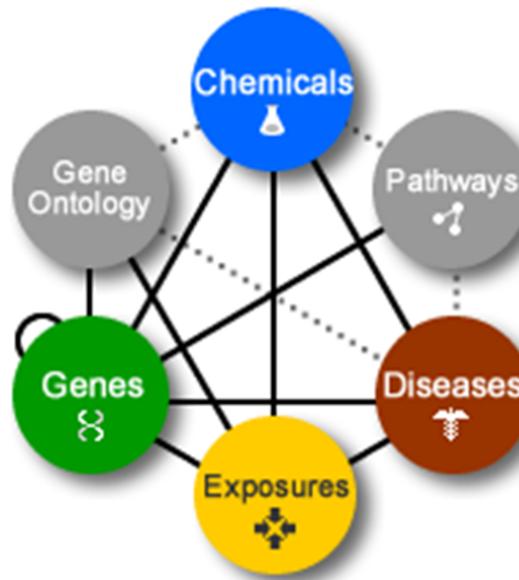


# BioCreative V Track 3: Chemical-Disease Relation and Disease Named Entity Recognition and Normalization



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*Earl Stadtman Investigator*

*Head, Biomedical Text Mining Group*

*National Center for Biotechnology Information*

**Thomas C. Wiegers**

*Research Bioinformatician*

*Comparative Toxicogenomics Database*

*North Carolina State University*

# Agenda

## Introduction and Background

- Introduction to CDR/DNER-based curation
- Explanation of why we chose these particular tasks:
  - Why is CDR/DNER text mining so important?
- Case study of how CDR/DNER curation is used

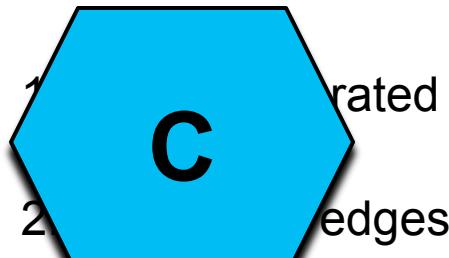
## Task Design

- DNER task examples
- CDR task examples
- Corpora descriptions
- Supporting resources
- Task evaluation

## Results

- Challenge results
- Survey results

# CTD is a Curated Knowledgebase



3. Controlled vocabularies

4. Import external attributes

MeSH



Chemical-Disease  
Relationships

chemical-gene  
interactions

chemical-disease  
relationships



Entrez Gene

D

D

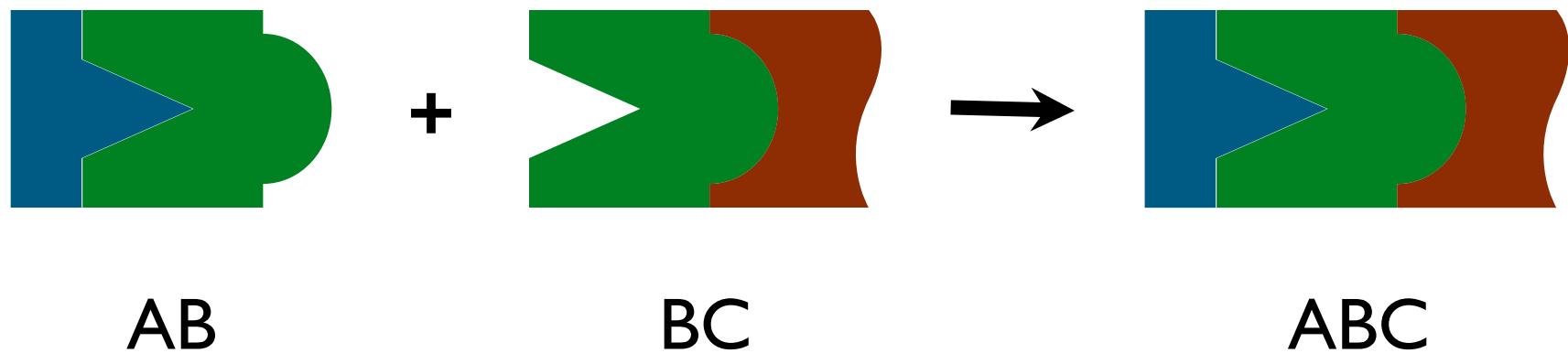
gene-disease  
relationships

# Chemical/Disease Relationships (CDRs)?

- **Marker:** A chemical that correlates with a disease (e.g., increased abundance in the brain of chemical X correlates with Alzheimer disease) or may play a role in the etiology of a disease (e.g., exposure to chemical X may play a role in causing lung cancer).
- **Therapeutic:** A chemical that has a known or potential therapeutic role in a disease (e.g., chemical X is used to treat leukemia).

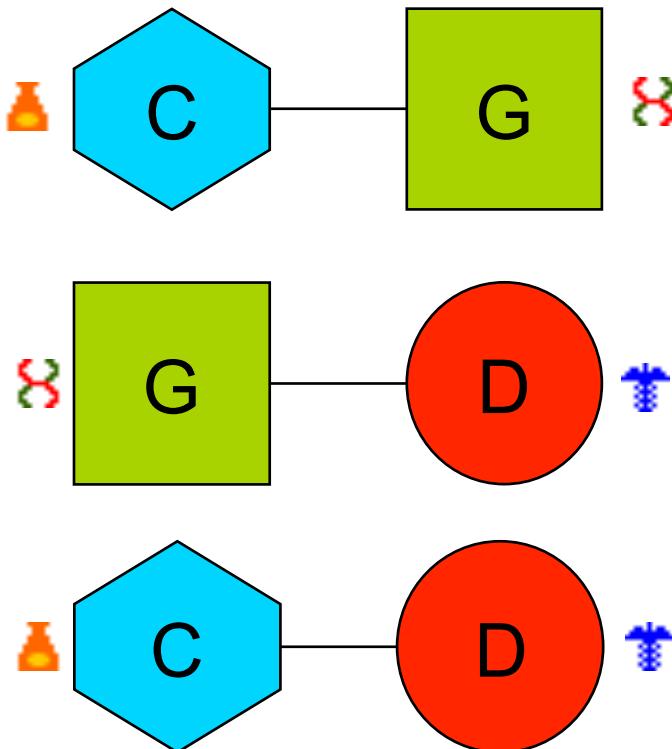
Why are these relationships important?

# Transitive Inference ABC Model of Hypothesis Generation



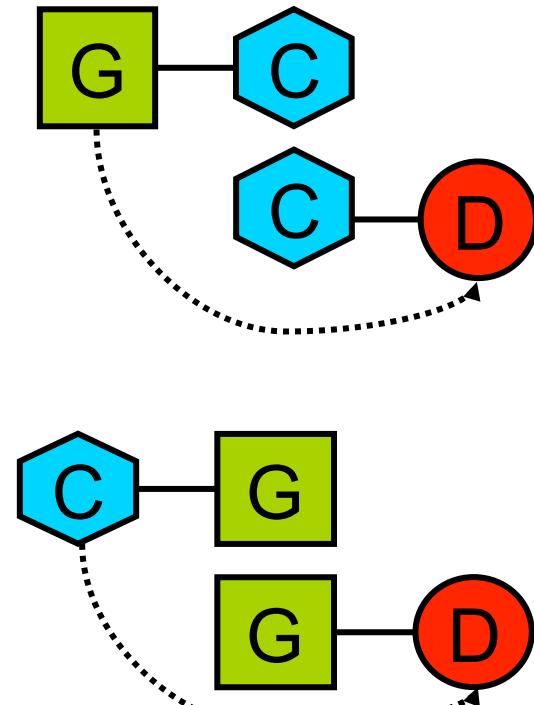
# Transitive Inference at CTD

Curated Facts



Inferred Discoveries

Integrate



CTD: Over 27 million  
Toxicogenomic Relationships



# Case Study: CTD/Pfizer Collaboration



PubMed.gov US National Library of Medicine National Institutes of Health

PubMed Advanced Search

Abstract Send to:

[Database \(Oxford\)](#), 2013 Nov 28;2013:bat080. doi: 10.1093/database/bat080. Print 2013.

**A CTD-Pfizer collaboration: manual curation of 88,000 scientific articles text mined for drug-disease and drug-phenotype interactions.**

Davis AP<sup>1</sup>, Wiegers TC, Roberts PM, King BL, Lay JM, Lennon-Hopkins K, Sciaky D, Johnson R, Keating H, Greene N, Hernandez R, McConnell KJ, Enayetallah AE, Mattingly CJ.

[+ Author information](#)

**Abstract**  
Improving the prediction of chemical toxicity is a goal common to both environmental health research and pharmaceutical drug development. To improve safety detection assays, it is critical to have a reference set of molecules with well-defined toxicity annotations for training and validation purposes. Here, we describe a collaboration between safety researchers at Pfizer and the research team at the Comparative Toxicogenomics Database (CTD) to text mine and manually review a collection of 88,629 articles relating over 1,200 pharmaceutical drugs to their potential involvement in cardiovascular, neurological, renal and hepatic toxicity. In 1 year, CTD biocurators curated 254,173 toxicogenomic interactions (152,173 chemical-disease, 58,572 chemical-gene, 5,345 gene-disease and 38,083 phenotype interactions). All chemical-gene-disease interactions are fully integrated with public CTD, and phenotype interactions can be downloaded. We describe Pfizer's text-mining process to collate the articles, and CTD's curation strategy, performance metrics, enhanced data content and new module to curate phenotype information. As well, we show how data integration can connect phenotypes to diseases. This curation can be leveraged for information about toxic endpoints important to drug safety and help develop testable hypotheses for drug-disease events. The availability of these detailed, contextualized, high-quality annotations curated from seven decades' worth of the scientific literature should help facilitate new mechanistic screening assays for pharmaceutical compound survival. This unique partnership demonstrates the importance of resource sharing and collaboration between public and private entities and underscores the complementary needs of the environmental health science and pharmaceutical communities. Database URL: <http://ctdbase.org/>

- Objective: Leverage Published Literature on Toxicity
  - Begin building comprehensive drug-event relationships database
  - Target: Cardiovascular, Neurological, Renal, and Hepatic Toxicity
- CTD Manually Curated 88,629 Articles:
  - 254,173 Toxicogenomic Interactions
  - 152,173 Chemical-Disease
  - 58,572 Chemical-Gene
  - 5,345 Gene-Disease
  - 38,083 Phenotypic

Why? Example: Investigation of Adverse Study-Related Findings

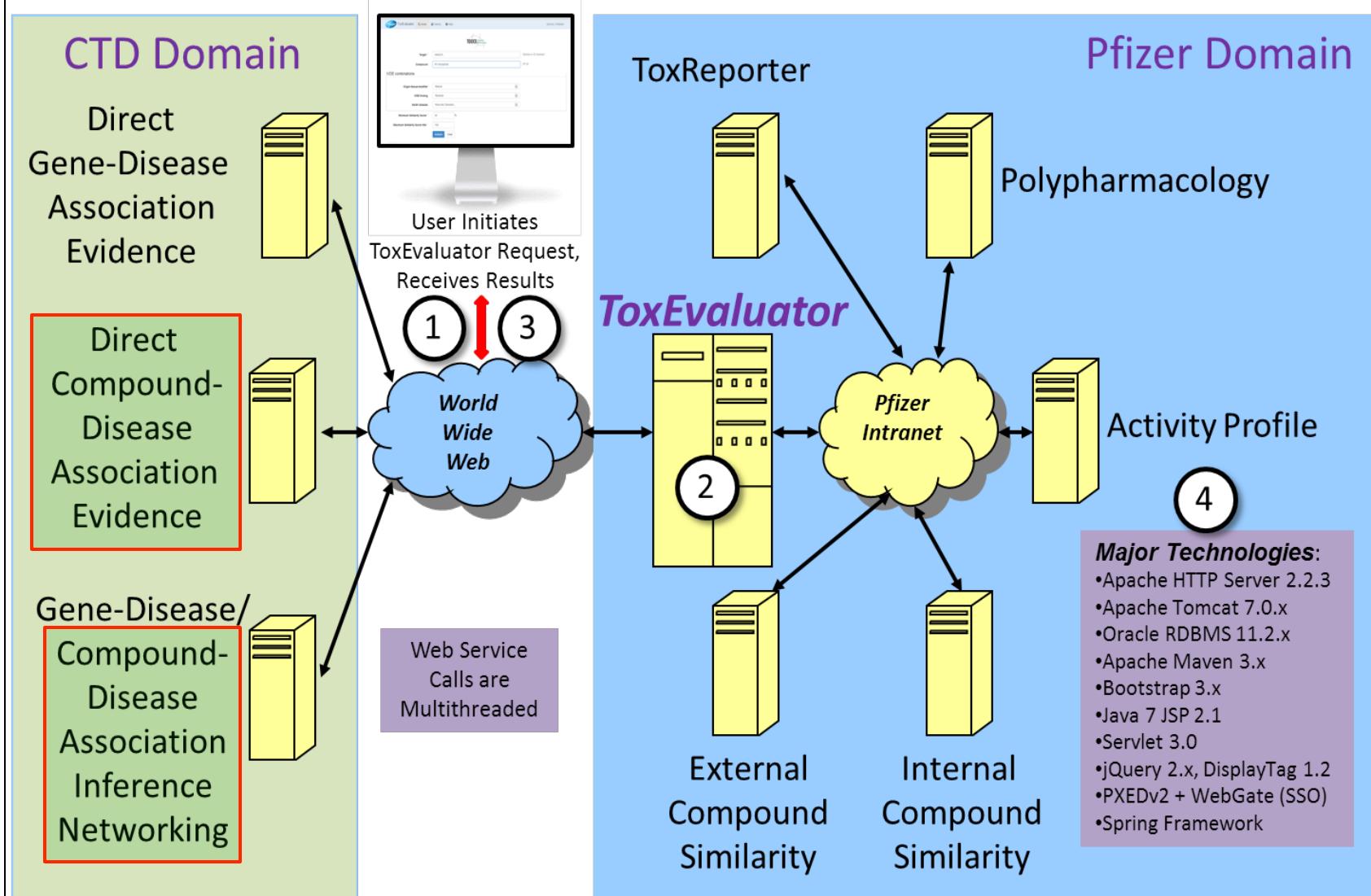
# Investigation of Adverse Findings Clinical and Pre-Clinical Studies

Pathologists and Toxicologists Investigate Adverse Findings in Animal Studies:

- Investigations Typically Begin With:
  - Chemical Structure
  - Intended Biological Target
  - Description of Observed Pathology:
    - Organ
    - Type of lesion
    - Species
- First Question: Is Lesion Related to the Study?
- If So, Is Lesion Related to:
  - The Chemical Structure Itself, i.e., the ***chemotype***
  - Pharmacological Response to Modulation of Target, i.e., ***mechanism***
  - Interaction with Related or Unrelated Biological Systems, i.e., ***off-target***
- Bottom Line: Are there Connections Between the:
  - Chemical Structure
  - Targeted Modulation
  - Observed Pathology

# Pfizer's ToxEvaluator

Real-Time, Multidimensional Querying Across Pfizer and CTD Web Services



# Pfizer's ToxEvaluator

**Pfizer ToxEvaluator** [Home](#) [History](#) [Help](#) Dennis J Pelletier

← Revise query: TARGET: HMGCR (3156) · COMPOUND: PF-00346556 · ORGAN-TISSUE-MODIFIER: Muscle · IVDE FINDING: Myolysis · MeSH DISEASE: Muscular Diseases · MAXIMUM SIMILARITY SCORE HITS: 100 · MINIMUM SIMILARITY SCORE: 50%

**Polypharmacology**

PF-00346556 was predicted to bind to **42** targets with probability **>0.7**.

Targets with **direct** Marker links to Muscular Diseases: **0 out of 42**

Targets with **inferred** links to Muscular Diseases: **42 out of 42**

**Activity Profile**

Targets showing activity for PF-00346556 **18** out of 17

Targets with **direct** Marker links to Muscular Diseases: **1 out of 18**

Targets with **inferred** links to Muscular Diseases: **17 out of 18**

**Internal Compound Similarity**

Number of compounds showing similarity to PF-00346556 > 0.5 = **0**

Similar compounds with findings of Myolysis: **0**

Similar compounds with other IVDE findings: **0**

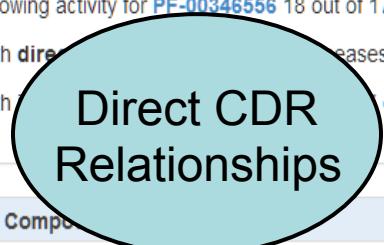
[Full Report](#)

**External Compounds**

Number of compounds showing similarity to PF-00346556 > 0.5 = **3**

Similar compounds with **direct** links to Muscular Diseases: **1 out of 3**

Similar compounds with **inferred** links to Muscular Diseases: **2 out of 3**




**ToxReporter**

Nervous\_System Summary Score: **0.7534**

No Tox Reporter Score found for Tox Term: Skeletal\_muscle; Scores returned from Tox Reporter: [Adipose:0.8322, Cardiovascular\_System:0.7532, Dermal:0.4734, Endocrine\_System:0.7403, Gastrointestinal\_System:0.7558, Hepatobiliary\_System:0.7754, Immune\_System:0.6272, Musculoskeletal\_System:0.6111, Nervous\_System:0.7534, Ocular:0.4666, Pulmonary\_System:0.6102, Renal\_System:0.5933, Reproductive\_System:0.7706, Carcinogenicity:0.5815, Inflammation:0.6453, Mitochondrial\_toxicity:0]

**CTD Gene**

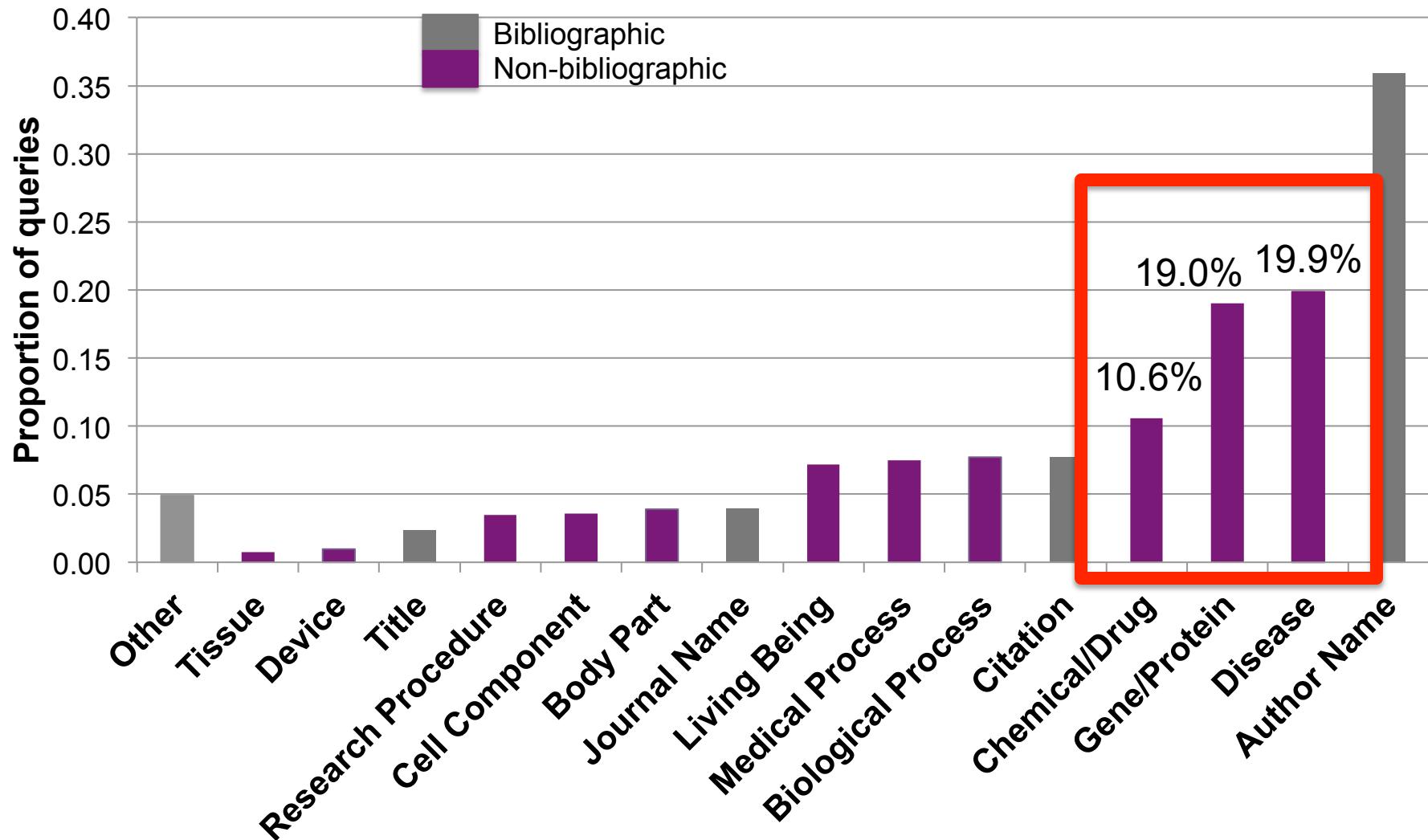
**Direct**

Evidence map: [View](#)  
References: **0**

**Inferred**

Inference Score: **78.43**  
References: **117**

# Most Searched Bio-Concepts in PubMed Queries



# Disease Named Entity Recognition (DNER) Task

Clin Oncol (R Coll Radiol). 1993;5(6):367-71.

## Liposomal daunorubicin in advanced Kaposi's sarcoma: a phase II study.

Money-Kyrle JF<sup>1</sup>, Bates F, Ready J, Gazzard BG, Phillips RH, Boag JC

 Author information

MeSH: D012514  
Sarcoma, Kaposi

### Abstract

We report a non-randomized Phase II clinical trial to assess the efficacy and safety of liposomal daunorubicin (DaunoXome) in the treatment of AIDS related Kaposi's sarcoma. Eleven homosexual men with advanced Kaposi's sarcoma were entered in the trial. Changes in size, colour and associated oedema of selected 'target' lesions were measured. Clinical, biochemical and haematological toxicities were assessed. Ten subjects were evaluated. A partial response was achieved in four, of whom two subsequently relapsed. Stabilization of Kaposi's sarcoma occurred in the remaining six, maintained until the end of the trial period in four. The drug was generally well tolerated, with few mild symptoms of toxicity. The main problem encountered was haematological toxicity, with three subjects experiencing severe neutropenia (neutrophil count < 0.5 x 10<sup>9</sup>/l). There was no evidence of cardiotoxicity. In this small patient sample, liposomal daunorubicin was an effective and well tolerated agent in the treatment of Kaposi's sarcoma.

PMID: 8305357 [PubMed - indexed for MEDLINE]

- Input: Raw PubMed Abstracts
- Output: Disease Mentions Normalized to MeSH IDs

# Relation Extraction Task

Clin Oncol (R Coll Radiol). 1993;5(6):367-71.

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PMID: 8305357 [PubMed - indexed for MEDLINE]

- Input: Raw PubMed Abstract
- Output: Chemical-Disease Pairs

Relation name	Relation type	Bio-entities
CID	Chemical_Disease	D003630(daunorubicin)   D009503(neutropenia)

# Challenges in Relation Extraction

Clin Oncol (R Coll Radiol). 1993;5(6):367-71.

## Liposomal daunorubicin in advanced Kaposi's sarcoma: a phase II study.

Money-Kyrle JF<sup>1</sup>, Bates F, Ready J, Gazzard BG, Phillips RH, Boag FC.

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PMID: 8305357 [PubMed - indexed for MEDLINE]

Not relevant;  
Therapeutic  
Relationship

Relation Across  
Sentence  
Boundaries

Negation

# Previous Work

- Related Resources
  - **Corpora**: NCBI Disease Corpus; CHEMDNER; ADE; EU-ADR
  - **Ontologies**: MeSH; OMIM; MEDIC; UMLS; Disease Ontology
  - **Tools**: DNorm; MetaMap; Peregrine
- Previous Challenges:
  - BioCreative 2012 and IV CTD Tasks (2012, 13)
  - ShARe/ CLEF eHealth Shared Task (2013- )
  - PPI, DDI

# Task Corpora

- Selected from CTD-Pfizer Set
  - Training Set : 500 Articles
  - Development Set: 500 Articles
  - Test Set : 500 Articles
- For Each Article, Annotated Data Includes:
  - Disease Named-Entity Annotations
  - Chemical Named-Entity Annotations
  - Relation Annotations Via CTD Curation
- Data Formats:
  - BioC XML
  - PubTator Text

# Chemical and Disease Annotations

- Task: Mark Up Chemicals & Diseases
  - Text spans
  - Normalized MeSH ids
  - All 1,500 articles
- Annotators
  - MeSH Indexers
  - Double Annotation performed
- Annotation Guidelines
  - What TO and NOT to Annotate
  - Largely Consistent with Previous Guidelines...
  - ...i.e., CHEMDNER and NCBI Disease Corpus
- PubTator-Assisted Annotation
  - For Improved Productivity
  - Used Pre-Annotations by DNorm & tmChem

The screenshot shows the PubTator web-based text-mining tool. At the top, there are buttons for 'Go back' (with a left arrow), 'Curatable' (radio button), 'Not Curatable' (radio button), and 'TBD' (radio button). To the right is the 'PubTator' logo. Below that is a 'Bioconcepts' section with checkboxes for 'Disease' (checked) and 'Chemical' (unchecked).

The main content area displays a medical abstract from the journal 'Chest'. The abstract discusses lidocaine-induced cardiac asystole in a 67-year-old man. Key terms like 'lidocaine', 'asystole', 'depression', 'bradycardia', and 'arrhythmias' are highlighted in green, indicating they have been annotated.

Below the abstract is a 'CTD\_GOLD' section with a table of annotations:

Entity type	Entity mention	Concept ID	Nomenclature	Delete	Evidence	Comment
Disease	asystole	D006323	MEDIC (Mention)	<input type="button" value="Delete"/>	<input type="button" value="Evidence"/>	
Disease	bradycardia	D001919	MEDIC (Mention)	<input type="button" value="Delete"/>	<input type="button" value="Evidence"/>	
Disease	depression	D019982	MEDIC (Mention)	<input type="button" value="Delete"/>	<input type="button" value="Evidence"/>	
Chemical	lidocaine	D008012	MESH (Mention)	<input type="button" value="Delete"/>	<input type="button" value="Evidence"/>	

At the bottom of the interface are buttons for 'Save Annotation Results' and 'Save & Export Annotation Results'.

# Relation Annotation Overview

- Leveraged CTD-Pfizer Curation...Except:
  - Removed Relations Involving Entities Not in Abstracts
  - Removed General Relations, e.g. Adverse Drug Event
  - Updated Some Relations Due to MeSH Update
- Performed new annotations for 100 test articles
  - Selected Using NCBI Similarity Indexing
  - Similar C/D/CID to Training and Development Sets
  - Done by CTD Curators
  - Used Normal CTD Curation Protocol
  - Not Released to Public until Testing Complete

# Annotation Results

- Corpus Overall Statistics

Data Set	# of articles	Chemical		Disease		CID relation
		Mention	ID	Mention	ID	
Training Set	500	5203	1467	4182	1946	1039
Development Set	500	5347	1507	4244	1838	1012
Test Set	500	5385	1435	4424	1988	1066

- Inter-Annotator Agreements in Jaccard Index

Data Set	IAA - Diseases	IAAs - Chemicals
Training Set	0.8600	0.9523
Development Set	0.8742	0.9577
Test Set	<b>0.8875</b>	0.9630
All Sets	0.8747	0.9605

# Task Supporting Resources

- BioC Library ([bioc.sourceforge.net](http://bioc.sourceforge.net))
- Data Visualization & Comparison in PubTator
- Freely Available NER Taggers
  - Disease: DNorm
    - 1<sup>st</sup> in ShARe/CLEF Disease Normalization Task (2013)
    - 80.9% in F1 Score
  - Chemical: tmChem
    - 1<sup>st</sup> in BioCreative IV CHEMDNER-CEM Task (2013)
    - 88.3% in F1 Score
  - Customized for CDR-task
    - Trained models with CDR data sets
    - Folder-pooling mechanism for instant response
- Evaluation Kit
- FAQs



# CDR Task Evaluation

- How? Through Web Services:
  - Representational State Transfer (RESTful) APIs
    - Client Sends Text Passages...
    - ...Using Simple HTTP Calls
    - Web Service Annotates Text Passages
    - Client Receives Automatic Mark-Ups
  - Requested Responses Within 30 Seconds
- Why Web Services?
  - Quick Access to Text-Mining Services
  - Simpler Text-Mining Pipeline Integration
  - Potential Improvement of System Scalability

# RESTful Web Service Support

## 1. RESTful Set-up Support:

- Executable programs
- Step-By-Step Instructions
- Or Create Your Own

## 2. API Testing

- Results Format Validation
- Response Time



**CDR API testing**

This webpage is for CDR API testing purposes only. Users can copy and paste their service POST url into the input box below. The web pages will show testing results.

**Input Format:**

**Task:**

**Run:**

**POST url:** (e.g., <http://104.154.76.254/cdr-ud-team/dner>)

---

**Report**

URL	<a href="http://104.154.76.254/cdr-ud-team/dner">http://104.154.76.254/cdr-ud-team/dner</a>
Status	VALID
Response time	2932 ms
Number of document	1

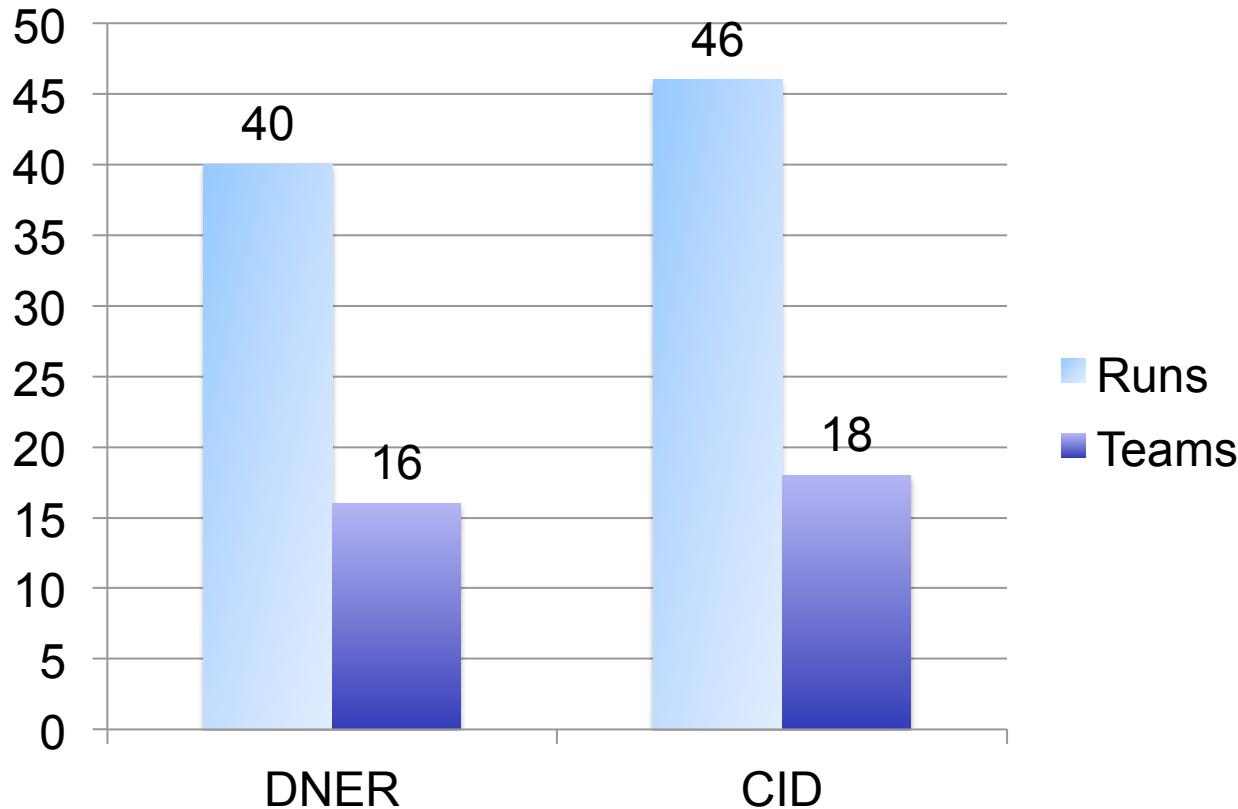
---

**Evaluation result:**

Class	gold (match)	answer (match)	recall	prec.	fscore
Disease					
Concept id matching	3 ( 0 )	0 ( 0 )	00.00	--	00.00
Mention matching	10 ( 0 )	0 ( 0 )	00.00	--	00.00

- DNER Task:
  - Dictionary Look-Up - Baseline
  - Out-Of-Box DNorm - Machine Learning
    - Used CDR Data with Default Settings
- CID Task:
  - Co-Occurrence Method
    - Used DNorm and tmChem
    - Two Versions:
      - Abstract-Level
      - Sentence-Level

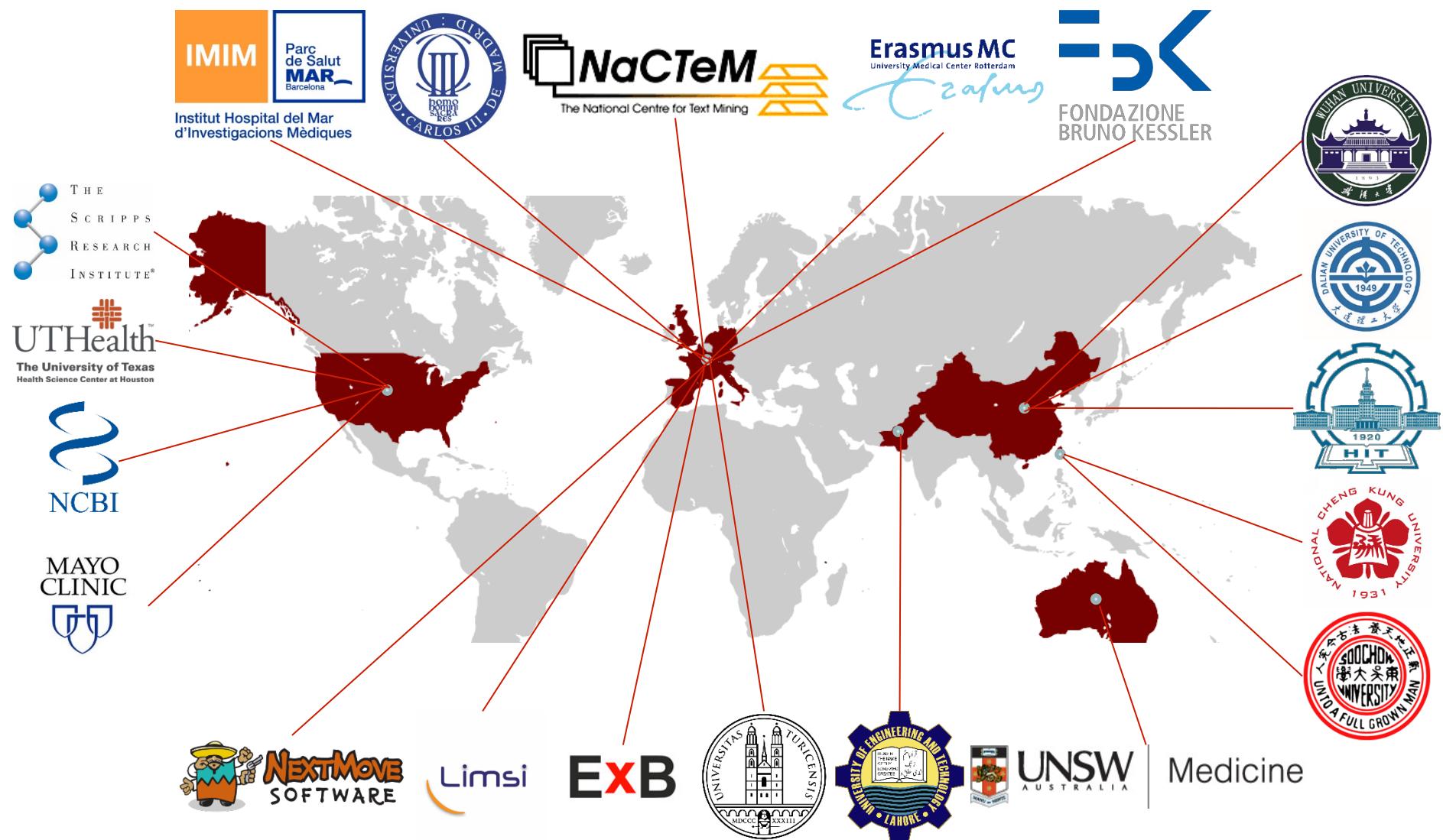
# Participating Teams Up to 3 Runs Per Team



In addition, a CID team submitted two manual runs via crowdsourcing



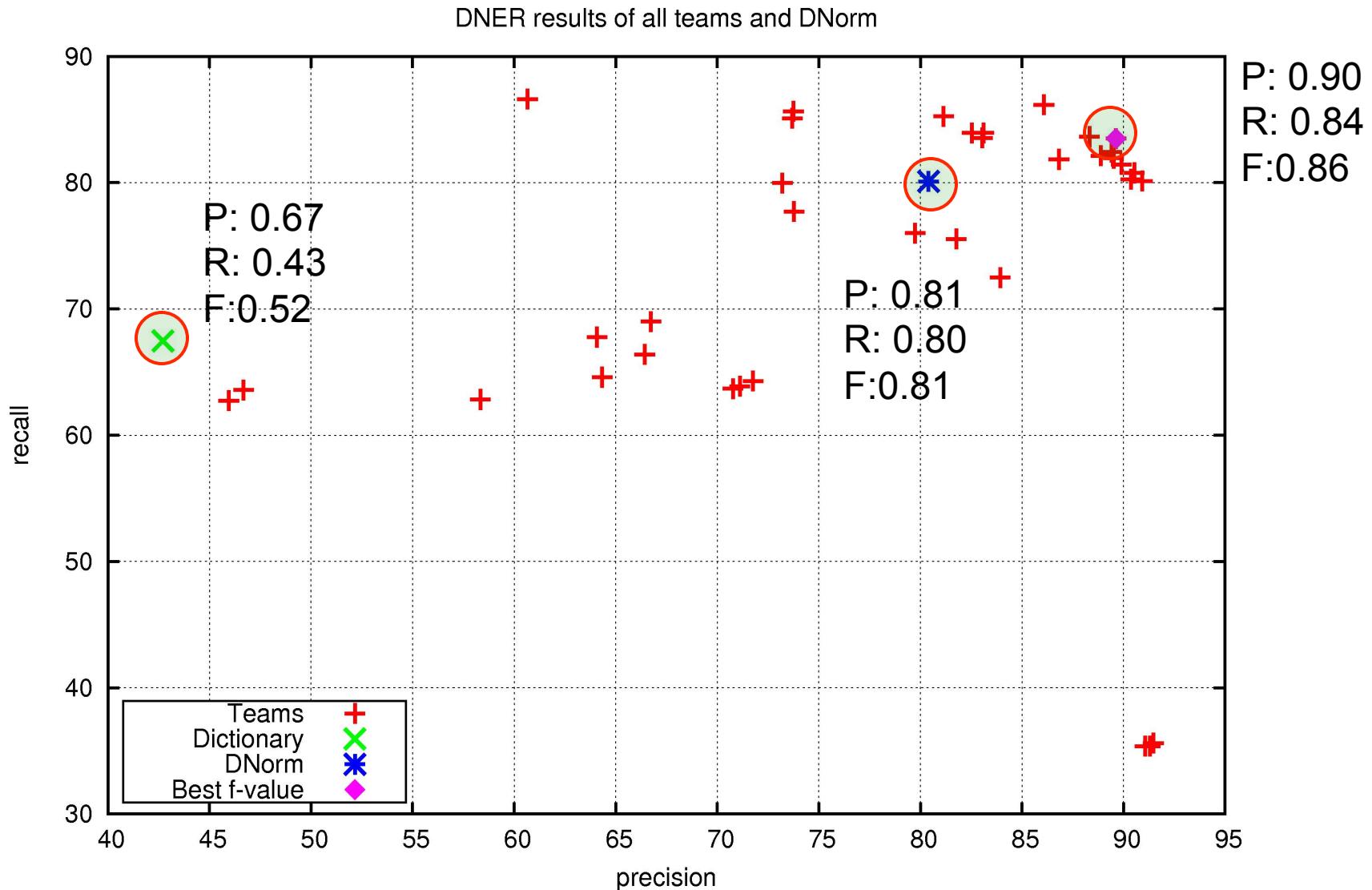
# Teams: 12 Countries/4 Continents



# Evaluation Metrics

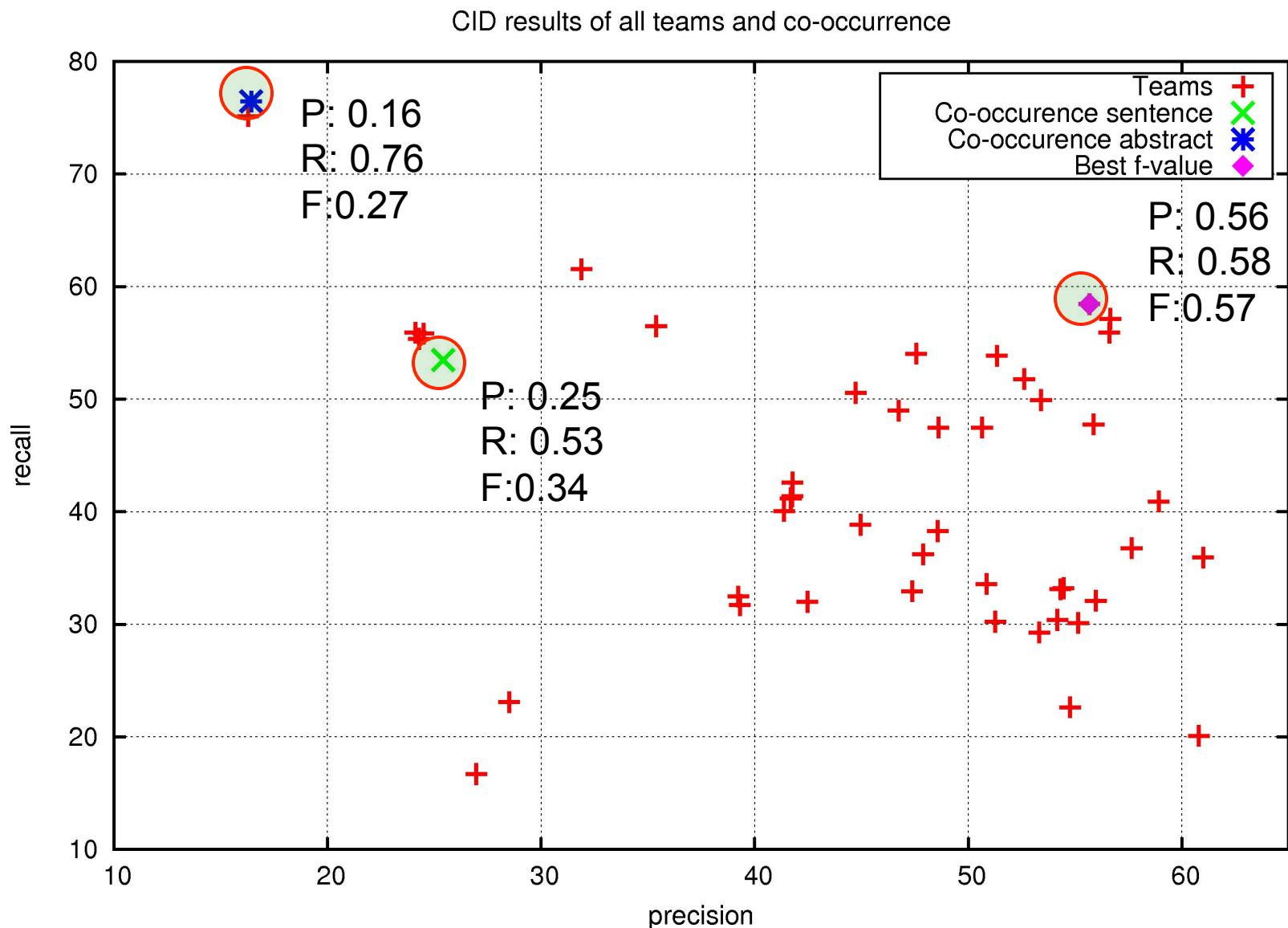
- DNER Task
  - Data Points: Document ID, Disease Concept ID
  - Precision, Recall, F-score
  - Mention-Level Accuracy Computed
- CID Task
  - Data points:
    - Document ID
    - Disease Concept ID
    - Chemical Concept ID
  - Precision, Recall, F-score
- Refer to Task Overview Paper

# DNER Results: 16 Teams, 40 Runs



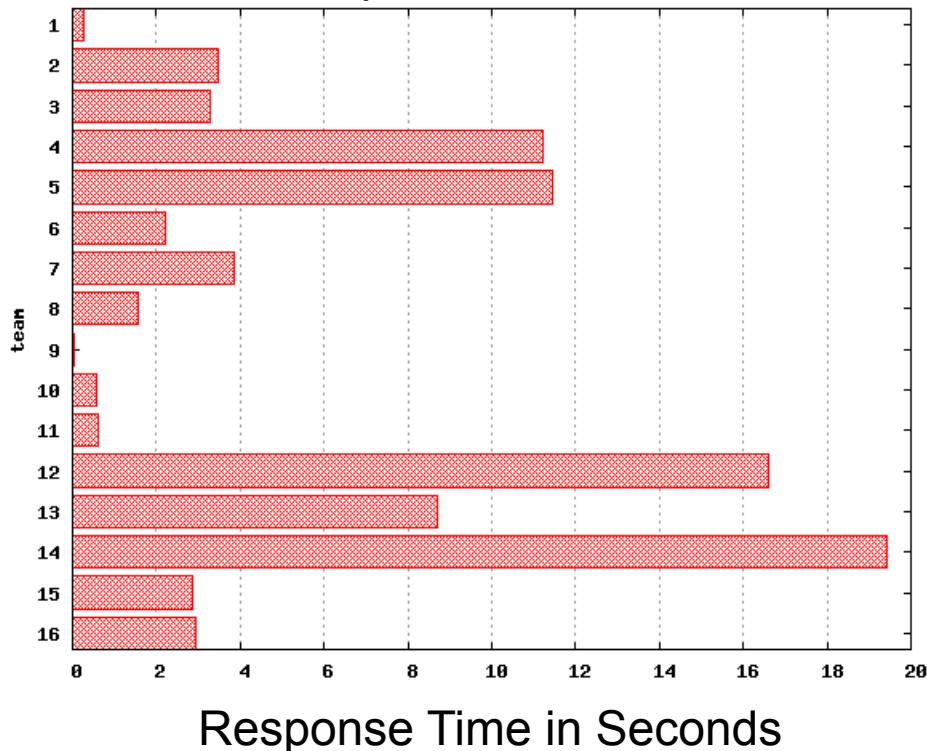


# CID Results: 18 Teams, 46 Runs

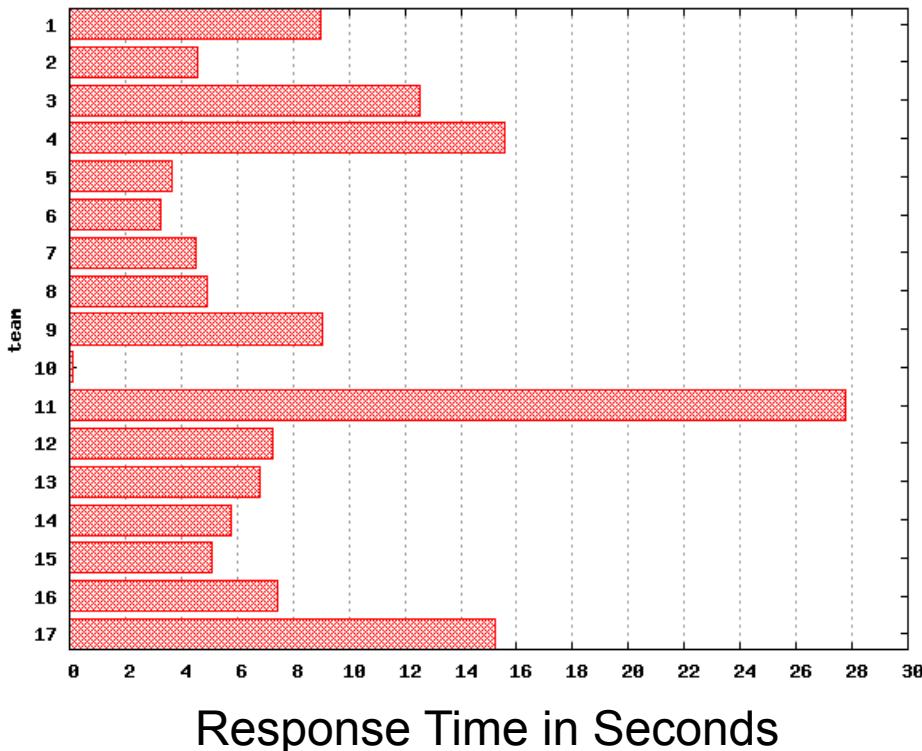


# System Response Time by Group

DNER Response Time – Best Run



CID Response Time – Best Run



# Post-Challenge Survey

- Use Task Supporting Tools?
- Use Additional Lexicon Resources?
- Use Machine Learning? Features?
- Use Existing BioNLP Tools?
- What General Software Tools?
- What Techniques?
- What Programming Language(s)?
- Pre/Post-Processing?

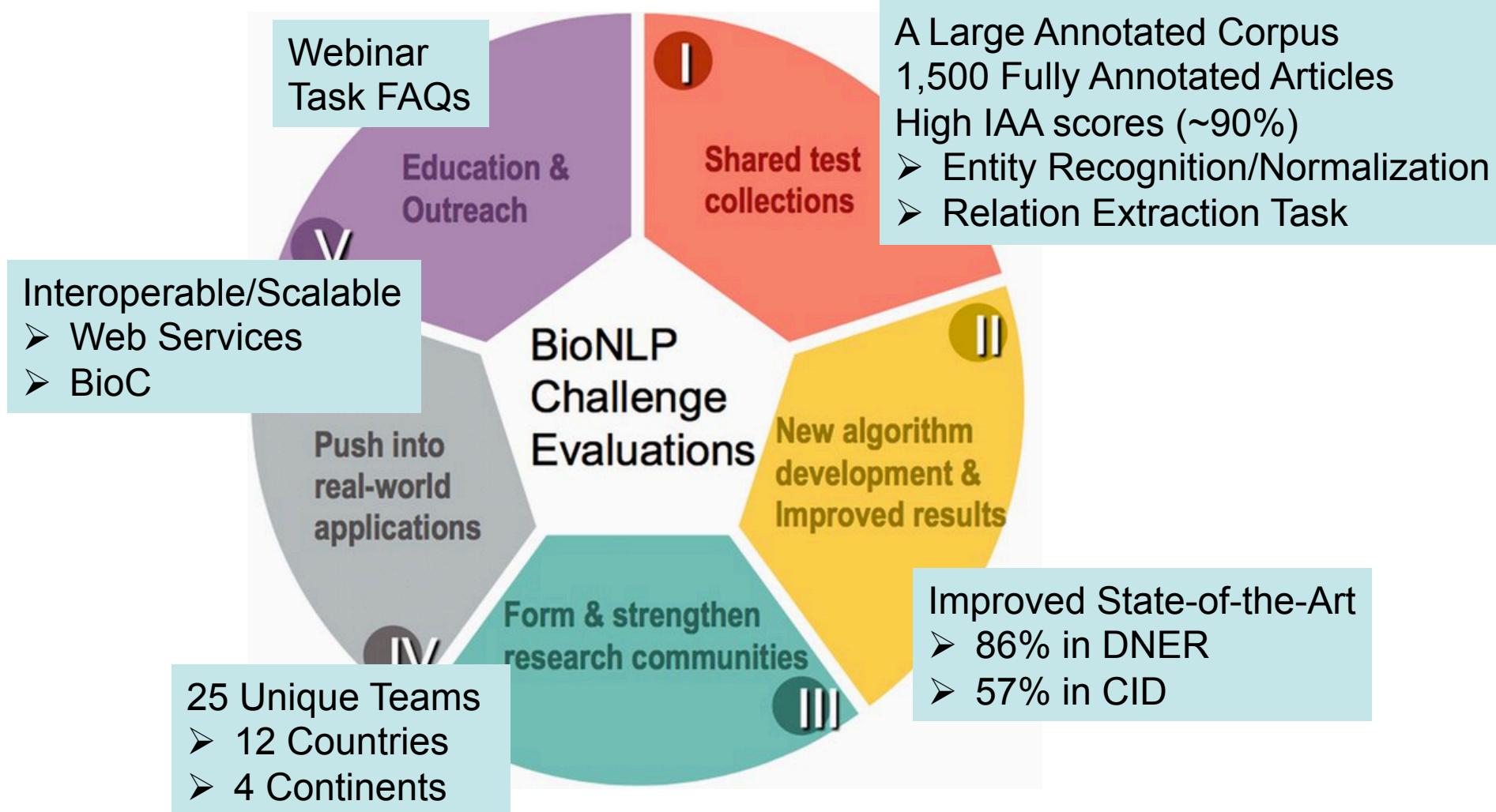
# Post-challenge Survey Results

Supporting Resources	DNER (16 teams)	CID (18 teams)
Web-service setup code	13	14
Evaluation scripts	11	10
DNorm	5	12
tmChem	0	13
PubTator (for data visualization & comparison)	7	4
BioC Library	3	4

# Survey of Team Methods



# Main Contributions



# Limitations

- Relation extraction is difficult!
- Other known issues:
  - Annotation is Imperfect (Human Errors)
  - No New IAAAs for Relation Annotations (High Cost)
    - Past CTD IAA Chemical-Gene Curation F1: 77%
  - Difficulty Running Web Services (in China)

# What's Next?

- 6 team presentations
  - Performance; Methods; Availability/Scalability
- Release CDR Test Set to Teams
- Invite Teams for Special Issue
- Propose a Meta Web Service for CIDs
  - Merging team results can improve performance
  - Make the results of this challenge more useful for the biocuration community and beyond



# Acknowledgements

*[ncbi.nlm.nih.gov](http://ncbi.nlm.nih.gov)*  
*[ctdbase.org](http://ctdbase.org)*



## ***NCBI Team:***

Zhiyong Lu  
Robert Leaman  
Yifan Peng  
Chih-Hsuan Wei

## ***CTD/NC State Team:***

Carolyn J. Mattingly  
Allan Peter Davis  
Thomas C. Wiegers  
Robin Johnson  
Daniela Sciaky

## ***CAM Team:***

Jiao Li  
Yueping Sun  
MeSH Indexers

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**NC STATE UNIVERSITY**

