

Applying Semantic and Network Methods in AOP Knowledge Discovery

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**Adverse Outcome Pathways: From Research to Regulation
NIH Workshop, Bethesda MD, September 3-5 2014**

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Purpose of this presentation and discussion

A view of what is possible when we bring together the emerging science of AOP's, and state of the art in the computational techniques of data science, semantic technologies and network science

For technical details, see SOT presentation at <http://djwild.info>



The Usborne Book of the Future, 1979

Semantic technologies and AOP's – a new opportunity

- Our understanding of the effects of chemicals on our body is moving from a reductionist approach to a system, network approach
- The impacts of a chemical on the body are complex
 - Multiple targets, pathways
 - Indirect cascade effects
 - Phenotype and genotype dependent
- Semantic technologies fit this model well, as a way to handle big, complex, networked data sets from multiple sources
 - Applications in drug discovery, safety and chemical toxicity

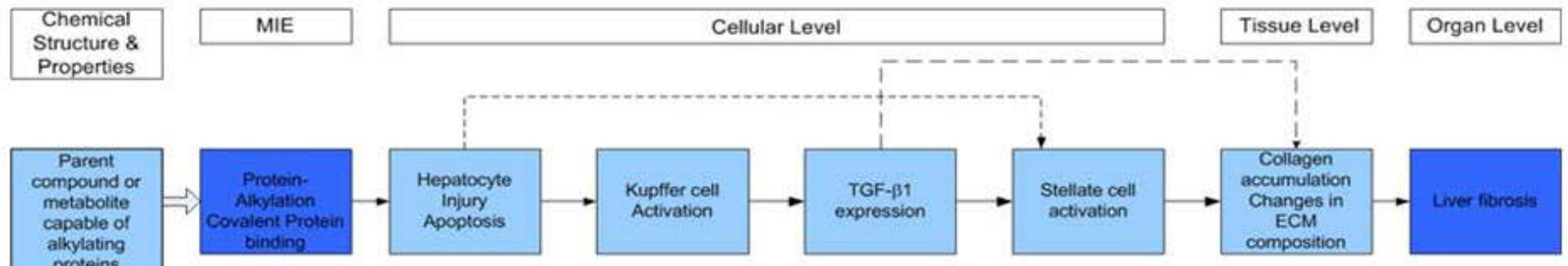
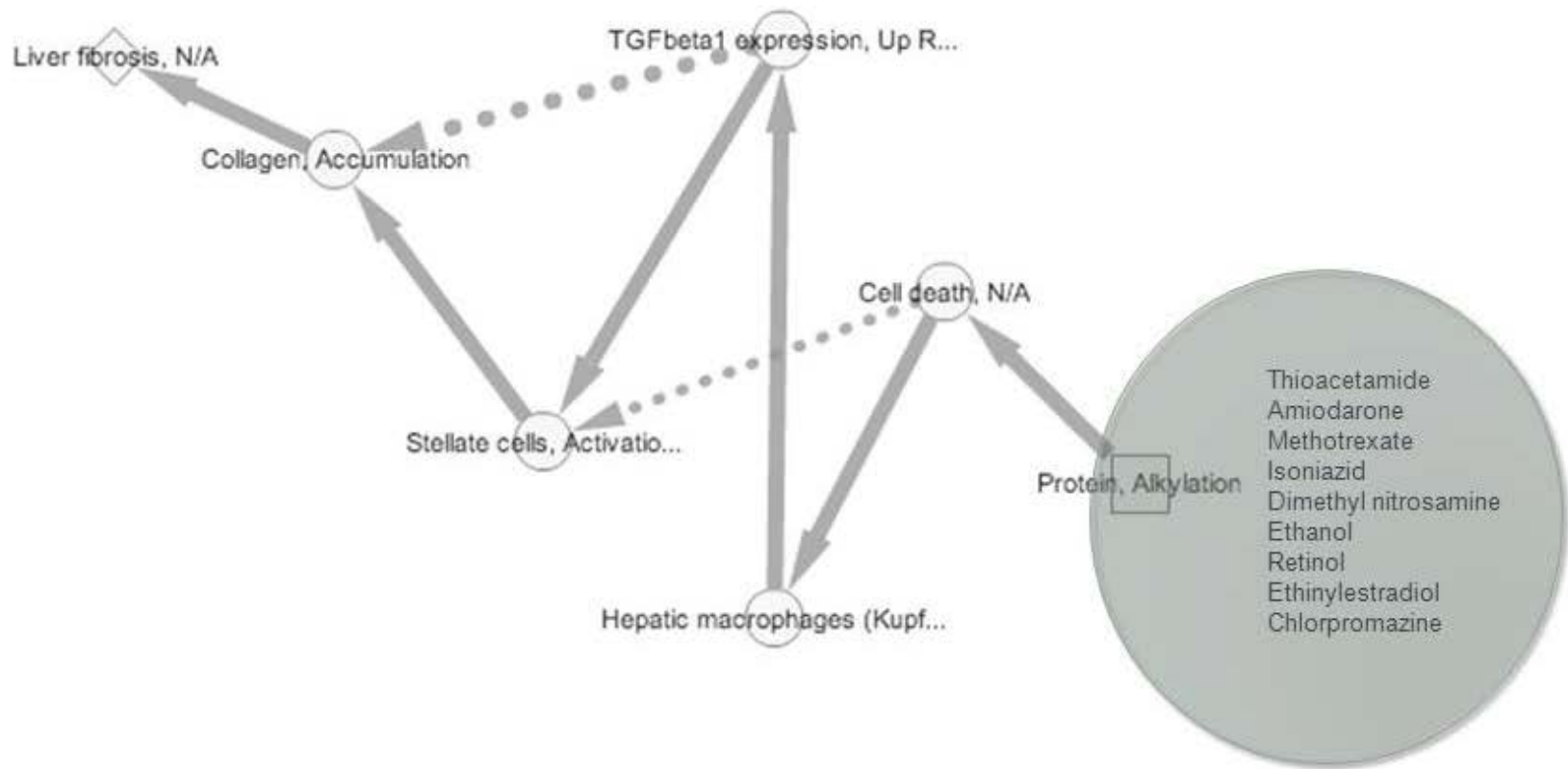
New “big” data approaches going mainstream in science

- NoSQL
 - Good for large amounts of **simple** or unstructured data
 - Very lightweight data structures e.g. tagging
- Semantic technologies
 - Good for large amounts of **complex** data
 - Represents data as networks rather than tables
 - Highly flexible in incorporating and linking many different kinds of data
 - Ontologies apply meaning to the data and relationships
 - Identified by Gartner as one of the top technology trends impacting information infrastructure in 2013:
<http://www.gartner.com/newsroom/id/2359715>
 - Now heavily used internally Google, Facebook, etc
 - Increasingly applied in scientific domains

Value proposition

- Semantic and network technologies could aid researchers in building AOP's and knowledge around AOP's
 - Predicting associations between compounds, targets and end points
 - Testing hypothesis
 - “Auto suggestion” of AOP associations
- Semantic and network technologies could help us apply established AOP's in problems like toxicity prediction
 - Profiling compounds across toxic end-points using computational representations of AOP's

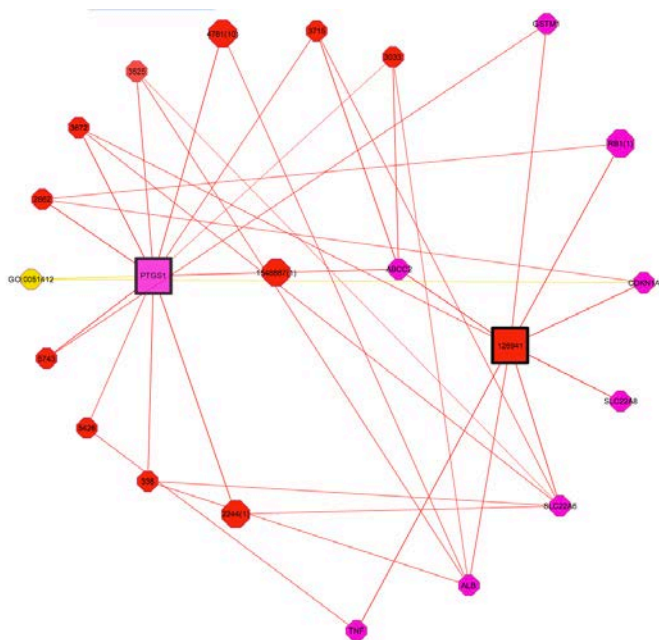
Example – Liver Fibrosis



Source: AOP Wiki

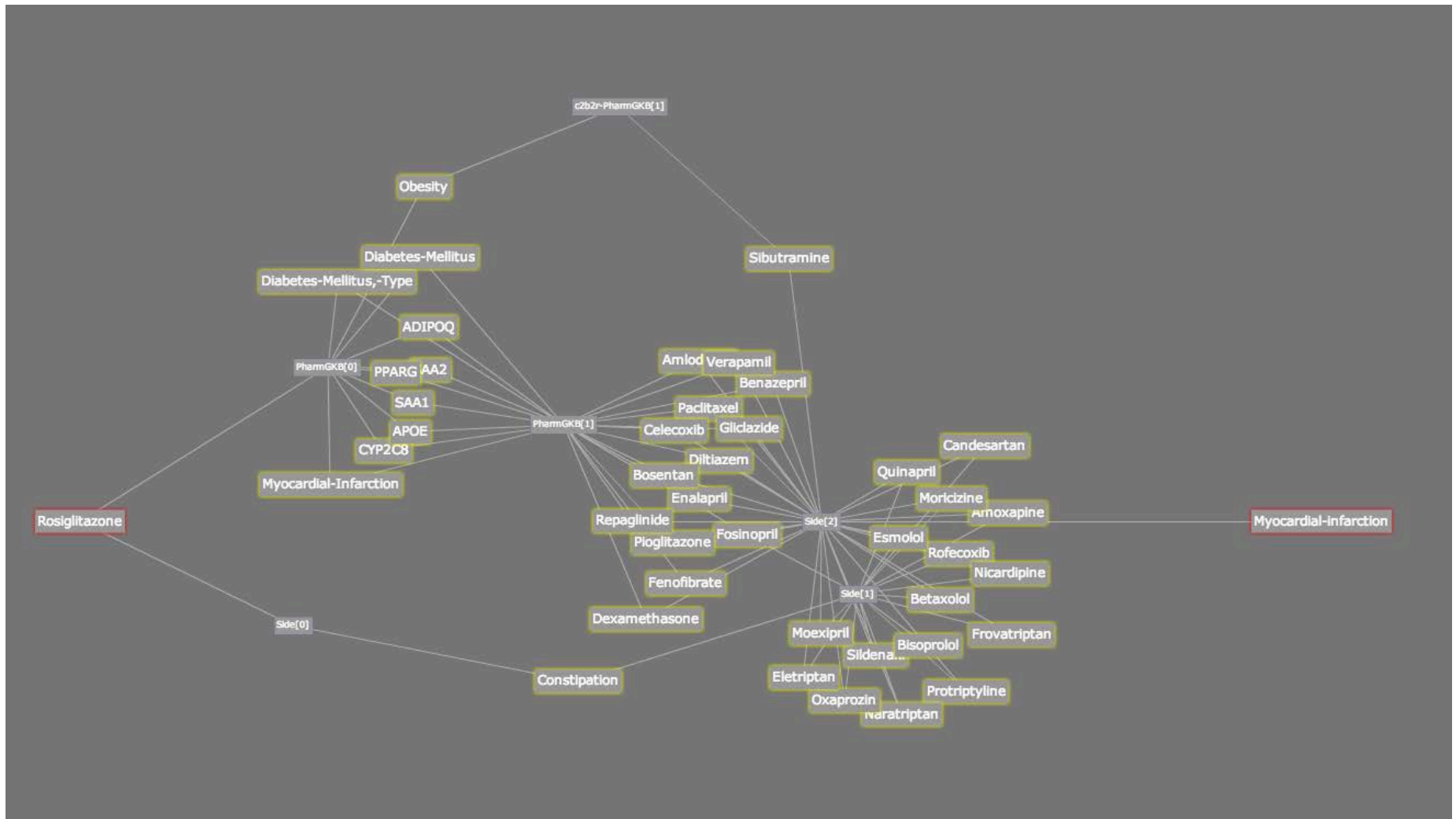
target	p value	score	type	chemohub
CYP2C9	3e-04	597.54	predicted	see paths
CYP2A6	0.002	266.26	predicted	see paths
CYP1A2	0.0027	228.29	predicted	see paths
HMOX1	0.0029	219.85	predicted	see paths
CYP2B6	0.0034	202.05	predicted	see paths
CYP3A4	0.0038	190.21	approved interaction	see paths
CYP17A1	0.0091	118.54	predicted	see paths
GSTA2	0.0432	43.14	approved expression	see paths
HMOX2	0.0744	28.28	approved expression	see paths
ABAT	0.1524	14.75	approved interaction	see paths
HPRT1	0.1983	11.16	approved expression	see paths
PLOD2	0.4368	3.81	approved expression	see paths

Isoniazid



PubChem CID	structure	Drug Name	Similarity	Related Diseases	ATC
3767		Isoniazid	1	Tuberculosis	J04AC01
3198		Econazole	0.915		D01AC03 G01AF05
5362440		Indinavir	0.894	HIV	J05AE02
6323497		Rifapentine	0.885	Tuberculosis	J04AB05
5342		Sulfapyrazole	0.873		M04AB02
4189		Miconazole	0.841		A01AB09 A07AC01 D01AC02 G01AF04 J02AB01 S02AA13
12560		Erythromycin	0.819		D10AF02 J01FA01 S01AA17
392622		Ritonavir	0.809	HIV Viral infection	J05AE03
2955		Dapsone	0.806	Leprosy	J04BA02
5281104		Paricalcitol	0.8	Hyperparathyroidism	A11CC07
4060		Mephenytoin	0.793	Epilepsy	N03AB04
55283		Itraconazole	0.783		J02AC02
5472		Ticlopidine	0.768	Stroke	B01AC05

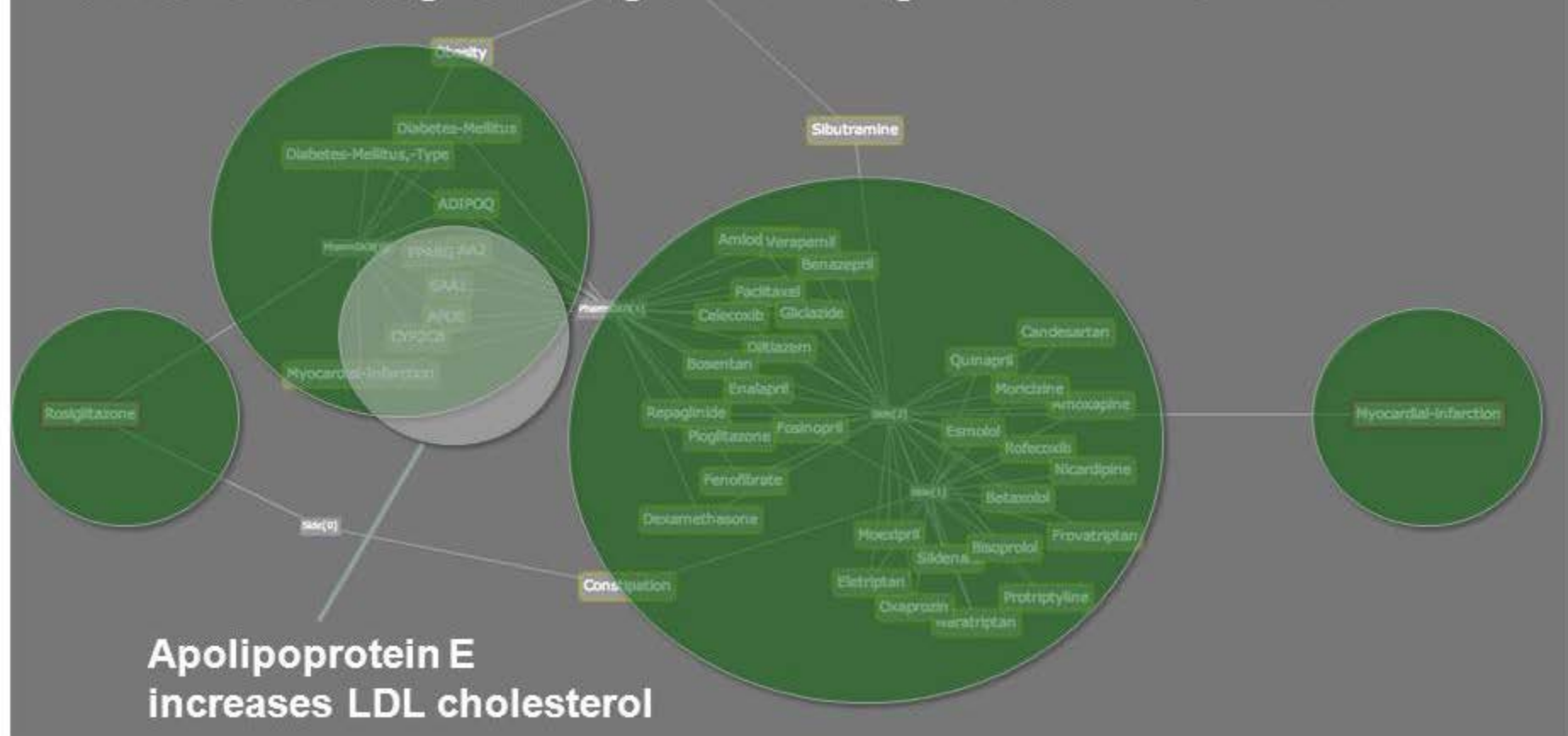
Association graph search – finding evidence paths



He, B., Tang, J., Ding, Y., Wang, H., Sun, Y., Shin, J.H., Chen, B., Moorthy, G., Qiu, J., Desai, P., Wild, D.J., **Mining relational paths in biomedical data** *PloS One*, 2011, e27506.

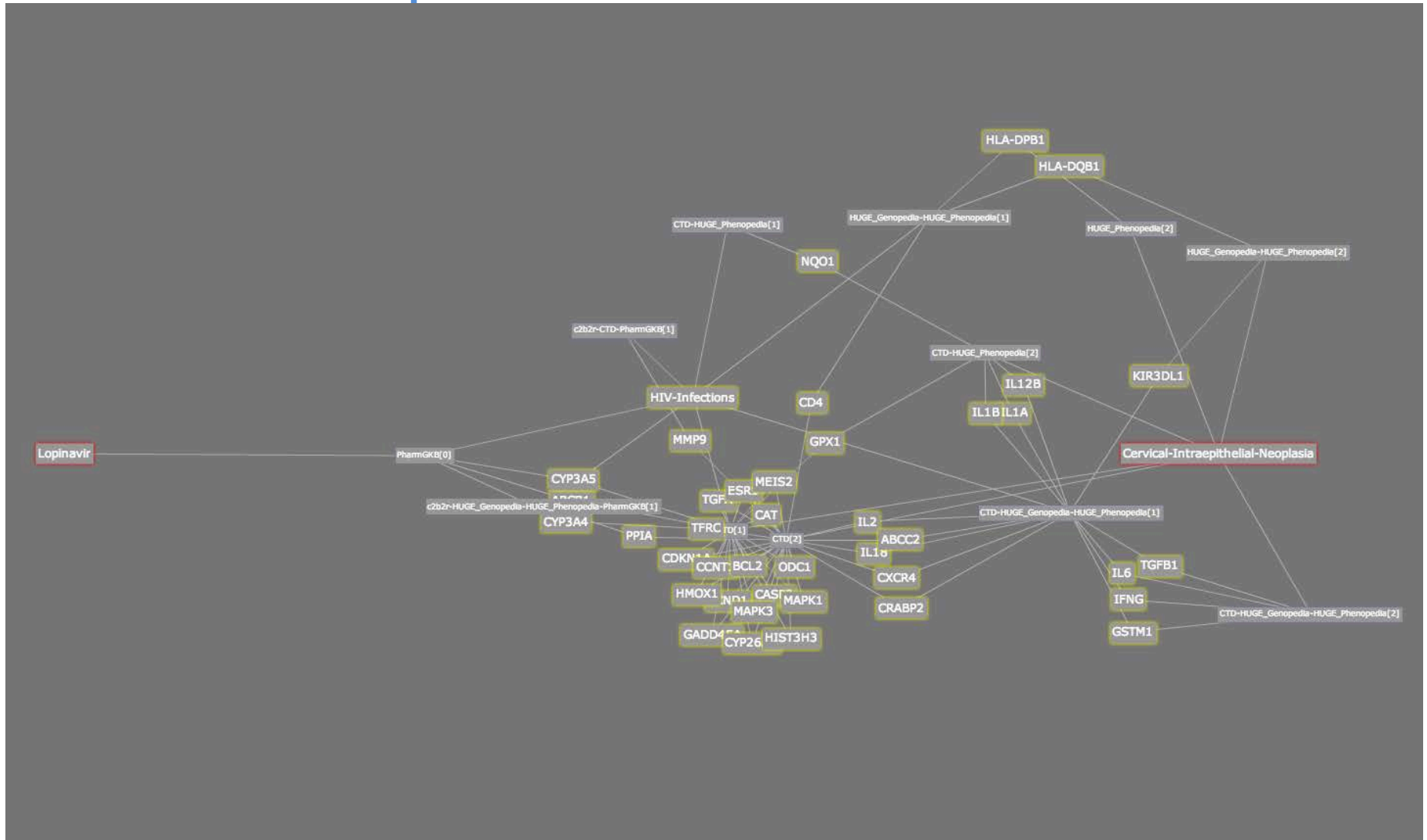
Association graph search – finding evidence paths

There exists a set of drugs with known MI side effect, that interact with a certain subset of genetic targets that Rosiglitazone also interacts with



He, B., Tang, J., Ding, Y., Wang, H., Sun, Y., Shin, J.H., Chen, B., Moorthy, G., Qiu, J., Desai, P., Wild, D.J., **Mining relational paths in biomedical data** *PloS One*, 2011, e27506.

Lopinavir – Cervical Cancer



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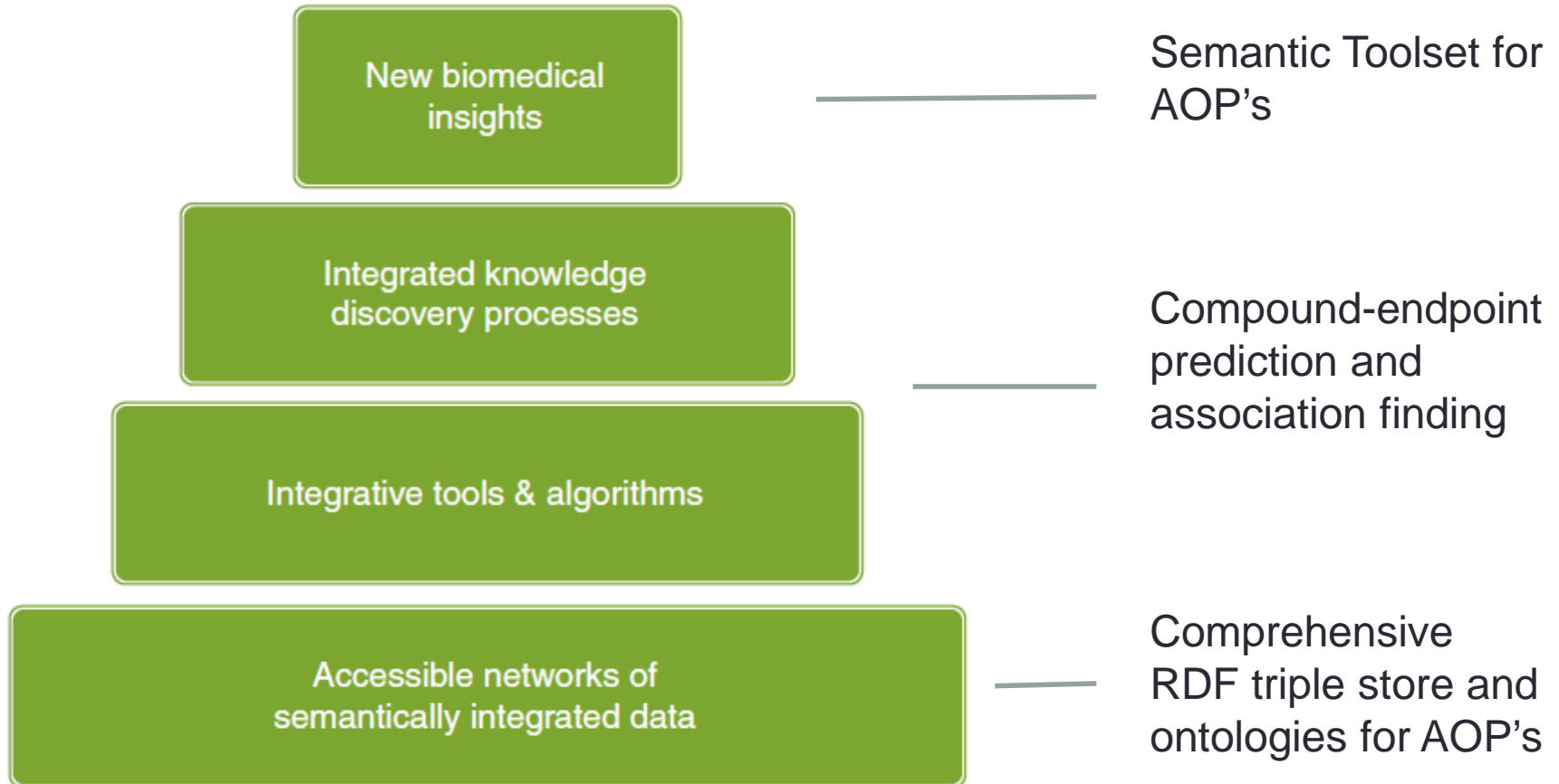
Application – Profiling Adverse Events

	hERG	Atrial Fib	Long QT	Tachycardia	Bradycardia	Cardiotoxicity	Cardiac failure	Cardiomyop.	Myocard. Inf.	Tachycardia	Cardiotoxicity	Long QT	Atrial Fib	hERG	Long QT	Tachycardia	Cardiotoxicity	Bradycardia	Tachycardia	Tachycardia	Cardiotoxicity	Tachycardia	Atrial Fib
A10366245		Orange			Red				Orange						Orange								
A10366585		Red			Red							Orange					Red			Red			
A49585949							Red															Red	
A48480494		Orange		Orange		Orange									Orange			Orange				Orange	
Aspirin								Red				Orange	Orange						Orange				
Rosuvastatin					Red		Orange																
Pioglitazone					Orange							Orange		Orange			Orange						
AGGREGATE		Red		Orange	Red	Orange	Red	Red	Orange			Orange	Orange	Orange	Orange	Orange	Red	Orange	Orange	Red	Orange	Red	

Why is semantic data powerful?

- Breaking down data and domain silos
 - Chemistry – biology – toxicology – adverse event - endpoint
 - Molecular – patient
 - Public – commercial – proprietary
- Easy to repurpose existing and harvest new data
 - RDF format is standard
 - Separation of the data from the structure of the data
- Semantic networks -> biological networks
 - Systems chemical biology / network biology
 - Move away from naïve drug/target or target/endpoint
 - Hugely powerful algorithms in networking community
 - Prediction, hypothesis testing, interpretation

Proposal: Semantic Toolkit for AOPs



Comprehensive semantic store for AOPs

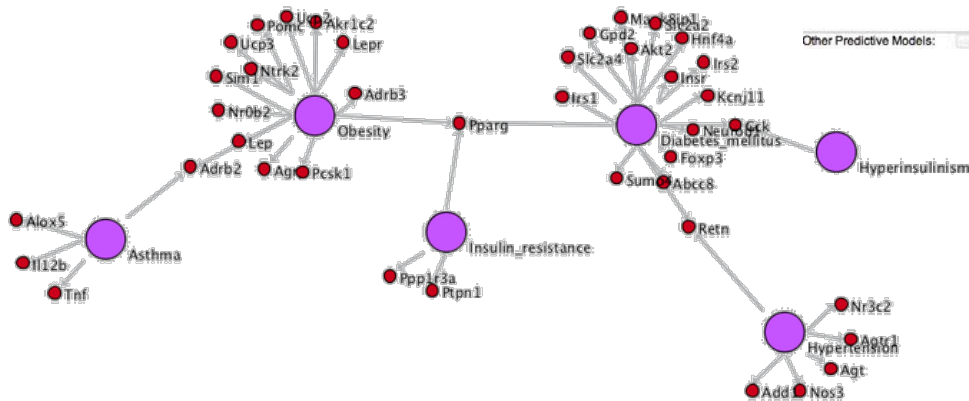
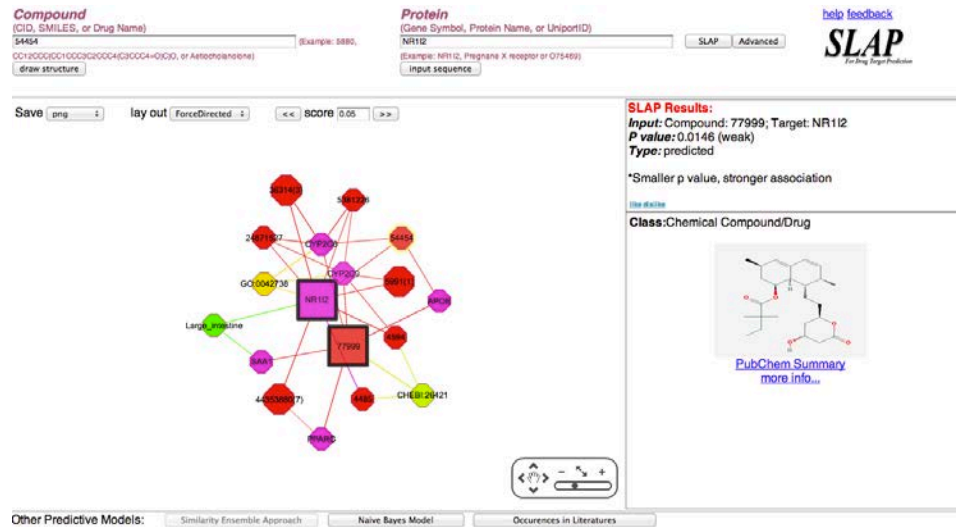
- Contains all public data of relevance, from compound to organism. As a start...
 - **OnTop***: PubMed, GO, KEGG, MeSH, NCI, UniProt, Entrez Gene, NCBO, CTD, ACToR, ToxRefDB, ToxMiner, ToxCat
 - **Chem2Bio2RDF/Chem2Bio2OWL**: 52 public datasets relating to compounds, genes, pathways, diseases and side effects
 - Other relevant sets – e.g. FDAERS, social media
- Ontologically mapped to concepts in AOPWiki
- SPARQL endpoint for searching

* Ontology for modeling adverse outcome pathways: semantic tools for systems tox. Imran Shah, EPA-NIEHS Advancing Environmental Health Data Sharing and Analysis: Finding a Common Language. June 25, 2013

Compound-Endpoint prediction & association finding

- Predicting compound-endpoint associations with SLAP
 - Modified version of current compound-target algorithm
 - Association score and p-value
- Automatic generation of preliminary AOP networks
 - Using SLAP significant subnetwork between compound and endpoint
 - “starting point” for understanding potential AOPs
- Generation of literature supported association networks
 - More open-ended association finding and visualization
- Random-Walk methods
 - Most recent research at IU

- Open toolset made available in association with AOP KB and integrated with other tools
- AOP prediction, exploration, hypothesis testing & application



Summary

- Semantic technologies becoming mainstream for big / complex data problems; increasing applications in science
- IU and EPA have demonstrated applicability of semantic technologies in chemical, biological data and for AOP's
- AOP's map particularly well onto the semantic approach
- Huge potential is realized when network and predictive algorithms are applied – the “semantic stack”
- Direct opportunity to engage semantic technologies in the emerging AOP KB / AOPWiki projects