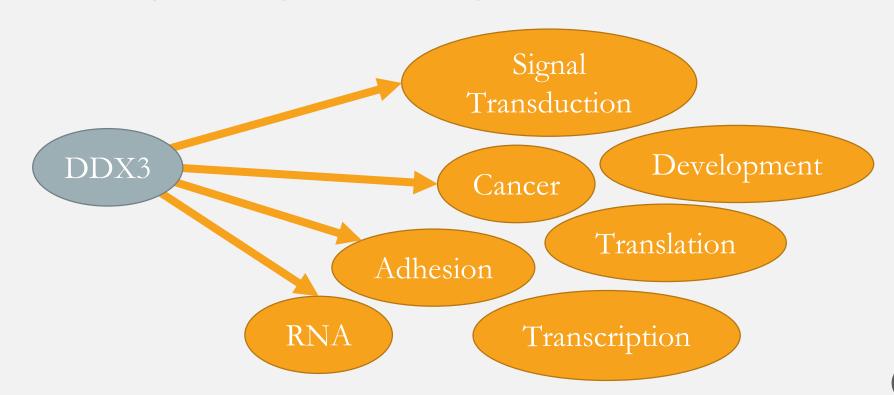
DRUG REPURPOSING EXAMPLE

- Drugs can be modified to treat new diseases
- Decreases drug development time and costs



DRUG REPURPOSING EXPERIMENT

Ran nearly 10,000 queries involving DDX3:



DRUG REPURPOSING EXPERIMENT

- Ran nearly 10,000 queries involving DDX3:
- Expecting:

Cell – Cell Adhesion WNT Signaling
Pathways

Cell – Matrix Adhesion

DRUG REPURPOSING RESULTS

cell-cell adhesion Cell – Cell regulation of cell-cell adhesion Adhesion cell-adhesion molecules signal-transduction associated kinases WNT Signaling cell adhesion kinase Pathways substrate adhesion Cell – Matrix RGD cell adhesion domain Adhesion cell adhesion factor focal adhesion kinase

APPLICATIONS

- Drug Repurposing
- Extensions to new domains
 - Patents, Economics, etc.
- Coping with Deadlines



OPEN RESEARCH QUESTIONS

- Result Interpretation
- System Verification
- Automatic Network Tuning
- Streaming Network Reconstruction
- Inclusion of Additional Data Sources

THANK YOU

J. Sybrandt, M. Shtutman, I. Safro "MOLIERE: Automatic Biomedical Hypothesis Generation System"

Code and Data: https://people.cs.clemson.edu/~isafro/software.html

Email: JSYBRAN@CLEMSON.EDU