

Big Data Science in Drug Discovery and Development

CIKM 2016 Tutorial Part II

October 24th, 2016

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Center for Computational Health

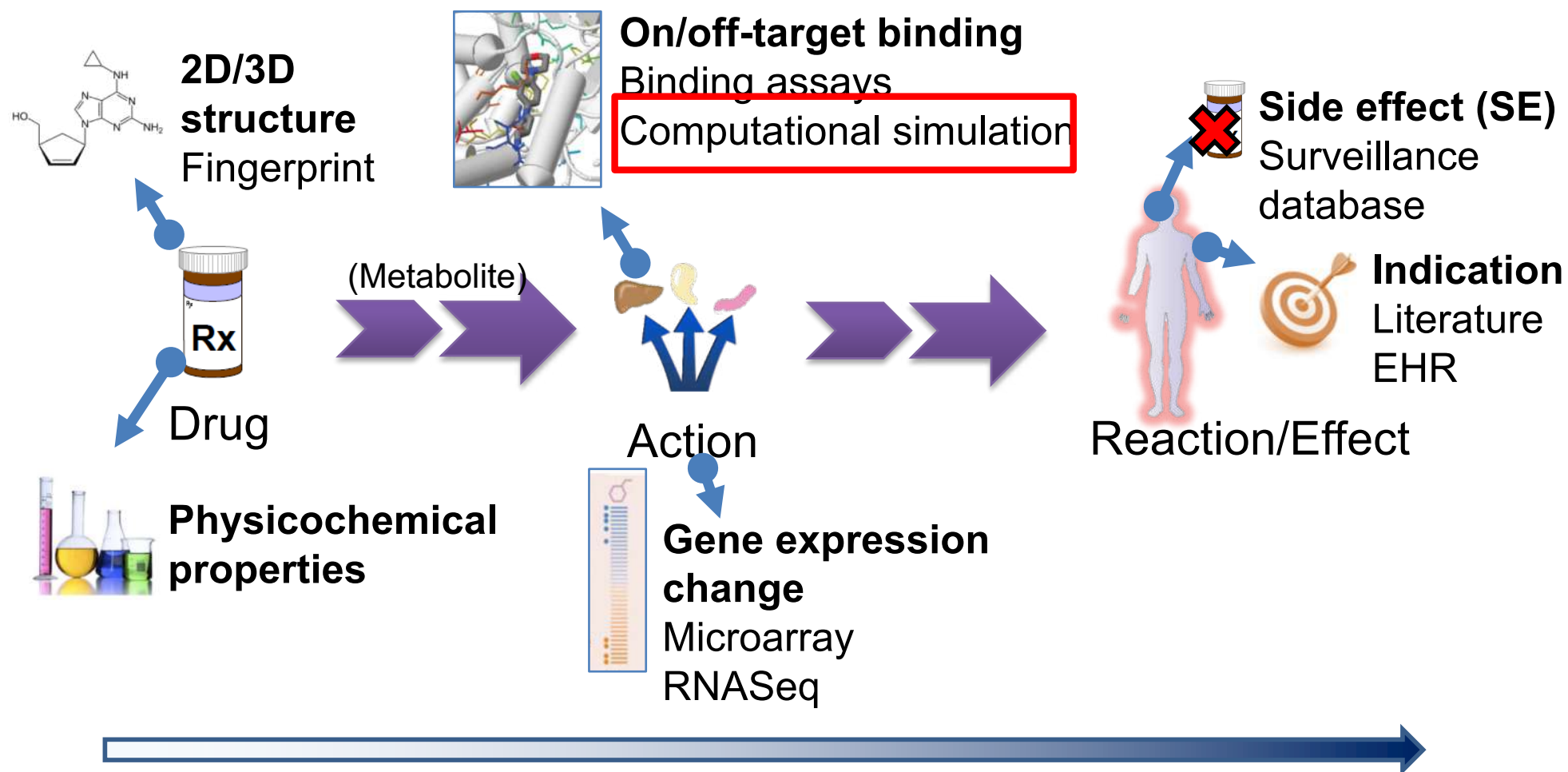
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Outline

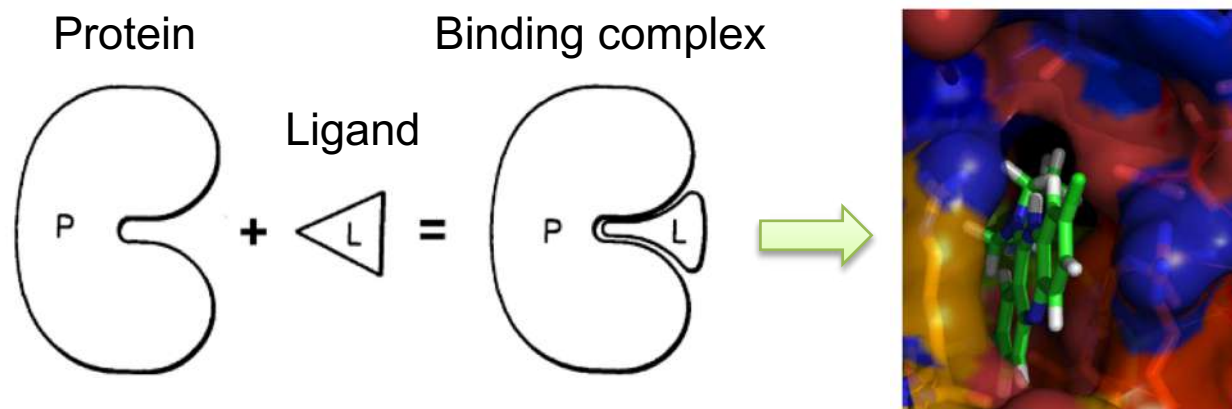
- Preclinical data analytics – chemical-protein interactome (CPI) as an example
 - Drug-drug interaction prediction
 - Drug repositioning
- Patient data analytics – real-world evidence (RWE) as an example
 - Drug safety signal detection from FAERS

Path from drug to effect



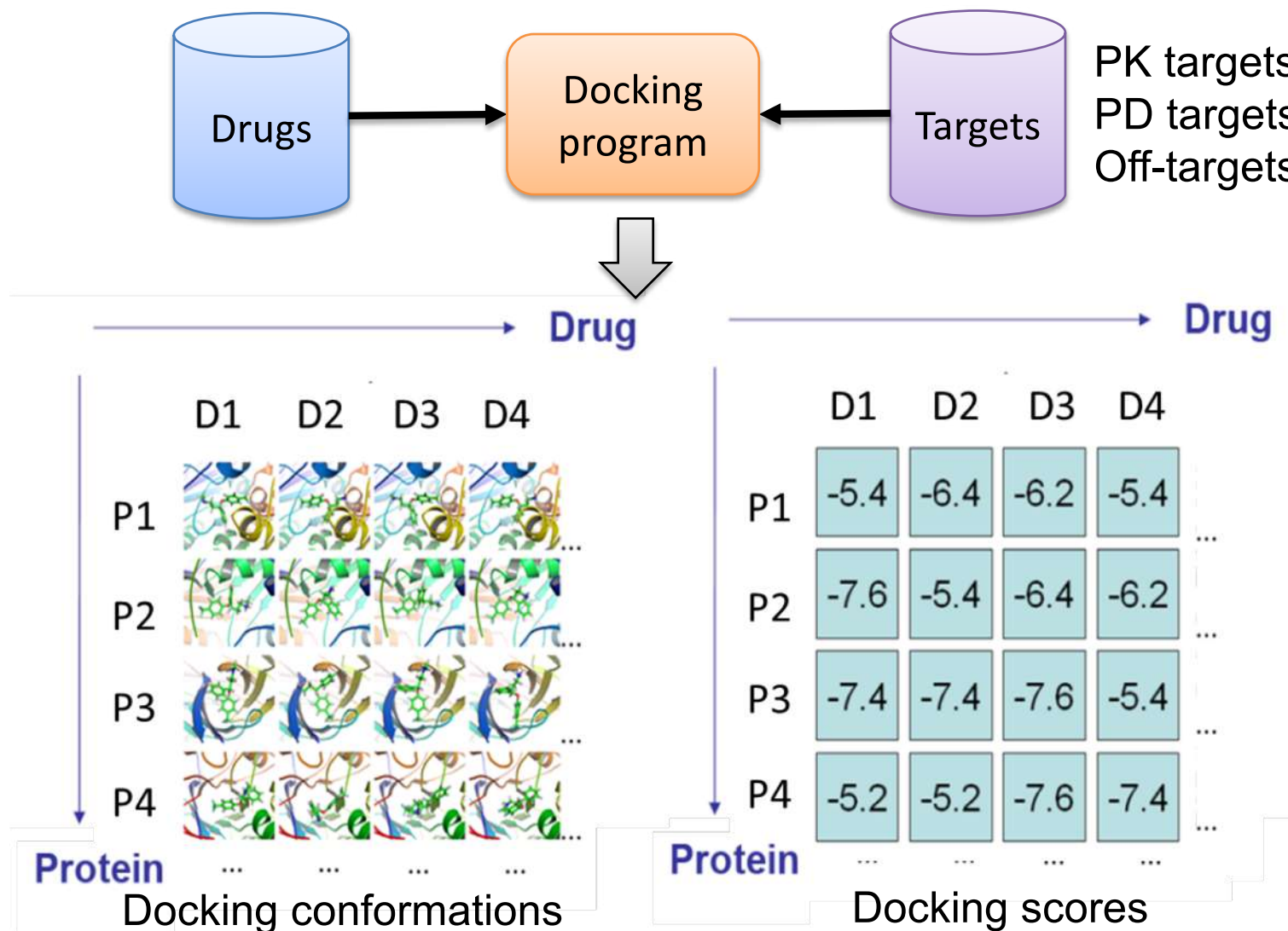
Molecular docking

- A docking program simulates the binding between a small molecule and a protein target.
 - Optimal binding position
 - Binding strength (docking score)



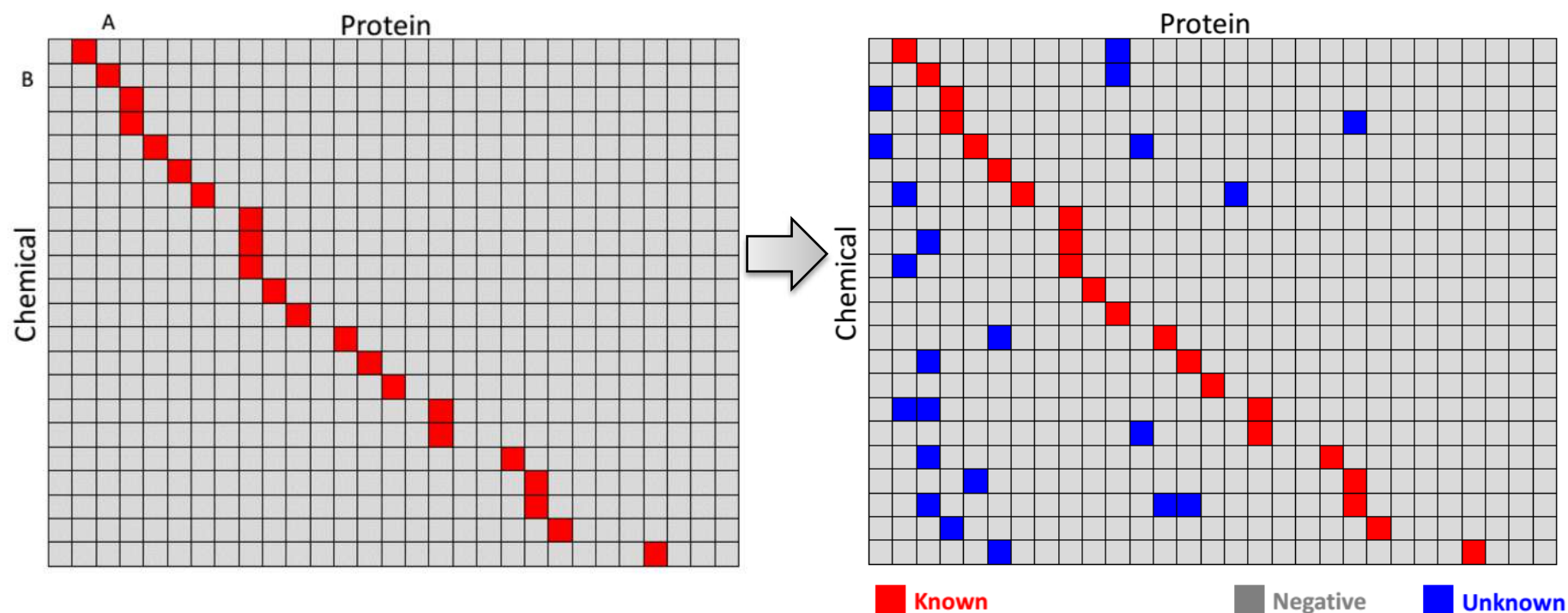
Drug binding inside the protein

Chemical-protein interactome (CPI)



Simulation of a chemical-protein interactome (CPI)

Why chemical-protein interactome?



- Expand the existing knowledge
 - Identify potential off-target binding
- Fast – 1 minute for a drug-protein pair
- Cheap – compared to wet-lab experiments

Application 1: Drug-drug interactions

- Older patients usually take more than one drug
- 1/25 individuals have adverse reaction caused by drug–drug interactions (DDIs)

Types of DDIs

- Potentiation
- Interference
- Antagonism
- Displacement



Two drugs compete for protein binding sites - a **major** cause for drug-drug interactions

Workflow of DDI-CPI server

Model training

(A) 12,656 drug pairs
(DrugBank)

Drug pair	DDI?
A and B	Yes
B and C	Yes
A and D	No

(B) Docking scores (2,515
drugs against 611 targets)

Drug	T ₁	T ₂	...
A	-9.3	-9.8	
B	-8.4	-10.1	
C	-7.3	-9.1	

Combine

(C) Training set

The sum and the absolute difference of the docking scores as features

Drug pair	T ₁		T ₂		DDI?
	Sum	Dif.	Sum	Dif.	
A and B	-17.7	0.9	-19.9	0.3	Yes
B and C	-15.5	0.9	-17.6	0.6	Yes
A and D	-19.7	1.1	-20.3	0.7	No

Train

(D)
Logistic
regression
models

Predict

Model prediction

(F) Docking score towards
611 targets

Drug	T ₁	T ₂	...
X	-6.0	-8.2	

Combine

(G) DDI predictions

Drug pair	T ₁		T ₂		DDI?	Probability
	Sum	Dif.	Sum	Dif.		
A and X	-15.3	3.3	-18.0	1.6	Yes	0.68
B and X	-14.4	2.4	-18.3	1.9	Yes	1.00
C and X	-13.3	1.3	-17.3	0.9	Yes	1.00



(E) Drug X

Submit


Calculate

DDI-CPI
server

Demo: DDI-CPI

Submit a molecule - DDI-CPI | Drug Interactions (Sertraline) ... | Binding pattern of User drug (...)

cpi.bio-x.cn/ddi/

 **DDI-CPI**, a server Predicting Drug-Drug Interaction via Chemical-Protein Interactome

cpi.bio-x.cn/ddi

Navigation: [Home](#) > [Submit a molecule](#) Welcome **guest!** [Submit a molecule/](#) [Log out](#)

- Home
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- How to cite

Submit a molecule

In order to protect privacy, your submissions will not be shown to others. ?

You can upload a single-molecule file to be processed by our server.

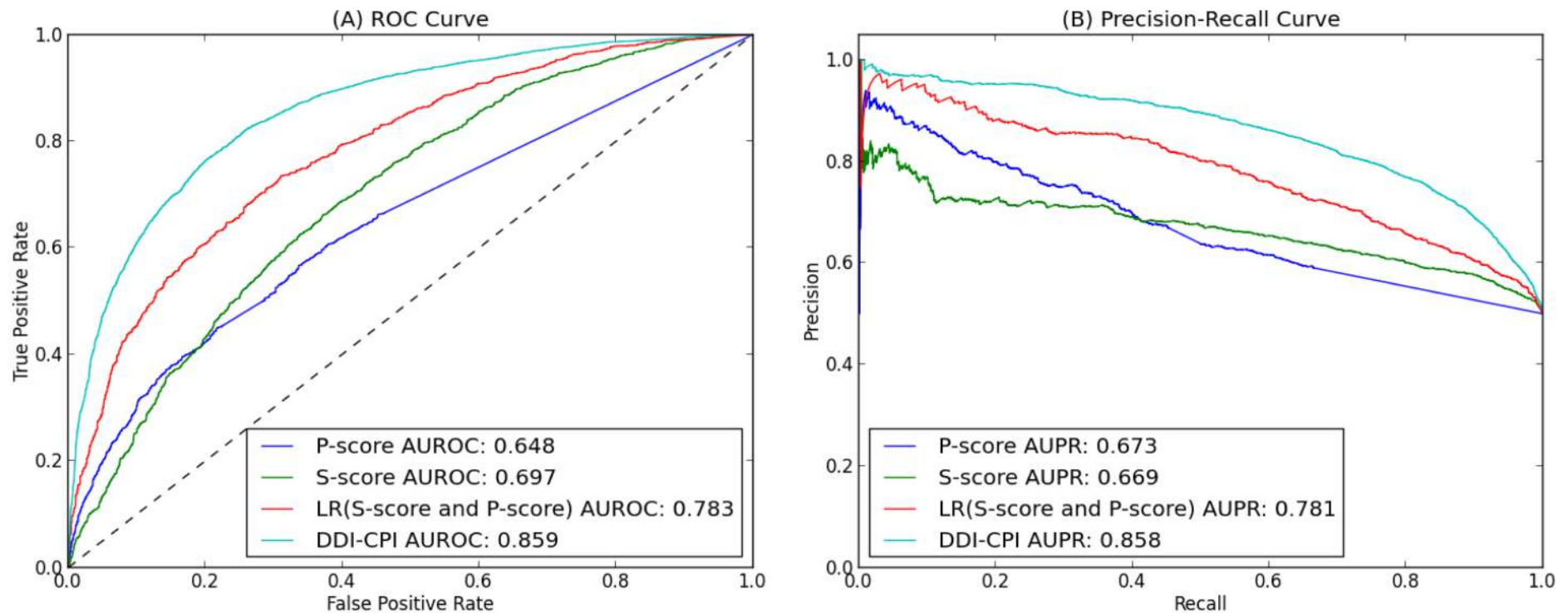
Here is [an example file](#), upload it and wait for about 15 mins to check the result.

Upload a molecular file:	<input type="button" value="Browse..."/> No file selected.	*type: mol/ml2/mol2/pdb/pdbqt/sdf/smiles Instructions to prepare a molecule file
Or input SMILES string*:	<input type="text"/>	<input type="button" value="Draw"/> Molecular name*: <input type="text"/>
E-mail Address: ?	<input type="text"/>	The access link will be sent in the email (optional)
Your remark:	<input type="text"/>	
<input type="button" value="Submit"/> <input type="button" value="Reset"/>		

Disclaimer: The server is for research purposes only and the authors and their organizations are excluded from all liability for any costs, claims, expenses, charges, losses, damages or penalties of any kind incurred directly or indirectly arising from the use of this server.

Recommended browsers: FireFox, Chrome or Internet Explorer 9 (HTML5 support), resolution: 1366*768 or higher

Results



The ROC and precision-recall curve comparison for different DDI prediction methods based on independent validation

P-score: uses side-effect similarities to predict target sharing (Campillos, et al. Science (2008), 321, 263-266.)

S-score: uses drug-target network to predict DDIs (Huang, et al. PLoS Comput Biol (2013), 9, e1002998)

LR(S-score and P-score): integrates P-score and S-score by a Bayesian probabilistic model

DDI-CPI: predicts DDI using machine learning models via CPI

MAOI: Monoamine oxidase inhibitor
SSRI: Selective serotonin reuptake inhibitor

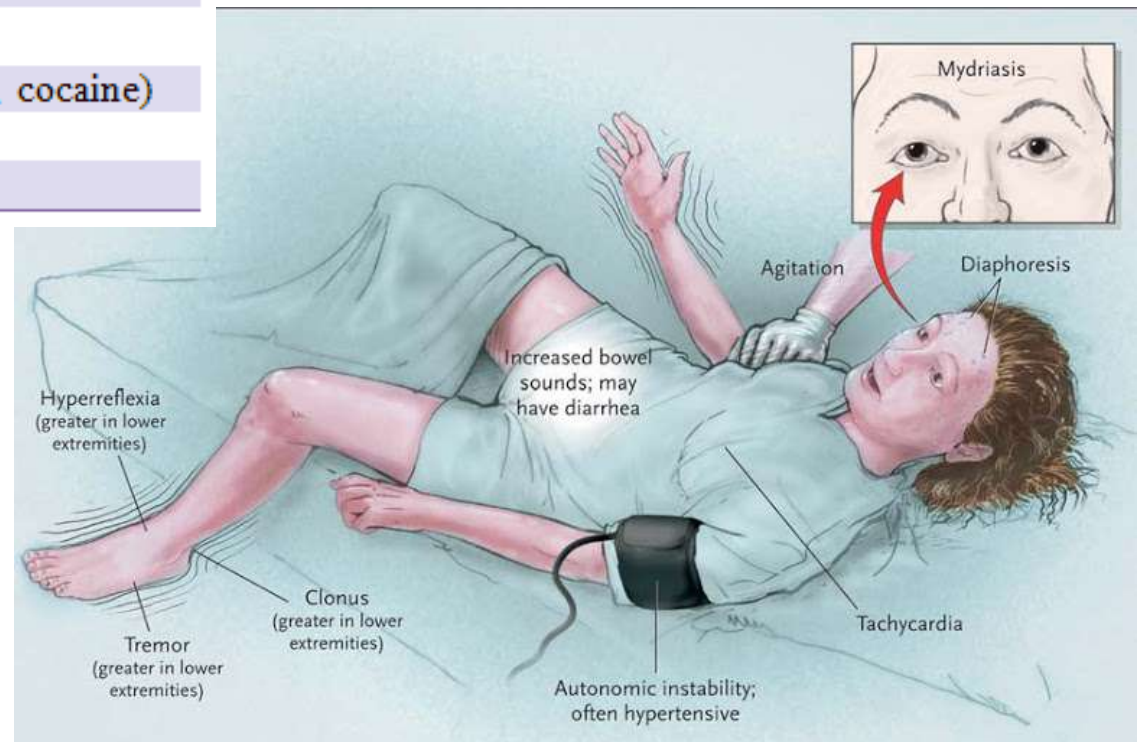
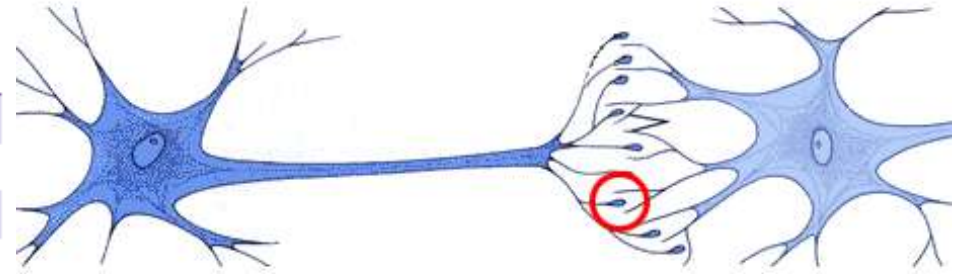
Case study - MAO-A inhibitors

Table 3 (adapted from reference 6,7)

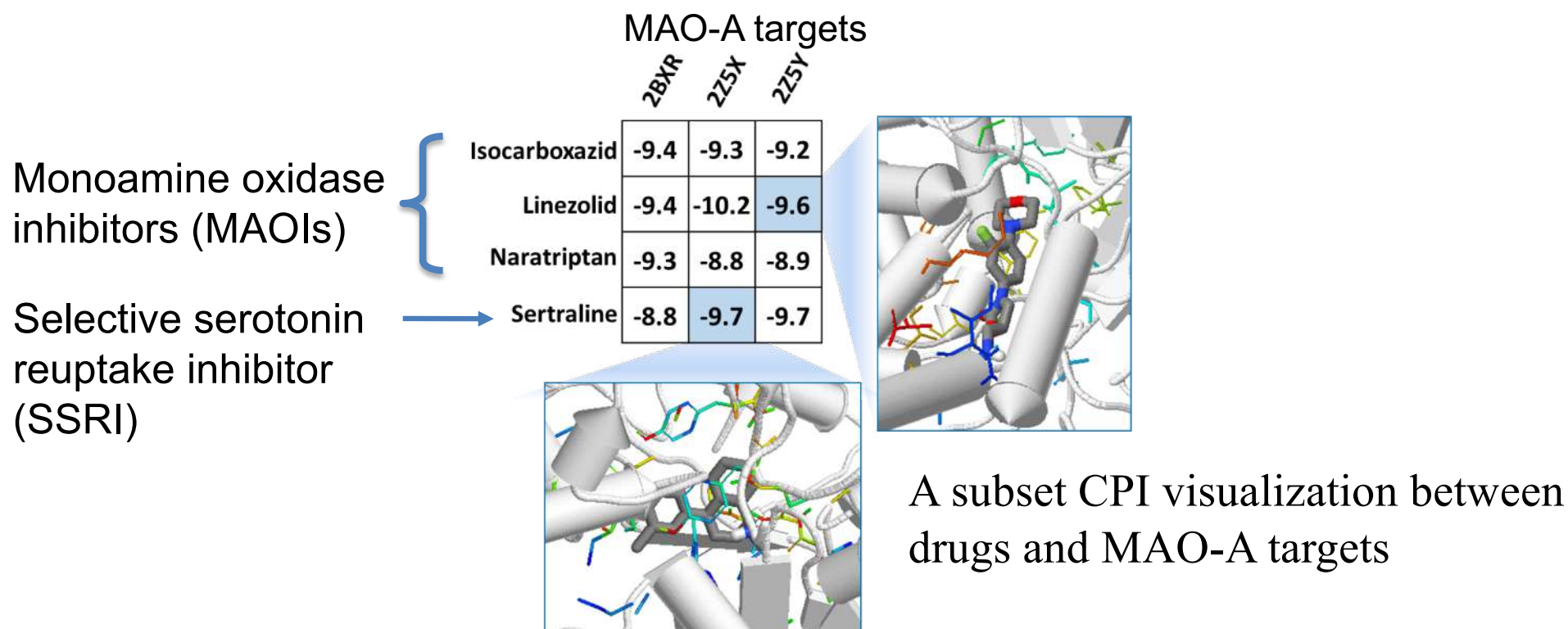
Drugs to Avoid When Taking MAOIs

Amphetamines	Bupropion
Cyclobenzaprine	Dextromethorphan
Linezolid	Meperidine
Methadone	Mirtazapine
<u>SSRIs/SNRIs</u>	TCA's
Triptans	Tramadol
Vasoconstrictors (pseudoephedrine, phenylephrine, cocaine)	
Chlorpheniramine, brompheniramine	
St. John's Wort	General anesthesia

- SSRI with MAOI results in high extracellular serotonin (5-HT) concentration – serotonin syndrome.



Case study - MAO-A inhibitors



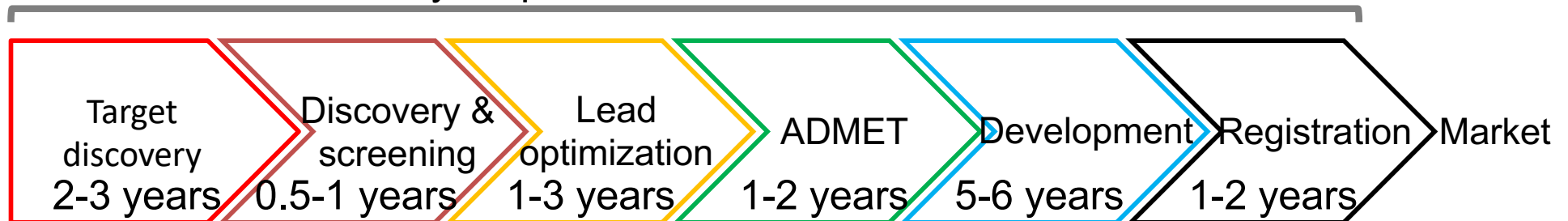
- The server predicts that sertraline may interact with isocarboxazid, linezolid, and naratriptan
- All of the predicted drugs can rank the MAO protein structures to the top 20% – possible mechanism suggested

Application 2: Drug repositioning

- Identify new indications for existing drugs.

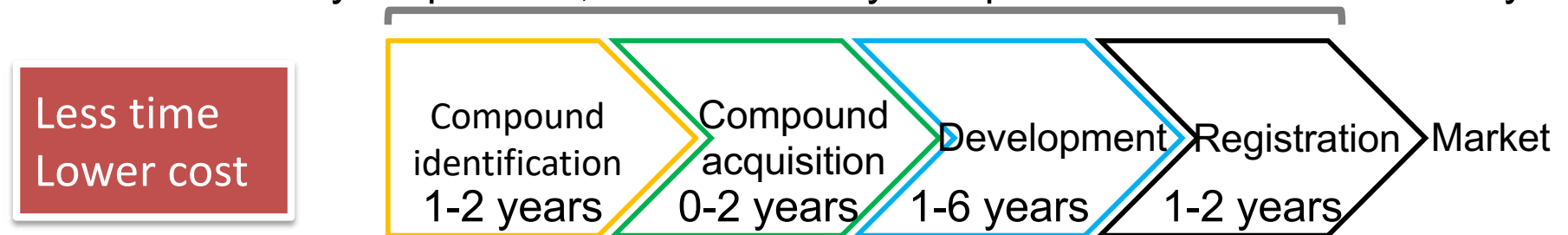
De novo drug discovery and development

10-17 year process, <10% overall success rate















Drug repurposing

3-12 year process, reduced safety and pharmacokinetic uncertainty

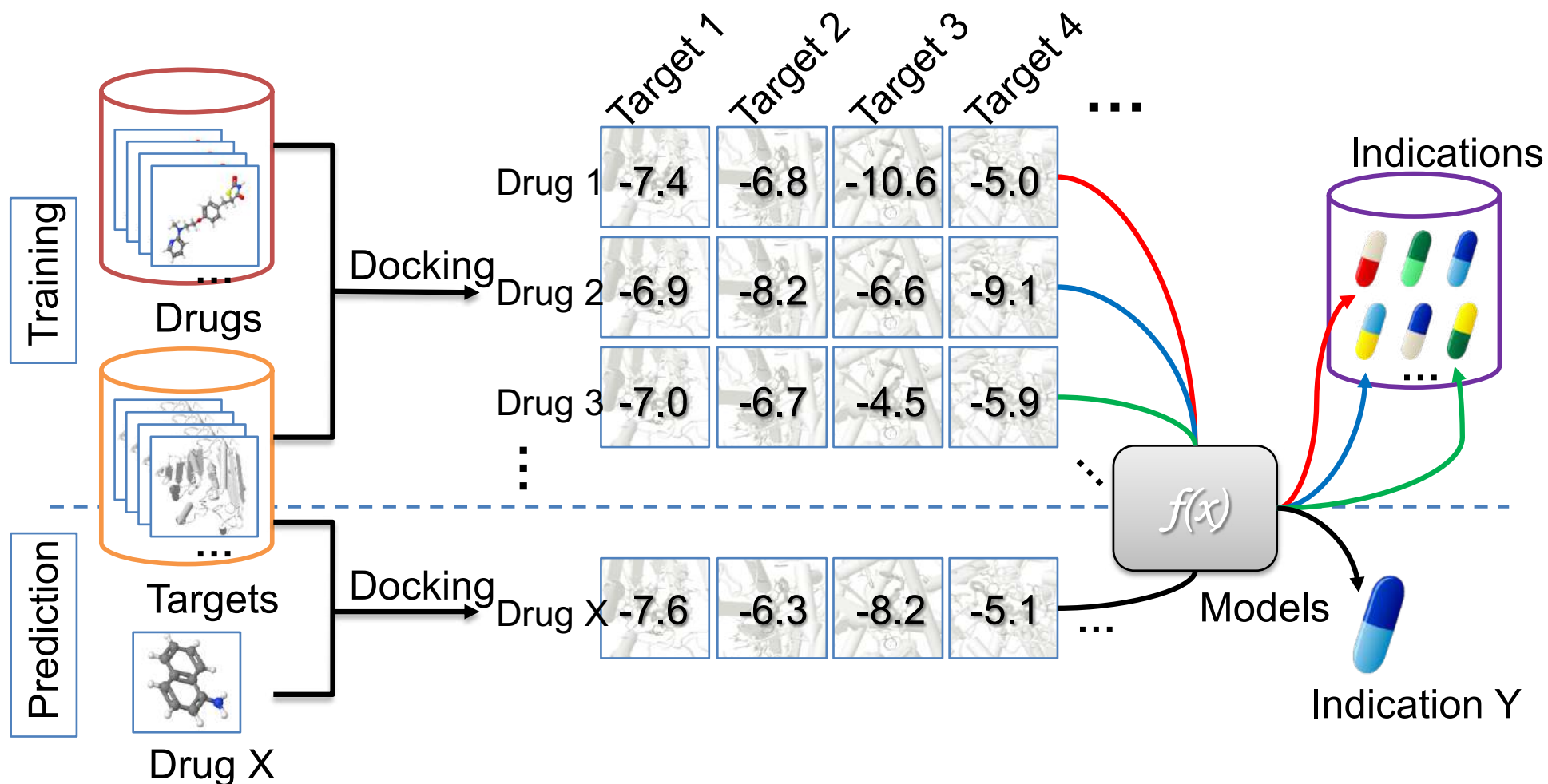


Revenue to pharmaceuticals

1.  77% \$\$ 54% Rx	2.  58% \$\$ 15% Rx	3.  50% \$\$ 20% Rx	4.  44% \$\$ 28% Rx
5.  37% \$\$ 24% Rx	6.  29% \$\$ 25% Rx	7.  24% \$\$ 17% Rx	8.  20% \$\$ 20% Rx
9.  19% \$\$ 21% Rx	10.  13% \$\$ 36% Rx	11.  9% \$\$ 47% Rx	12.  6% \$\$ 15% Rx

Contribution of the repositioned indications to the sales in 2011

Workflow of DPDR-CPI Server



Demo: DPDR-CPI

Home - DPDR-CPI Drug indication predictions (rosiglitazone) - DPDR-CPI Binding pattern of User drug Rosiglitazone with Amine oxidase [flavin-containing] A - DPDR-CPI

 **DPDR-CPI** , a server predicting Drug Candidate Positioning and Drug Repositioning via Chemical-Protein Interactome

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Navigation: [Home](#) Welcome guest! [Submit a molecule/](#) [Log out](#)

Welcome to our site, **guest!** (Click here to Log out)

View: Submission center (Submit a molecule) - Backward

Introduction

Background

Identifying the best indications for pipeline drug candidates (drug candidate positioning) and recycling old drugs for new indications (drug repositioning, or repurposing) are now attracting great interests in the pharmaceutical industry and academia due to the high attrition rate of developing a new drug *de novo* and looking for the right indication. Inspired from our previous server ([server link](#) and [publication link](#)) on predicting drug repositioning via Chemical-Protein Interactome (CPI), now we made significant changes and improvements to introduce the new DPDR-CPI server for both drug candidate positioning and repurposing.

What can DPDR-CPI do for you?

When you submit a molecule in in MOL/MOL2/PDB/SDF/SMILES format, the server will suggest potential indications with estimated confidence across 963 different disease indications by machine learning models. The server will also suggest putative targets and their docking conformations across 611 pharmacokinetic (PK) and pharmacodynamic (PD) targets. The server can visualize each ligand-protein binding pattern, with amino acid residues around 6.4 Å of the ligand highlighted

Server work flow



```
graph TD; A([Upload a drug molecule]) --> B[Wait the interactome of your drug molecule to be constructed (several hours)]; B --> C[Check the predicted indications for your molecule]; C --> D[Check the candidate off-targets tend to interact with your molecule];
```

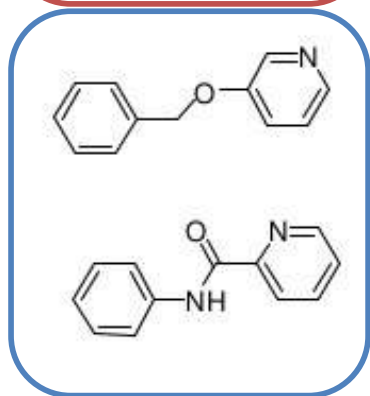
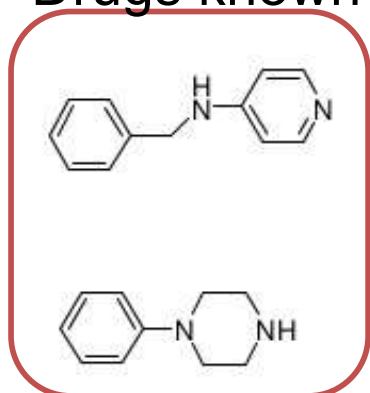
Click here to see
[\[Sample drug predicted by our server\]](#)

RSS feeds: [Target updates](#), [Library drug updates](#)

State-of-the-arts: Various fingerprints

Positive set

Drugs known to treat disease Y



Negative set

Drugs unknown with disease Y

Descriptors

001010010010101010101
110010101010100001001
010100001010101010101
000010010100101001010

Treat Disease Y?

Yes
Yes
No
No

$$f(x)$$

Machine learning
model for disease
Y

PubChem 881

Fingerprints

Bit Position Bit Substructure

0 ≥ 4 H

1 ≥ 8 H

4 ≥ 1 Li

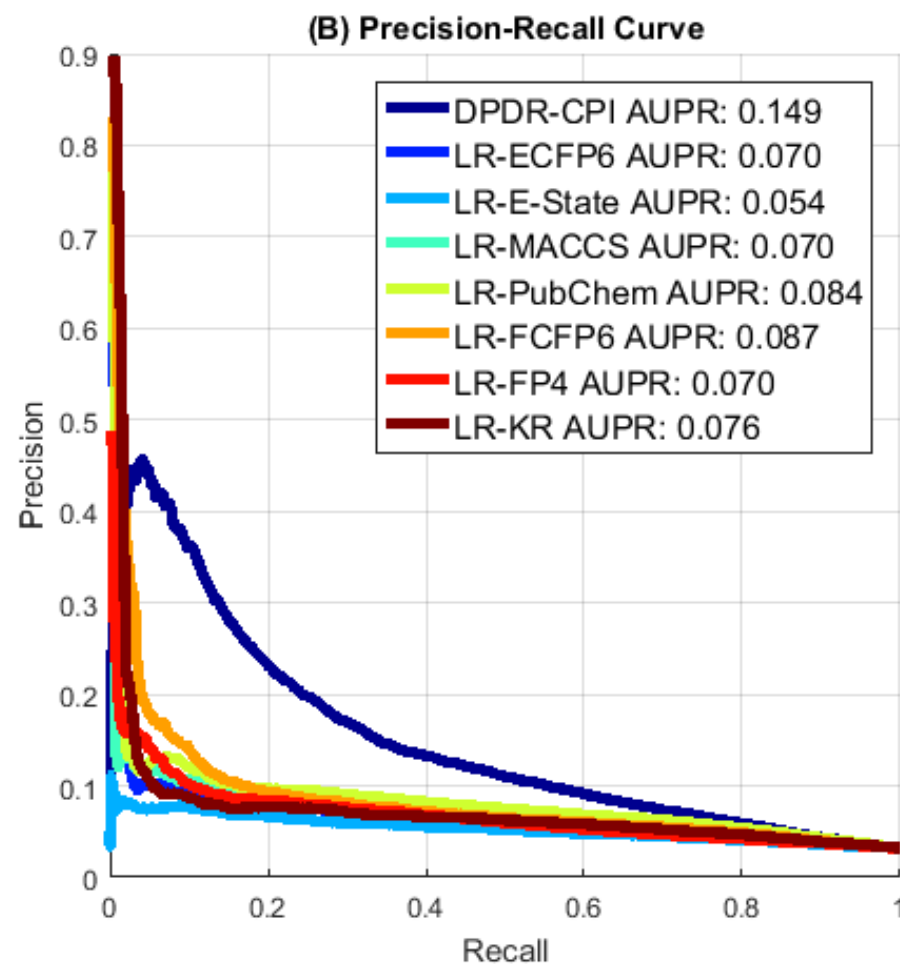
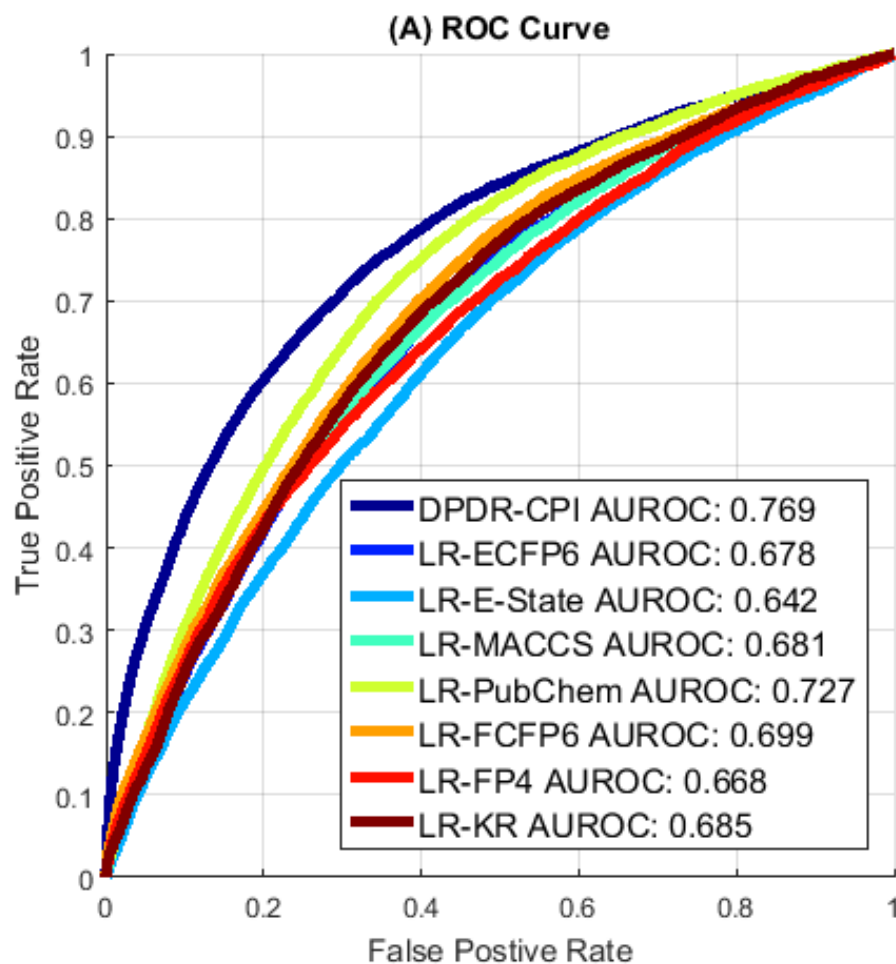
5 ≥ 2 Li

9 ≥ 2 C

10 ≥ 4 C

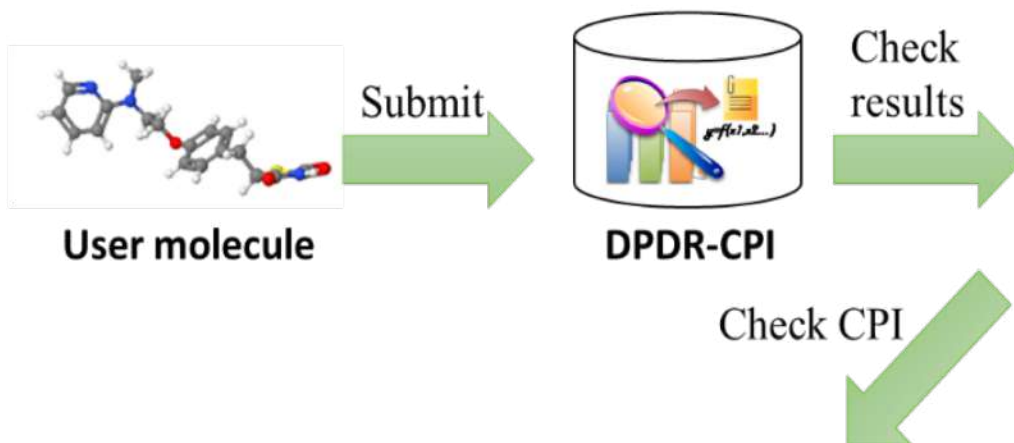
...

Performance comparison



Performance comparison between DPDR-CPI and chemical structure-based predictors based on independent validation set

Case Study - Rosiglitazone



DPDR-CPI, a server predicting Drug Positioning and Drug Repositioning via Chemical-Protein Interactome

Home Submit a molecule Log out Help Contact us

Navigation: Home > Submit a molecule > User drug: rosiglitazone > CPI Welcome guest! Submit a molecule/ Log out

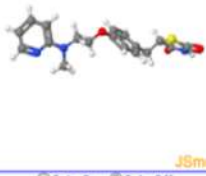
Chemical-Protein Interactome (rosiglitazone)

Interactome of your molecule across the targetable proteins (download)

Total: 611 Page: 1 / 31 Search one word: Go Reset

No	PDB ID	Class	Target name	Function	Docking Score	Visualization
1	2BZG	PK	Thiopurine S-methyltransferase	Catalyzes the S-methylation of thiopurine drugs such as 6-mercaptopurine.	-10.5	Visualization
2	1FCY	PD	Retinoic acid receptor gamma	Receptor for retinoic acid. Retinoic acid receptors binds heterodimers to their target response elements in response to their ligands, all-trans or 9-cis retinoic acid, and regulate gene expression in various biological processes. The RAR/ROR heterodimers bind to the retinoic acid response elements (RARE) composed of tandem 5'-AGGTCA-3' sites known as DR1-DR3. In the absence of ligand, acts mainly as an activator of gene expression due to weak binding to corepressors. Required for limb bud development. In concert with RARA or RARB, required for skeletal growth, matrix homeostasis and growth plate function (by similarity).	-10.3	Visualization
3	2H11	PK	Thiopurine S-methyltransferase	Catalyzes the S-methylation of thiopurine drugs such as 6-mercaptopurine. Catalyzes the oxidative deamination of biogenic	-10.2	Visualization

Molecule information



JSmol

☐ Spin On ☒ Spin Off
background white
[Reset to original orientation](#)
[Download molecule](#)

Target binding predictions

DPDR-CPI, a server predicting Drug Positioning and Drug Repositioning via Chemical-Protein Interactome

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Navigation: Home > Submit a molecule > User drug: rosiglitazone Welcome guest! Submit a molecule/ Log out

Drug indication predictions (rosiglitazone)

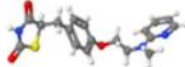
Please bookmark this page for later access.

Predicted indications (download)

Total: 54 Page: 1 / 3 Search one word: Go Reset

No	Disease family
1	251: Other disorders of pancreatic internal secretion (0.95) • 251.2: Hypoglycemia, unspecified (0.83)
2	250: Diabetes mellitus (0.93) • 250.1: Diabetes with ketoacidosis, type II or unspecified type, not stated as uncontrolled (0.93) • 250.10: Diabetes mellitus with ketoacidosis, type I [juvenile type], not stated as uncontrolled (0.89) • 250.00: Diabetes mellitus without mention of complication, type II or unspecified type, not stated as uncontrolled (0.81) • 250: Diabetes mellitus (0.80)
3	362: Other retinal disorders (0.91) • 362.83: Retinal edema (0.73) • 362.10: Background retinopathy, unspecified (0.62) • 362.9: Unspecified retinal disorder (0.62)
4	277: Other and unspecified disorders of metabolism (0.87) • 277.85: Disorders of fatty acid oxidation (0.81)

Molecule information

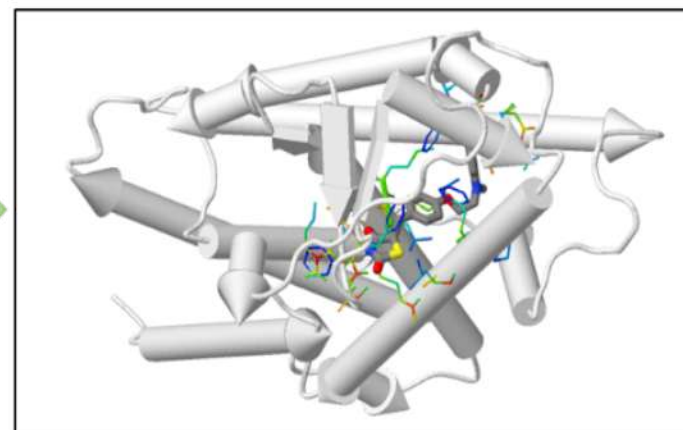


JSmol

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background white
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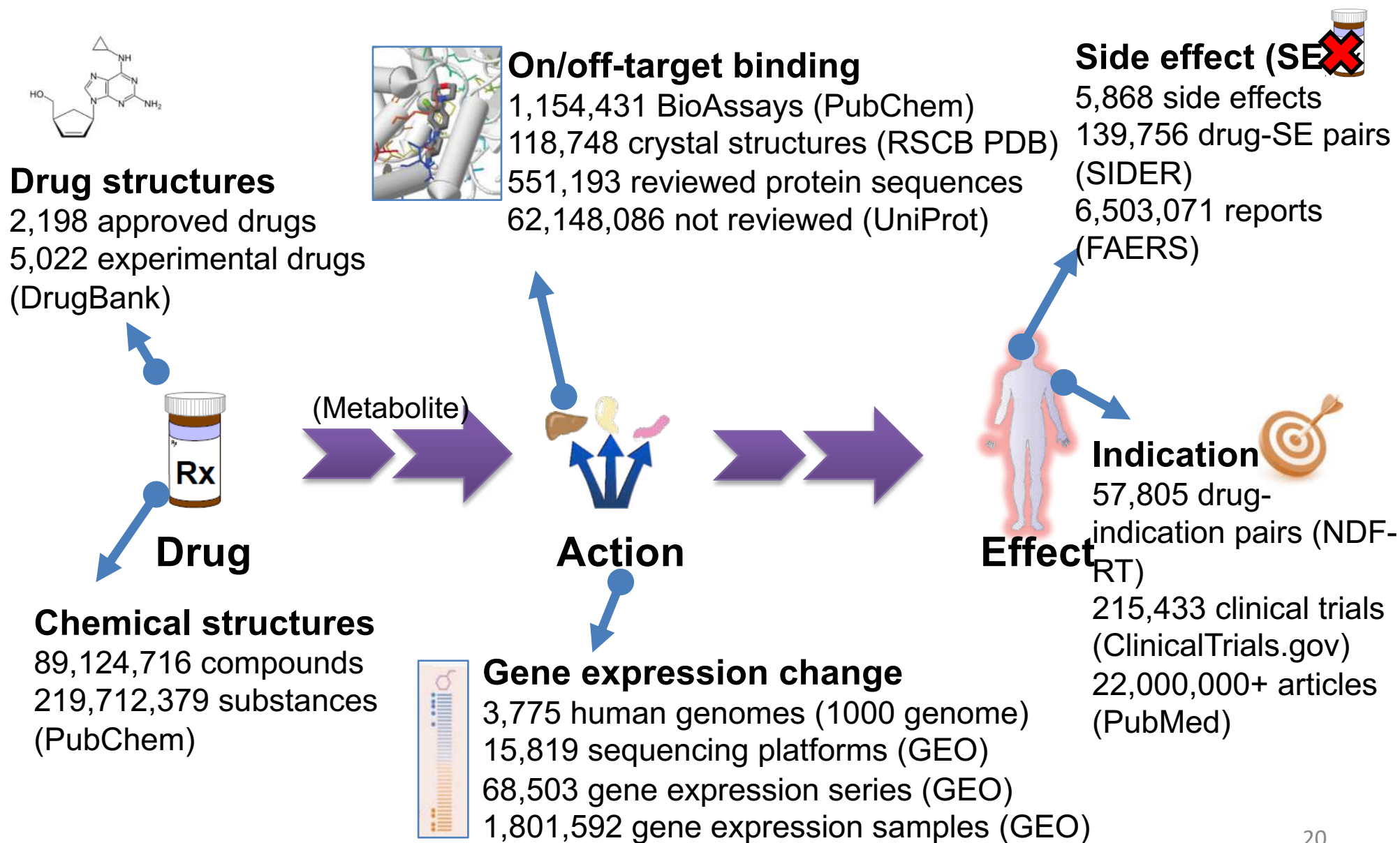
Indication predictions

Visualize binding

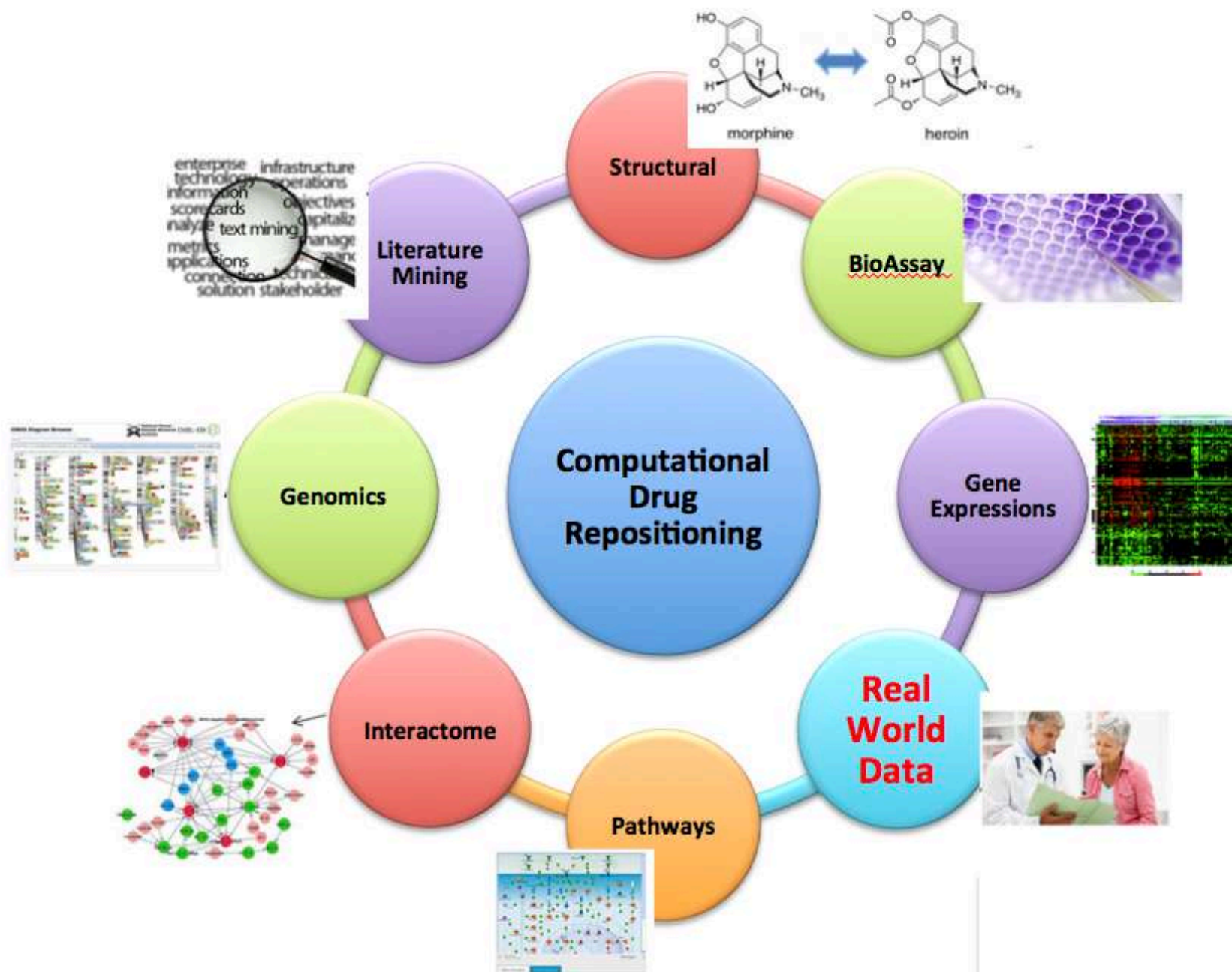


Binding conformation

Free big data in the domain



Next: Multi-channel detailed computational hypothesis generation



And even beyond the hypothesis generation...

biology

chemistry

dmpk

pharmacology

toxicology

Home » Pharmacology » Diabetes and Obesity » Obese Mice

ob/ob Diabetes Model - 16 Mice

Service Description

Provider: is a US company with laboratories in Hangzhou, China. The laboratory has been offering exploratory (non-GLP) pharmacology services to US and Chinese biopharma since 2004.

Background: The obese mutant mouse model was first reported by Ingalls A *et al* from the Jackson Laboratory in 1951 (*Obese, a New Mutation in the House Mouse* [164 KB]). The obese mouse resulted from a spontaneous mutation in a gene that was named *ob* in the V stock. Mice homozygous for the obese spontaneous mutation, (*Lep^{ob}ob*; commonly referred to as *ob* or *ob/ob*), are first recognizable at about 4 weeks of age. Homozygous mutant mice gain weight rapidly and may reach three times the weight of wild-type controls. In addition to obesity, mutant mice exhibit hyperphagia, a diabetes-like syndrome of hyperglycemia, glucose intolerance, elevated plasma insulin, subfertility, impaired wound healing, and an increase in hormone production from both pituitary and adrenal glands. Friedman J *et al* reported leptin in 1994, and demonstrated that leptin, the product of the *ob* gene, was produced in white adipose tissue and served as the peripheral signal to the central nervous system of nutritional status.

Service Details: This service offers a 28 day db/db mouse model of T2DM and obesity. Customer has various options that are conveyed to Links Biosciences using a Service Order Form. Customer assigns up to 16 mice to


\$9,000.00 USD


per service


9 week

turn around time

Provided By


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Holger Wesche, Principal Scientist, Large Pharma

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Big data researchers
will have a higher
impact in biomedicine
😊

Validation methods are increasingly commoditized

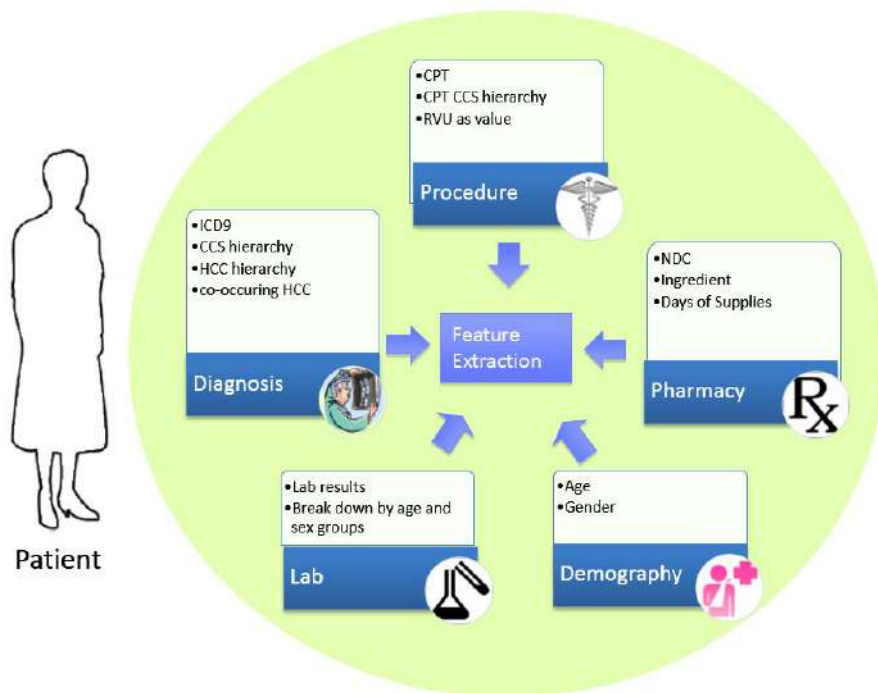
22

Outline

- Preclinical data analytics – chemical-protein interactome (CPI) as an example
 - Drug-drug interaction prediction
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- Patient data analytics – real-world evidence (RWE) as an example
 - Drug safety signal detection from FAERS

What is “Real World Evidence” (RWE)

- RWE is clinical observations other than randomized clinical trials (RCT).
 - RWE are large-scale clinical observations from population
 - RCT are expensive and in far smaller scale
- RWE is observations on human in the clinical stage
 - Less of a translational issue
 - Other than "omics", numerous external factors (e.g., environment, diet and exercise) affect response to medication
- RWE is not only vast but also varied in type and source: electronic medical records (EMR), claims data, and even social media.



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twitter



Application 3: Adverse drug reactions (ADRs)

- Post-approval ADRs remain a significant source of mortality and morbidity around the world
 - 2 million potentially preventable injuries, hospitalizations, and deaths each year in US alone
 - Associated cost estimated at \$75 billion annually

The New York Times

F.D.A. Issues New Alerts About Cholesterol Drugs


By GARDINER HARRIS

Published: February 29, 2012

CORRECTION APPENDED

Federal health officials on Tuesday added new safety alerts to the prescribing information for statins, the cholesterol-reducing medications that are among the most widely prescribed drugs in the world, citing rare risks of memory loss, diabetes and muscle pain.

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**SOUND OF
IN THEATRES**

Statins are considered some of the safest drugs

Merck Pulls Arthritis Drug Vioxx from Market

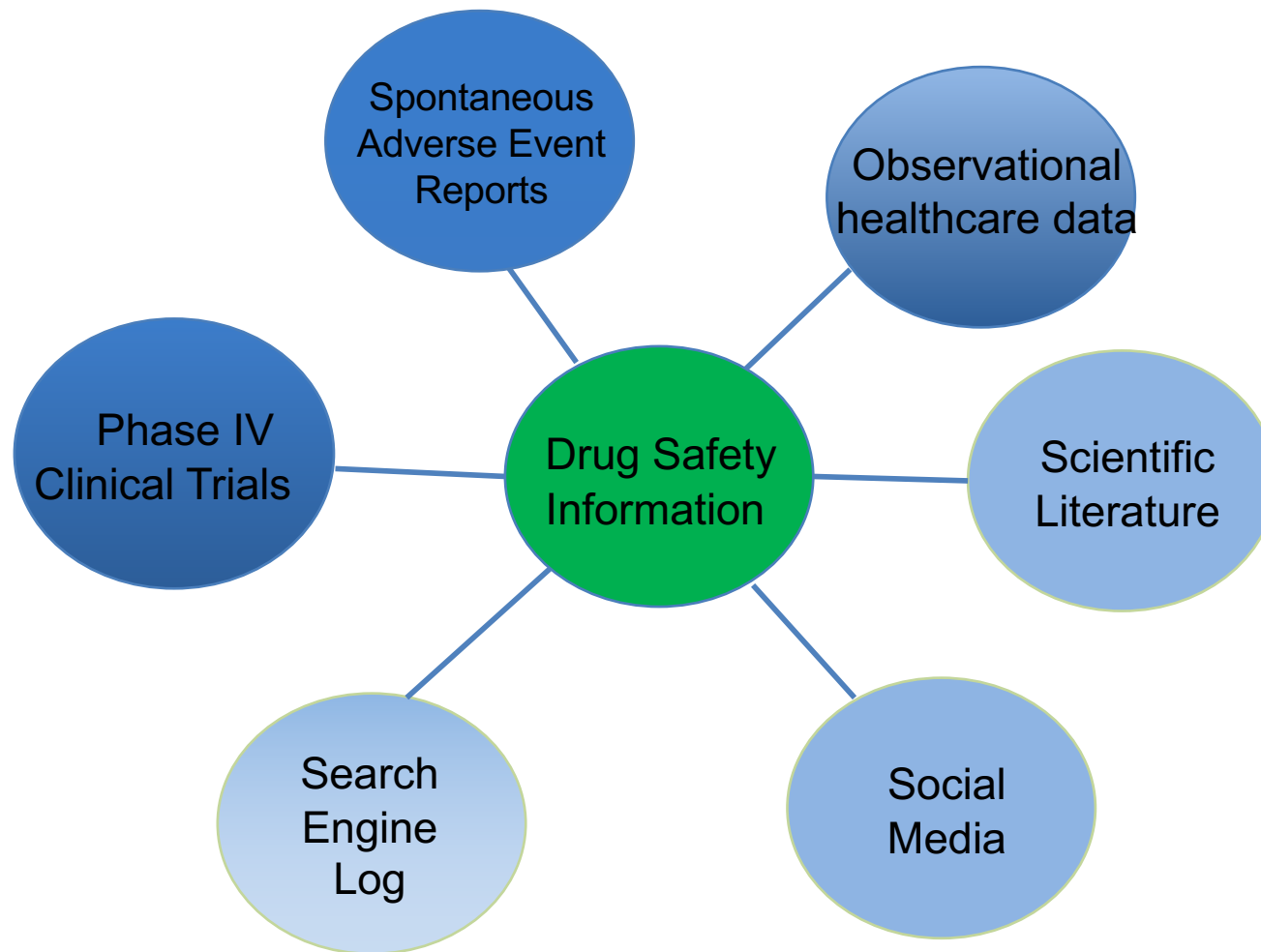
by RICHARD KNOX

September 30, 2004 12:00 AM ET

Pharmaceutical giant Merck & Co. is pulling its arthritis drug Vioxx from the market after a study confirmed earlier concerns that it raises the risk of heart problems, including heart attacks and stroke. Vioxx is currently used by 2 million people worldwide and has been used by more than 84 million people around the world, according to Merck.

- More than 140,000 cases of serious heart disease
- \$4.85 billion for legal claims from US citizens

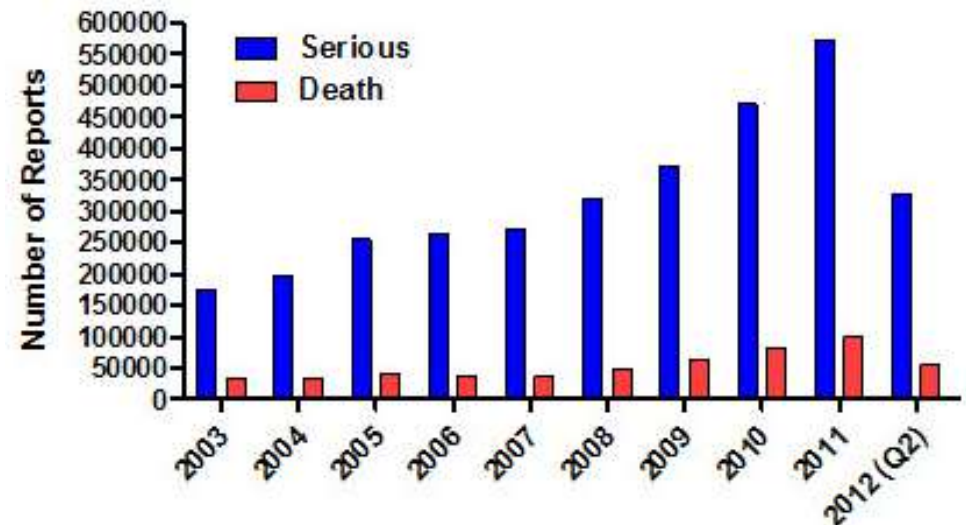
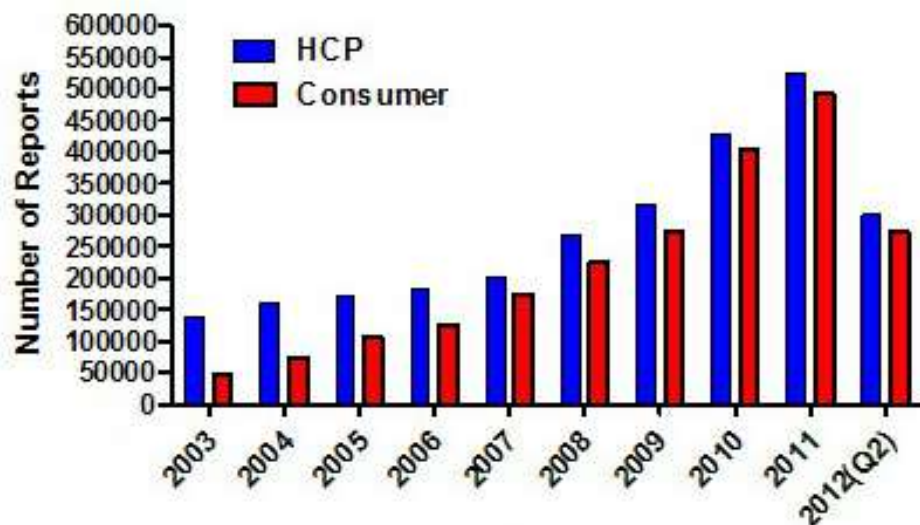
Data sources of drug safety information in post market stage



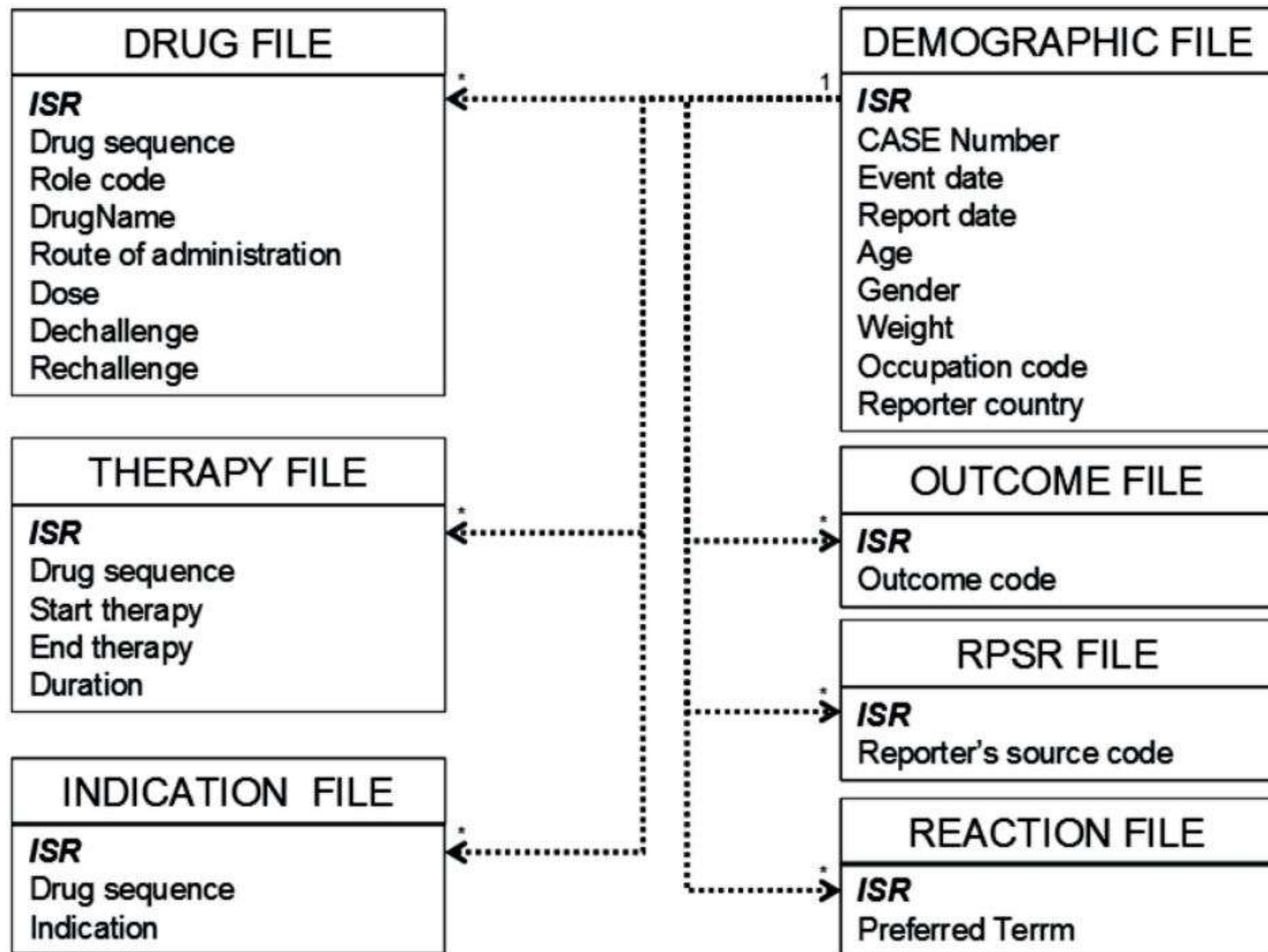
FDA Adverse Event Reporting System (FAERS)

- FDA Adverse Event Reporting System (FAERS)
 - FDA has maintained AERS since 1968
 - Spontaneous reports of suspected ADRs collected from healthcare professionals, consumers, and pharms
 - Data (from Jan 2004 to June 2016) is publicly available at FDA's website!
- Over 5 million reports collected so far:
 - patient: age, sex, weight, country
 - drugs they are taking
 - diseases they were being treated for
 - the adverse events that occurred to that patient

Often sparsely collected



FAERS database structure

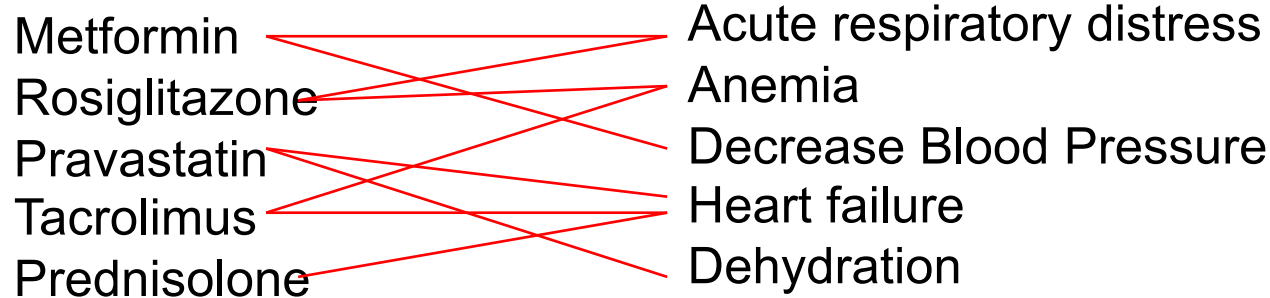


Interpreting those FAERS reports is hard

- Many drugs, many adverse events
 - what causes what?
 - Most of these red lines are false - which are true?

Drugs

Adverse Events



- Signal detection algorithms for FAERS
 - Quantify “unexpectedness”: to identify drugs that have a greater proportion of a particular event compared to the proportion seen for other drugs
 - Sampling variance
 - Underreporting
 - Over reporting
 - Selection biases
 - Causative covariates other than drug under analysis

Disproportionality analysis

	reports w ae	reports w/o ae	Total
reports w drug	a	b	a+b
reports w/o drug	c	d	c+d
Total	a+c	b+d	a+b+c+d

Measure of association	Formula	Probabilistic interpretation
Relative reporting (RR) ¹	$\frac{a(a+b+c+d)}{(a+c)(a+b)}$	$\frac{\Pr(\text{ae} \mid \text{drug})}{\Pr(\text{ae})}$
Proportional reporting rate ratio (PRR)	$\frac{a(c+d)}{c(a+b)}$	$\frac{\Pr(\text{ae} \mid \text{drug})}{\Pr(\text{ae} \mid \sim \text{drug})}$
Reporting odds ratio (ROR)	$\frac{ad}{cb}$	$\frac{\Pr(\text{ae} \mid \text{drug}) \Pr(\sim \text{ae} \mid \sim \text{drug})}{\Pr(\sim \text{ae} \mid \text{drug}) \Pr(\text{ae} \mid \sim \text{drug})}$
Information component (IC) ²	$\log_2 \frac{a(a+b+c+d)}{(a+c)(a+d)}$	$\log_2 \frac{\Pr(\text{ae} \mid \text{drug})}{\Pr(\text{ae})}$

1. The RR, when implemented within an empirical Bayesian framework, is known as empirical Bayes geometric mean (EBGM); 2. The IC is a logarithmic RR metric that is implemented in a Bayesian framework.

- Modern signal detection algorithms (e.g., EBGM, IC) could address sampling variance
 - Estimate confidence intervals (CIs) for disproportionality statistics
 - Dampen drug-event signals that have little evidence to support them
- How to address selection biases?

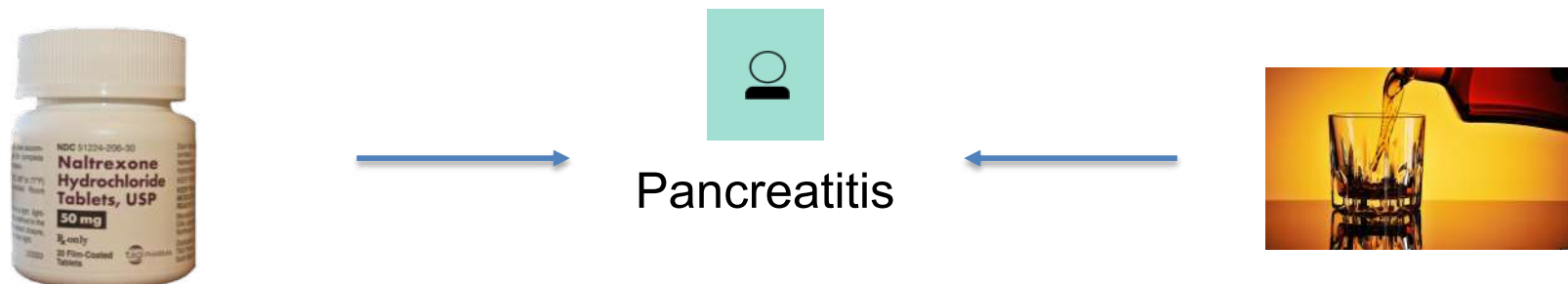
The Confounding Effect poses many challenges for ADR detection of real world events

Co-Prescription Confounders



Mary has arthritis, and has to take painkillers everyday. She has been taking both Aspirin and Vioxx. Which drug caused her heart attack?

Drug Indicator Confounders



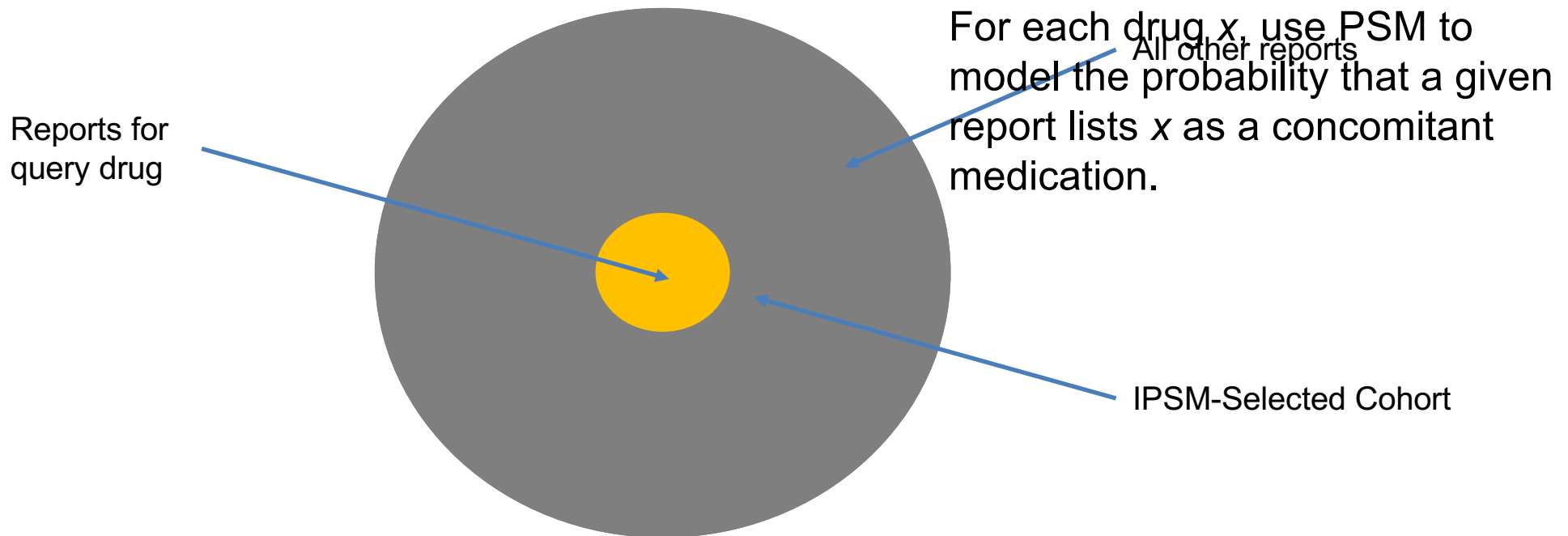
Joe is an alcoholic who develops Pancreatitis. He has been drinking daily and taking Naltrexone. What caused the Pancreatitis?

Selection biases in FAERS reports

- Selection biases introduce “synthetic associations”
 - (e.g.) from concomitant drug use (co-Rx effect)
 - drugs co-prescribed with **Vioxx** more likely to be associated with **heart attacks**
 - (e.g.) from indications (indication effect)
 - drugs given to **diabetics** more likely to be associated with **hyperglycemia**
 - (e.g.) co-Rx effect and indication effect extend to other covariates
 - Patients reported to be taking a cholesterol-lowering agent are more likely to be older, and this may cause these drugs to be synthetically associated with age-related effects, such as hypertension or myocardial infarction (age bias).
- Propensity score matching (PSM) corrects for bias of MEASURED covariates
 - Identify matched controls for the studied cases in observational clinical studies
 - Model the likelihood of a case being selected based on the covariates
 - $PS = \text{Estimated Pr(Exposed+ | covariates)} \sim \text{age} + \text{sex} + \text{weight} + \dots$
 - Match each case with one or more controls with the same likelihood
 - **However, PSM requires the covariates to be both known and measured; neither parameter is guaranteed to be present in FAERS**

Implicit Propensity Score Matching (IPSM)

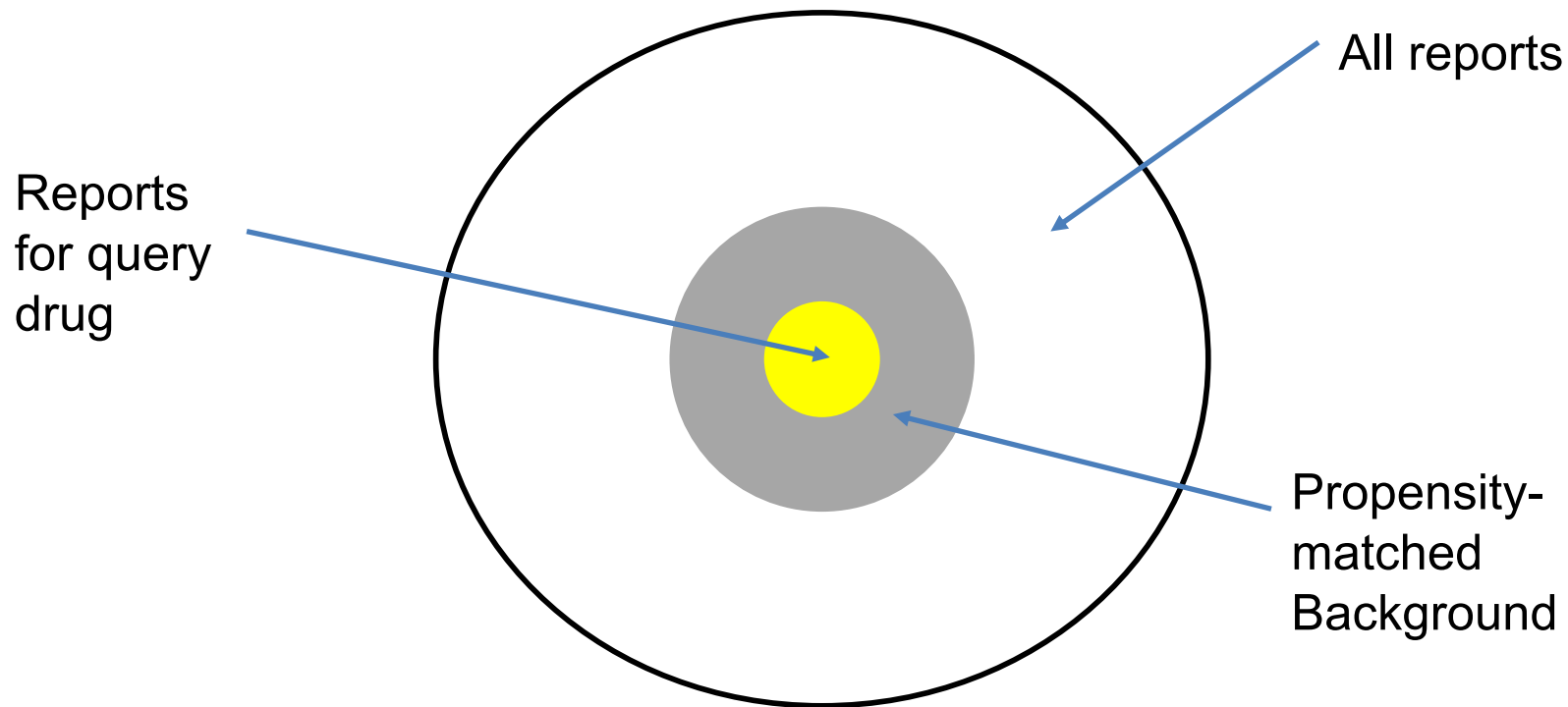
- Invented by Tatonetti NP et al. *Sci Transl Med*. 2012;4(125):125ra31.
- Assumes combination of co-reported drugs and co-indications describes all patient covariates. Generate a probability of a patient receiving a drug given co-prescribed medications and comorbidities.



- First, reduce to only those reports that have co-prescribed prescriptions listed
- Second, reduce to only those reports that have correlated indications listed

Takes advantage of co-Rx and indication variables likely to co-vary with unmeasured covariates

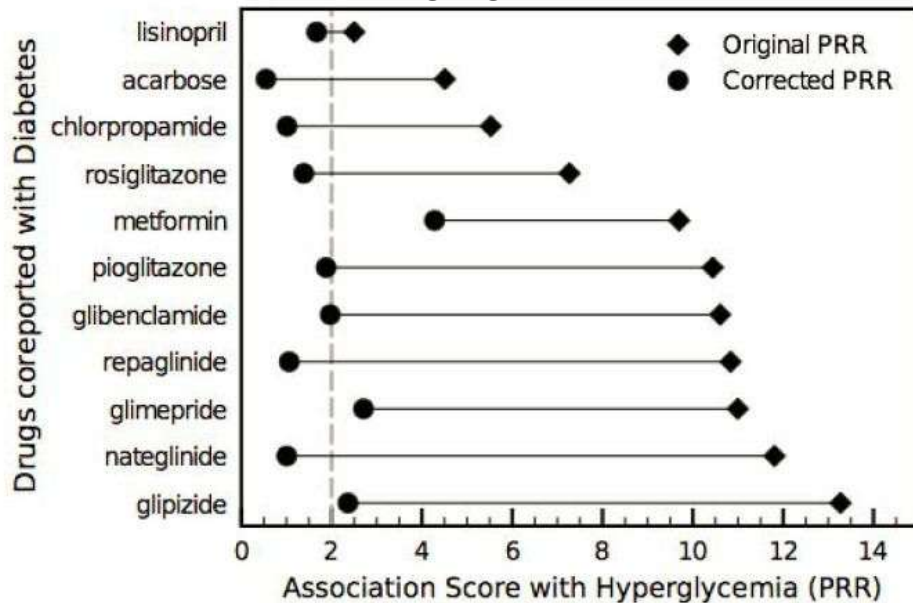
IPSM produces better estimates of expected values



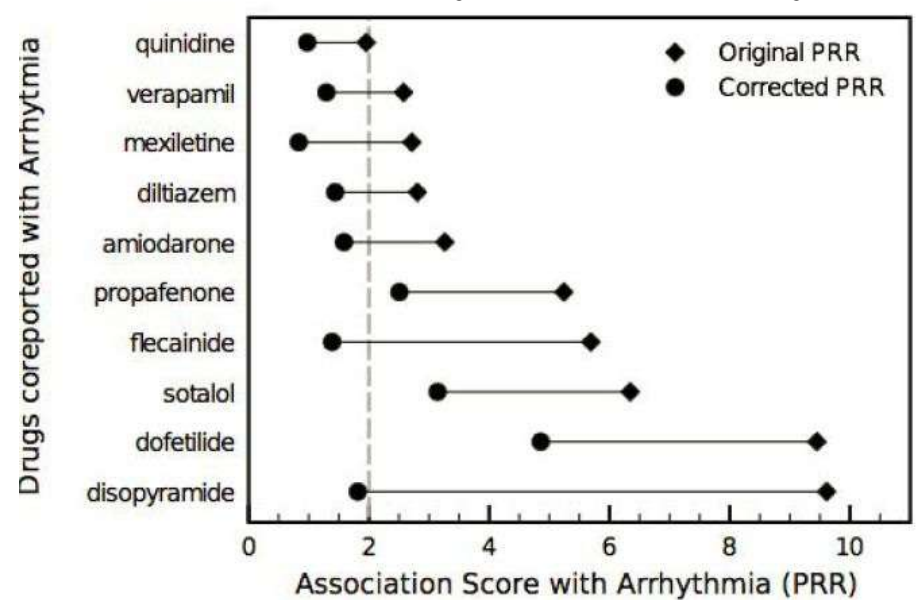
- Example: Reporting of **hyperglycemia** with **diabetes drugs**
- **Observed** reporting frequency: 17.7%
- **Expected** Estimates:
 - Entire database expected frequency: 1.5%
 - $PRR = 17.7\% / 1.5\% = 11.8!!!!$
 - IPSM-derived expected frequency: 17.6%
 - $PRR = 17.7\% / 17.6\% = 1.0 \dots$

IPSM corrects for indication and co-Rx biases

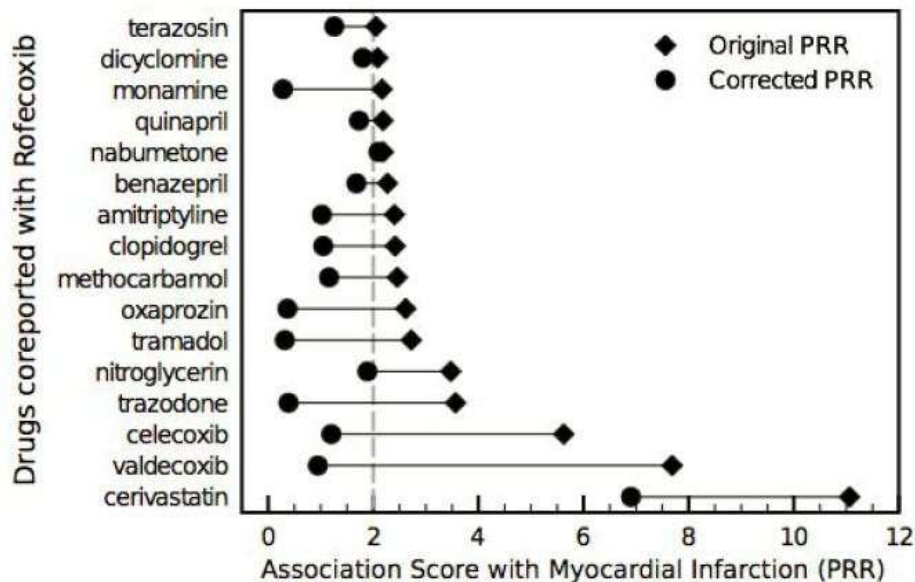
Drugs given to Diabetics



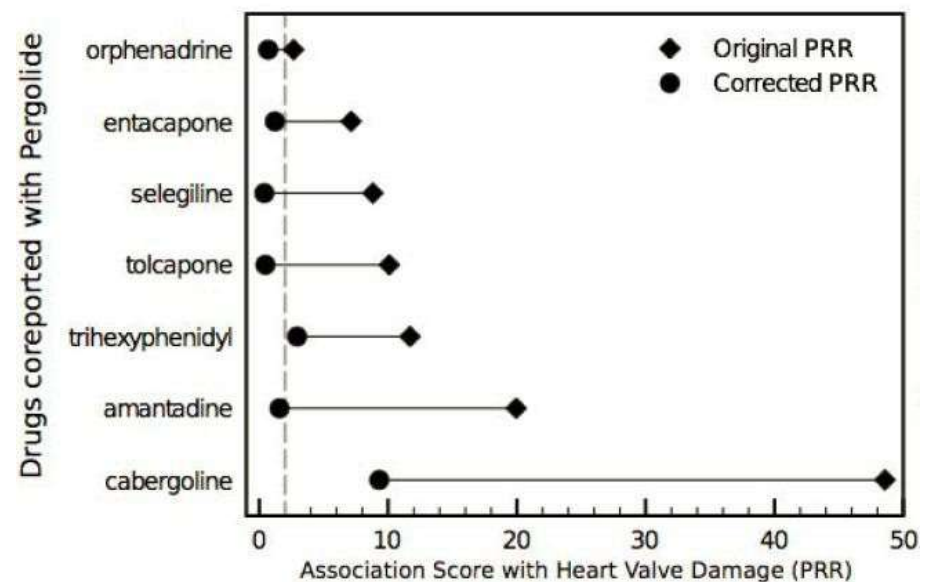
Anti-arrhythmics and Arrhythmia



Drugs co-reported with rofecoxib (Vioxx)

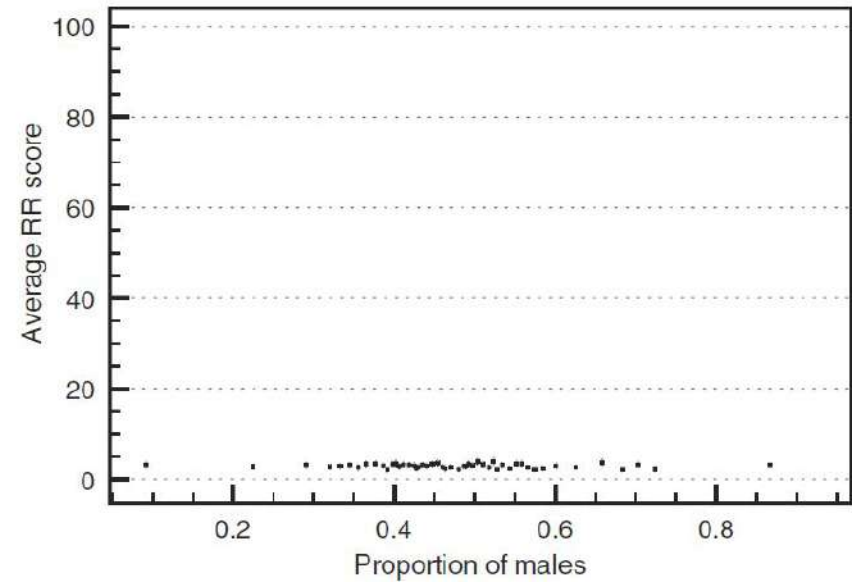
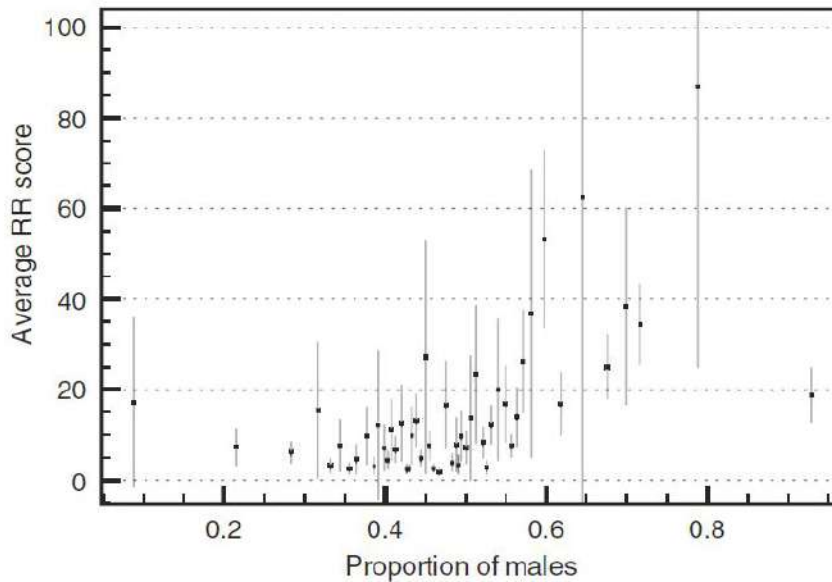


Drugs co-reported with pergolide

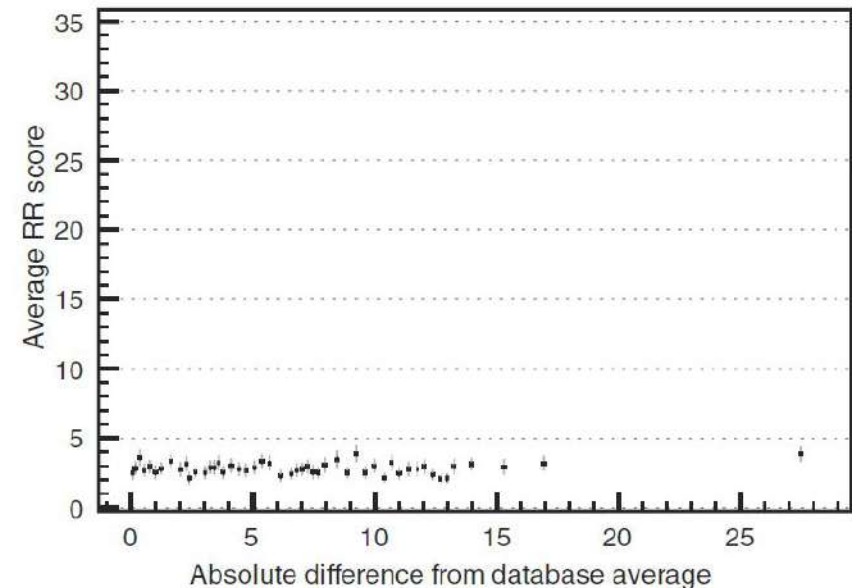
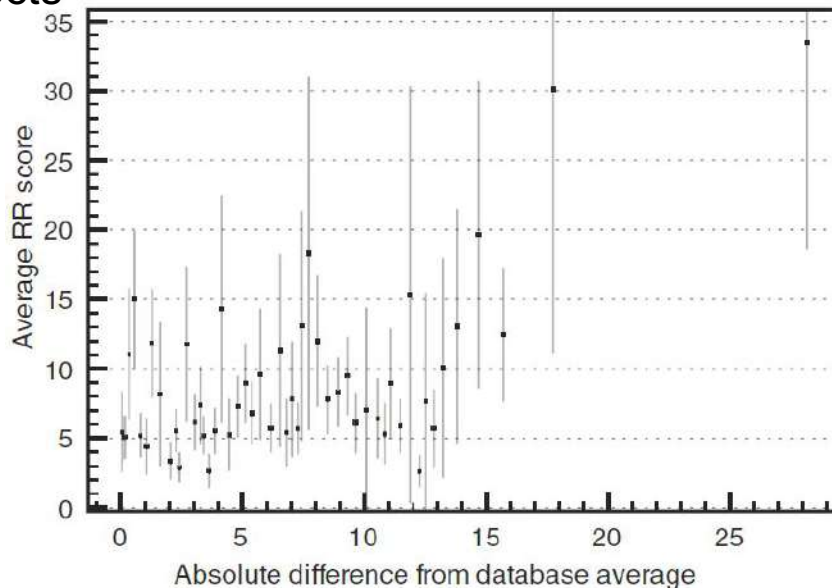


IPSM implicit correction for other biases

Drugs preferentially with males are more likely to be associated with 33 sex-related (male) effects



Drugs preferentially with young/old patients are more likely to be associated with 48 age-related effects



Performance of Pharmacovigilance Signal-Detection Algorithms for FAERS

- Data: FAERS data covered the period from 1968 through 2011 Q3, totaling 4,784,337 reports.

	Method name	Signal score computed
Disproportionality Analysis	Multi-item Gamma Poisson Shrinker (MGPS)	EBGM (empirical Bayes geometric mean): a centrality measure of the posterior distribution of the true observed-to-expected in the population EB05: lower 5th percentile of the posterior observed-to-expected distribution
	Proportional Reporting Ratio (PRR)	PRR: point estimate (mean) of the relative risk reporting ratio distribution PRR05: lower 5th percentile of the relative risk reporting ratio distribution
	Reporting Odds Ratio (ROR)	ROR: point estimate (mean) of the reporting odds ratio distribution ROR05: lower 5th percentile of the reporting odds ratio distribution
Multivariate Modeling	Logistic Regression (LR)	LR: point estimate of the odds ratio distribution obtained from logistic regression LR05: lower 5th percentile of the odds ratio distribution obtained from logistic regression
	Extended Logistic Regression (ELR)	ELR: point estimate of the odds ratio distribution obtained from extended logistic regression ELR05: lower 5th percentile of the odds ratio obtained from extended logistic regression

The application of biomedical gold standards

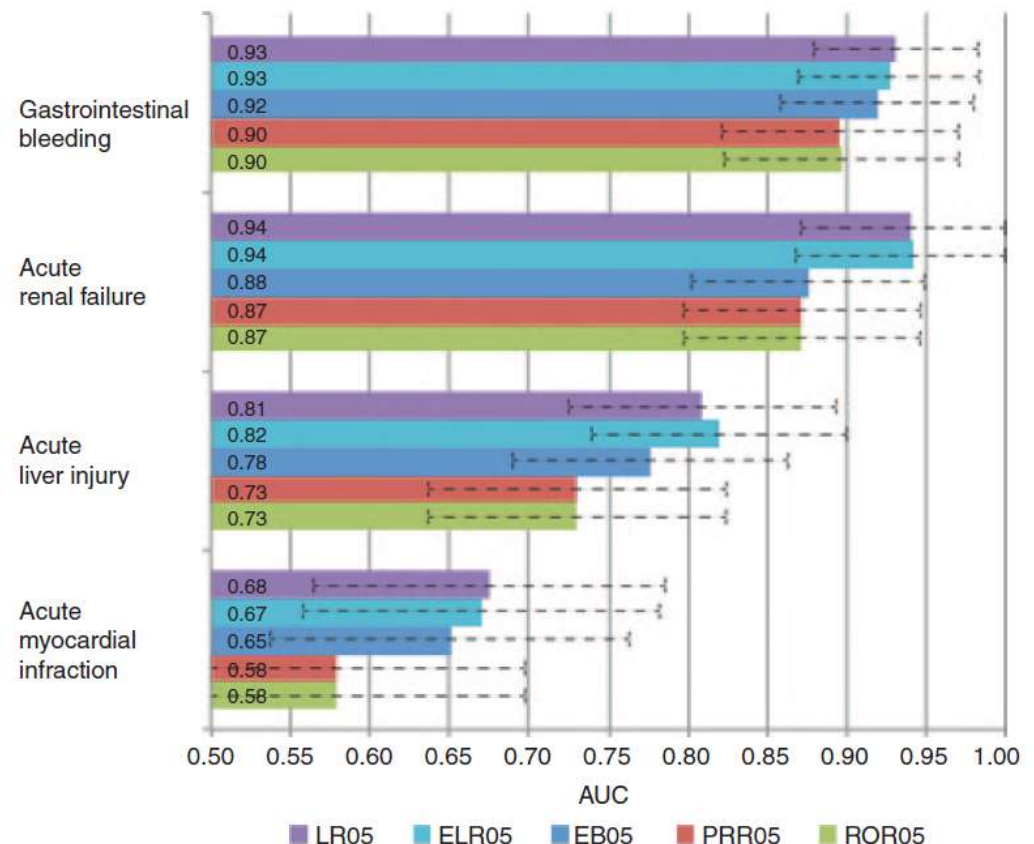
Positive Drug Set for an ADR:

- Event listed in Boxed Warning or Warnings/Precautions section of active FDA structured product label
- Drug listed as 'causative agent' in Tisdale et al, 2010: "Drug-Induced Diseases"[35]
- Literature review identified no powered studies with refuting evidence of effect









































Negative Set:

- Event not listed anywhere in any section of active FDA structured product label
- Drug not listed as 'causative agent' in Tisdale et al, 2010: "Drug-Induced Diseases"[35]
- Literature review identified no powered studies with evidence of potential positive association

Event	Positive Cases	Negative Case	Total
Gastrointestinal Bleeding	24	67	91
Acute Liver Injury	80	37	117
Acute Myocardial Infarction	36	66	102
Acute Renal Failure	24	64	88
Total	164	234	398



Summary - strengths and weaknesses of notable signal detection methods

	PRR	ROR	MGPS	BCPNN	LR
Simple to use					
Applicable to low event counts					
Easy to interpret					
Usable with SRS data					
Accounts for confounding factors					
Sensitivity					
Specificity					
					

Triaging to select signals and follow up

QUANTITATIVE “RULES”

- Apply fixed thresholds
 - $EB05 \geq 2$; $EBGM \geq 2$; $EBGM \geq 4$;
 - $PRR \geq 2$; a number of reports (N) ≥ 3 ; a Chi-square ≥ 4
 - Lower 95% CI of $PRR \geq 1$
 - Lower 95% CI of $ROR \geq 1$
 - $IC025 > 0$
- Apply flexible thresholds
 - Estimate the false discovery rate (FDR) to decide threshold on a signal-by-signal basis

QUALITATIVE “RULES”

- Novel
 - Not currently known and on drug label
 - New adverse event or new drug (“early warning”)
- High potential relevance
 - Public health issue – e.g. important drug (serious indication, widely used), serious reaction, many cases
 - Change in merit/harm
- Strong evidence
 - Exposure-response relationship (site, time-to-onset, dose, reversibility in dechallenge/rechallenge)
 - Reasonable from a biological mechanism perspective
- Time trend
 - Surge in recent reporting, notable increase in reporting over time

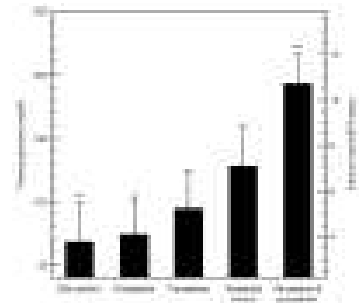
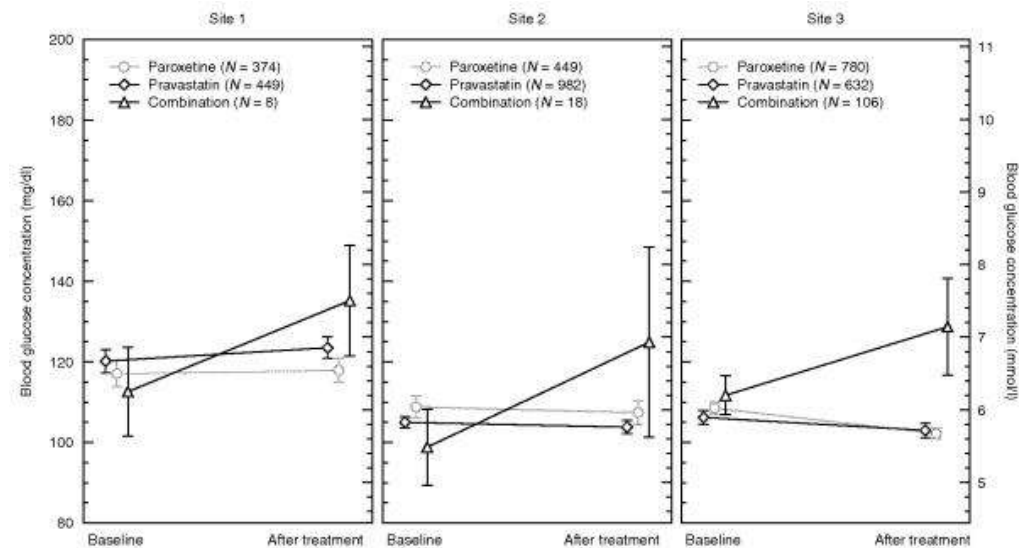
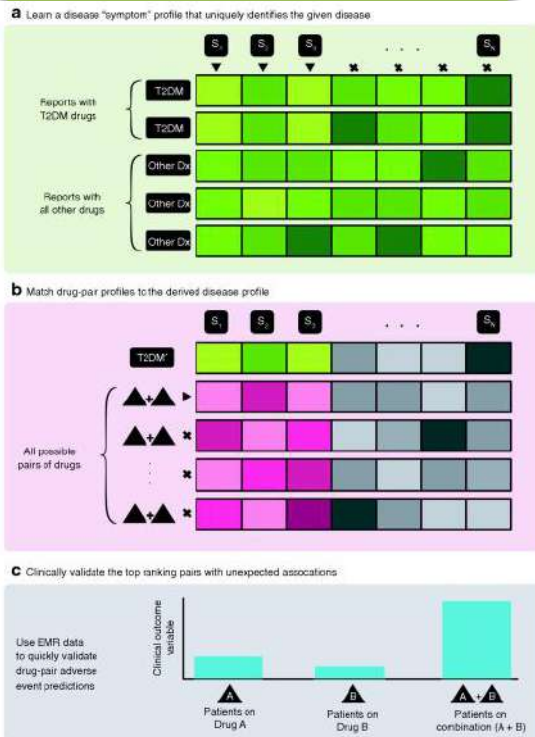
Beyond ADR detection

Common drug combo increases diabetes risk

Hypothesis generation based on FAERS

Signal validation based on EHR databases

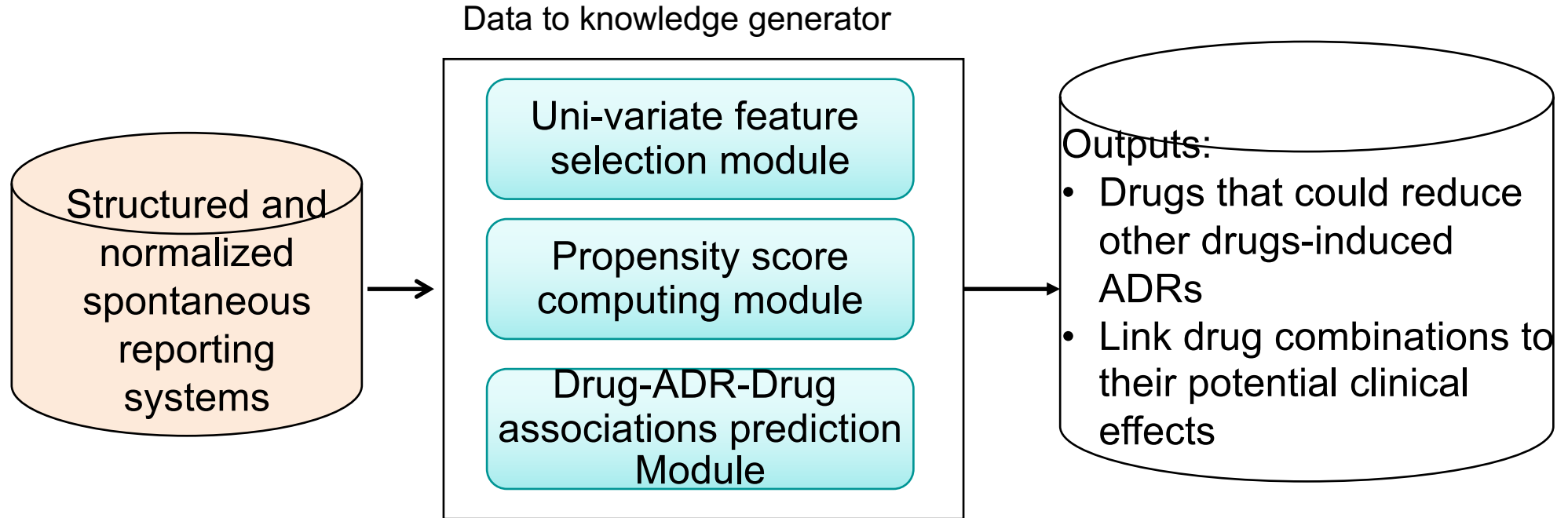
Mice model validation



A combination of two common drugs – paroxetine (one an antidepressant), pravastatin (the other used to lower blood cholesterol) – that caused blood sugar to rise, may put people at risk of developing diabetes.

Beyond ADR detection

Common drug combo decreases ADRs



$$\text{logit}(P(ADR = 1)) = \beta_0 + \beta_1 DrugA + \beta_2 P_1 + \beta_3 DrugB + \beta_4 P_2 + \beta_5 DrugA * DrugB + \lambda |\beta|_1$$

The novel regularized logistic regression is able to reveal two different mechanism of drug combinations

- $(\beta_3 + \beta_5)$: the degree that a patient who is on Drug A could benefit or suffer from taking Drug B for the ADR of interest
- β_5 : the degree that the interaction effect between Drug B and Drug A on the ADR

Clinical Validation

Pamidronate is used to treat high blood calcium levels

List of 15 predicted beneficial drug combinations and their ADR reduction

Drug A name	ADRs associated with drug A	Drug B name	Predicted beneficial score	Common ATC code	Evidence for combined use
benazepril	DIZZINESS	amlodipine besylate	-0.57	yes	F
atovaquone	PYREXIA	proguanil	-0.36	yes	F
rofecoxib	MYOCARDIAL INFARCTION MYOCARDIAL	pamidronate	-0.33	yes	
rosiglitazone	INFARCTION	exenatide	-0.32	yes	
progesterone	BREAST CANCER	adalimumab	-0.27	no	
trimethoprim	PYREXIA	sulfamethoxazole	-0.17	yes	F
exemestane	ARTHRALGIA	everolimus	-0.16	yes	III
amoxicillin	DIARRHOEA	clavulanic acid	-0.15	yes	IV
ampicillin	PYREXIA	sulbactam	-0.15	yes	F
desmopressin	HYPONATRAEMIA	somatropin	-0.15	yes	
sertraline	ANXIETY	nicotinic acids	-0.14	no	
sumatriptan	MIGRAINE	naproxen	-0.14	no	F
olanzapine	DIABETES	biperiden	-0.13	yes	
clindamycin	MELLITUS	benzoyl	-0.13	yes	F
fluticasone	DIARRHOEA	salmeterol	-0.13	yes	F
	DYSPNOEA				

F: FDA approved drug combination; III: phase III clinical trial; IV: phase IV clinical trial

a NSAID. On September 30, 2004, Merck withdrew rofecoxib from the market because of concerns about increased risk of heart attack and stroke associated with long-term, high-dosage use.

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Ginni Rometty
IBM Chairman, President and CEO
April 16, 2015



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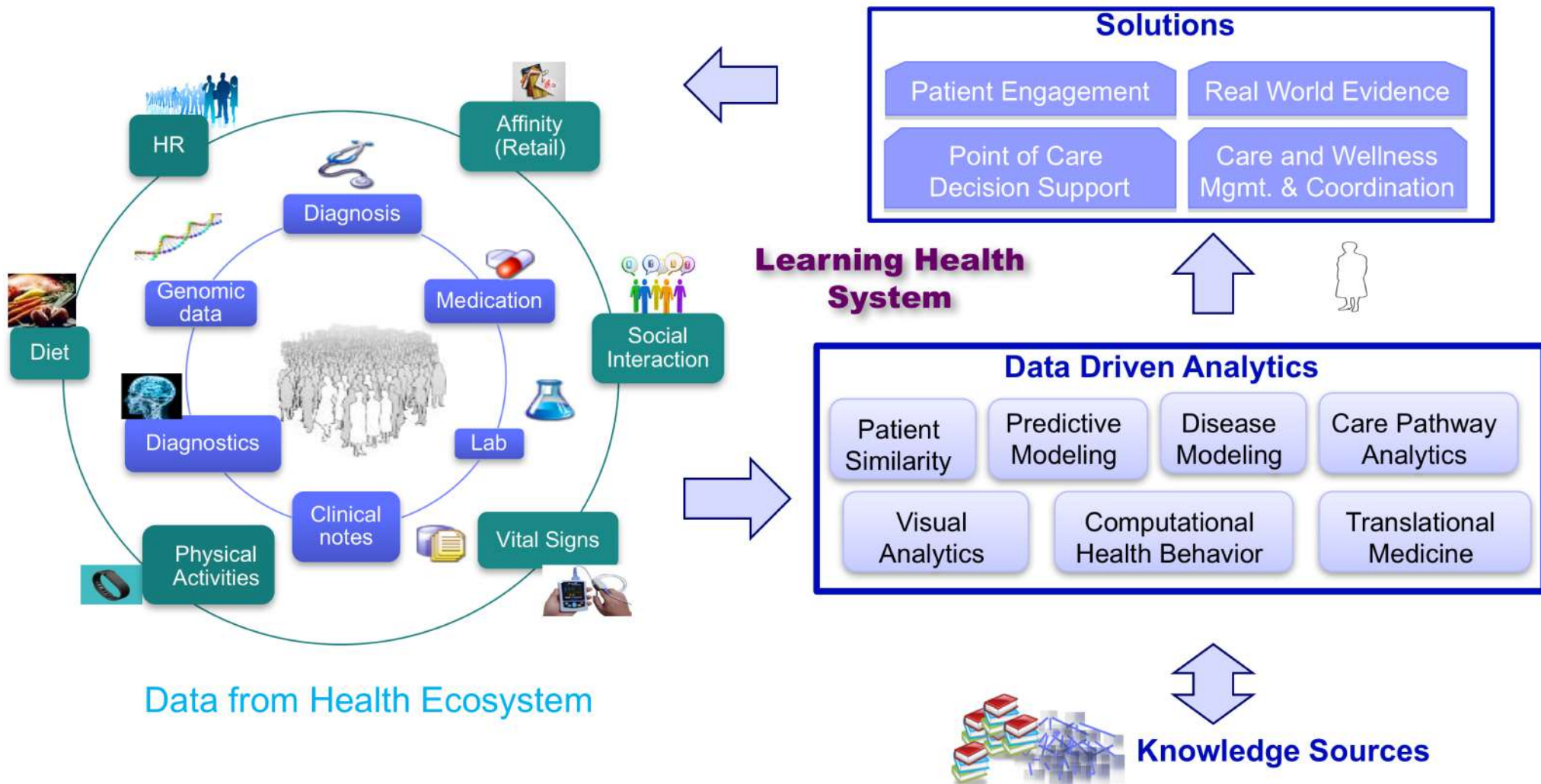
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