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ImmunoX

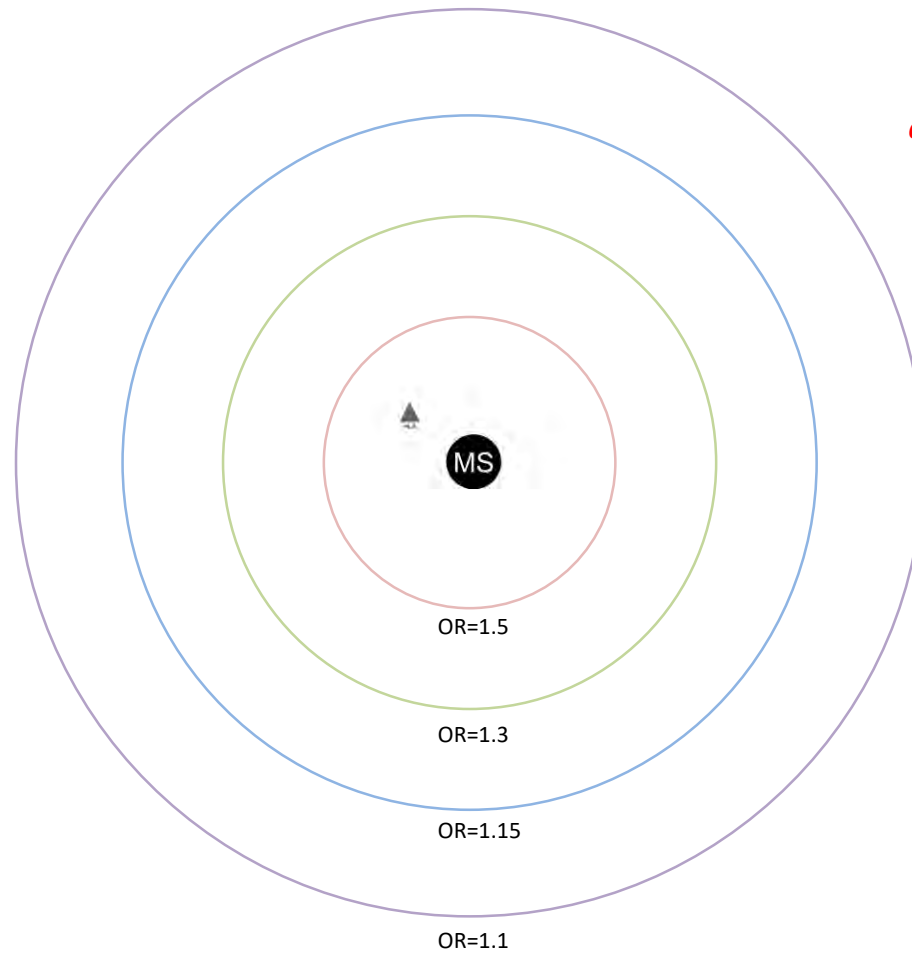
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University of California San Francisco

# The value (and challenges) of shared work

- International multiple sclerosis genetics consortium
- MultipleMS Consortium
- BRAVEinMS Consortium
- International MS microbiome study

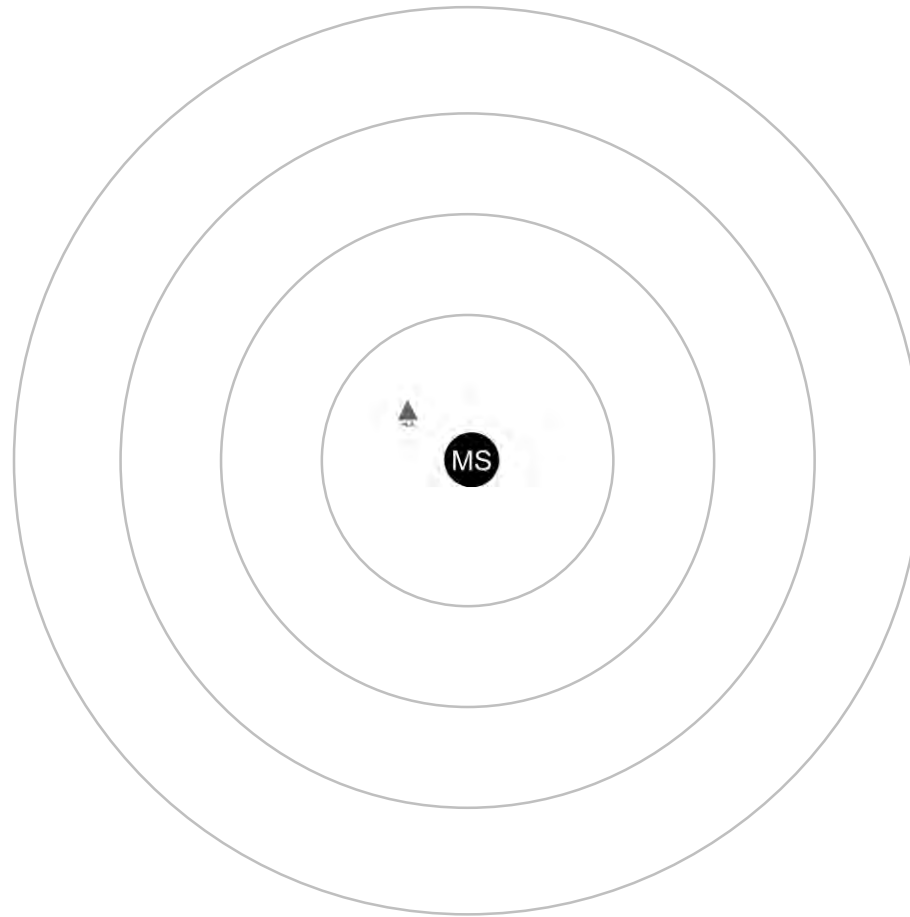
# The Genetic Landscape of MS



**1972**  
*First reported  
association between MS  
and HLA*

Jersild et al. *Lancet* 1:1240, 1972

# The Genetic Landscape of MS



**1996**

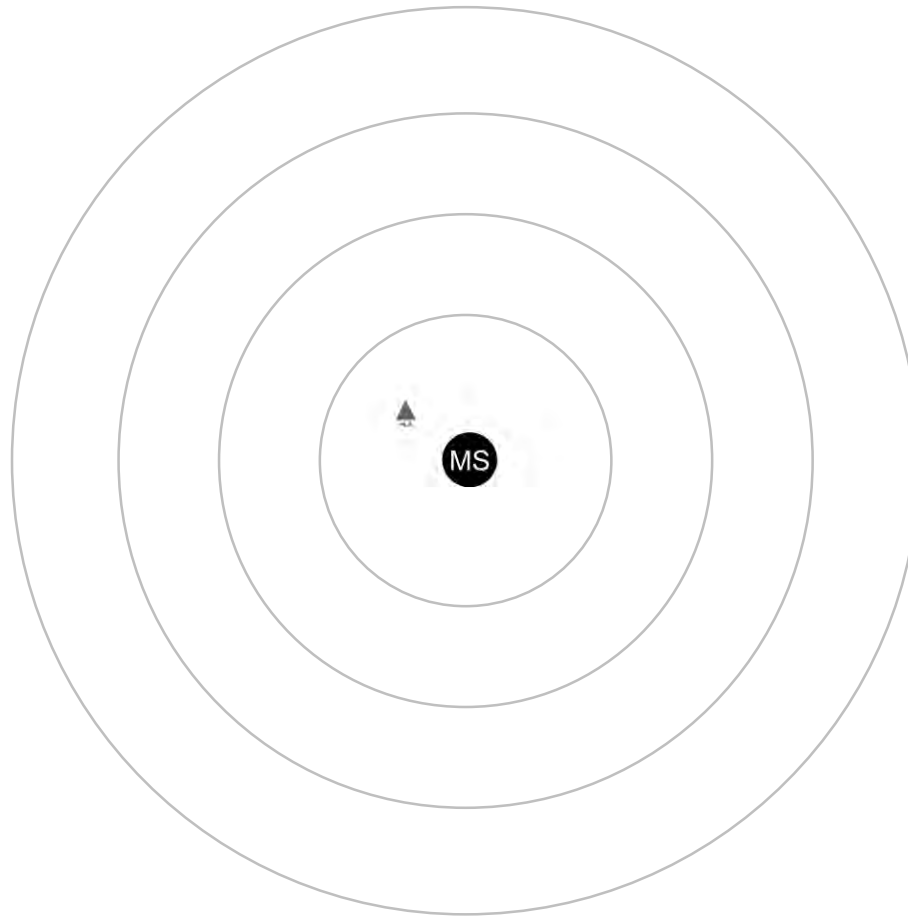
*First generation  
genome-wide linkage  
studies  
(400 markers)*

Sawcer et al. *Nat Genet* 13:464, 1996

Haines et al. *Nat Genet* 13:469, 1996

Ebers et al. *Nat Genet* 13:472, 1996

# The Genetic Landscape of MS



**2005**

*Second generation  
genome-wide linkage  
study (5000 markers)*

Sawcer et al. *Am J Hum Genet* 77:454, 2005

Incidence (per 100,000 inhabitants)

More than 45

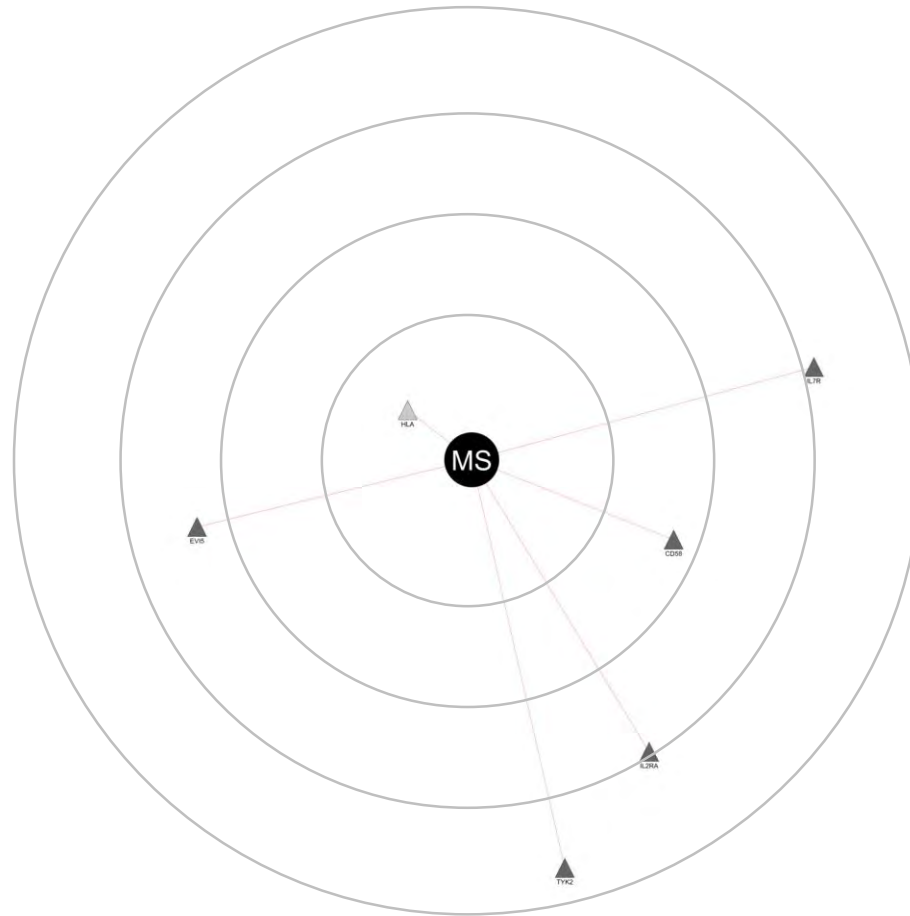
Less than 15

45

15

Group photo of researchers

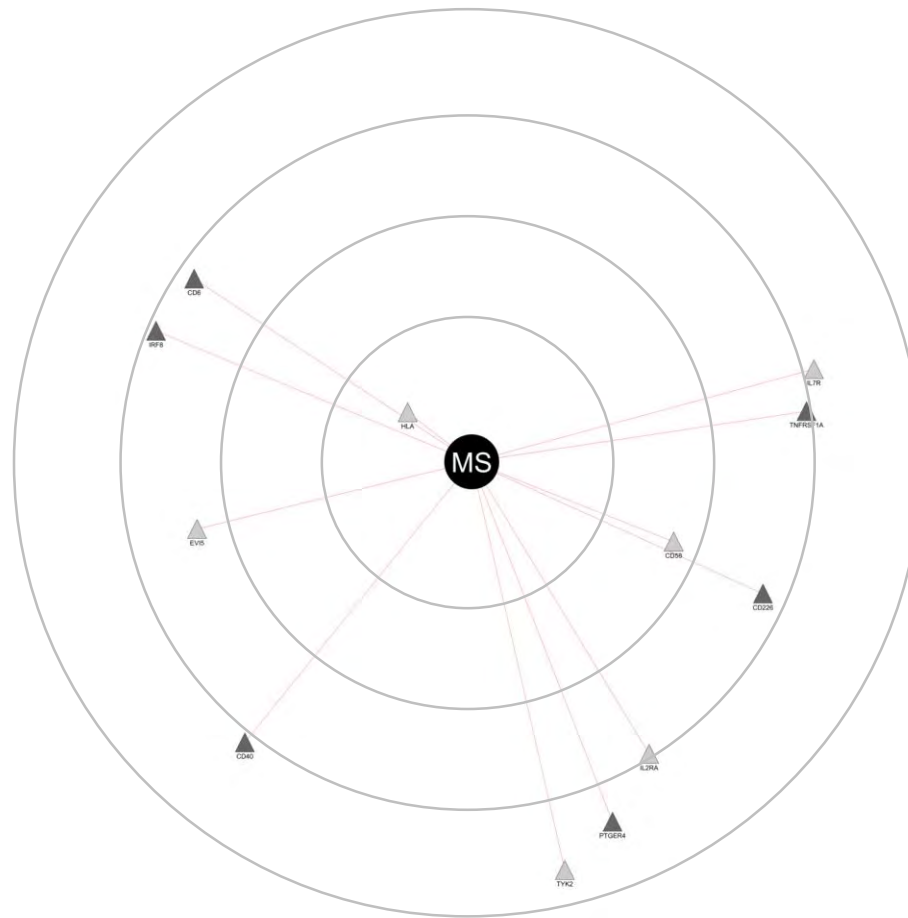
# The Genetic Landscape of MS



**2007**  
***First generation GWAS***  
***(1000 patients)***

IMSGC. *N Engl J Med* 357:851, 2007

# The Genetic Landscape of MS



**2009**  
*Meta-analysis of GWAS*

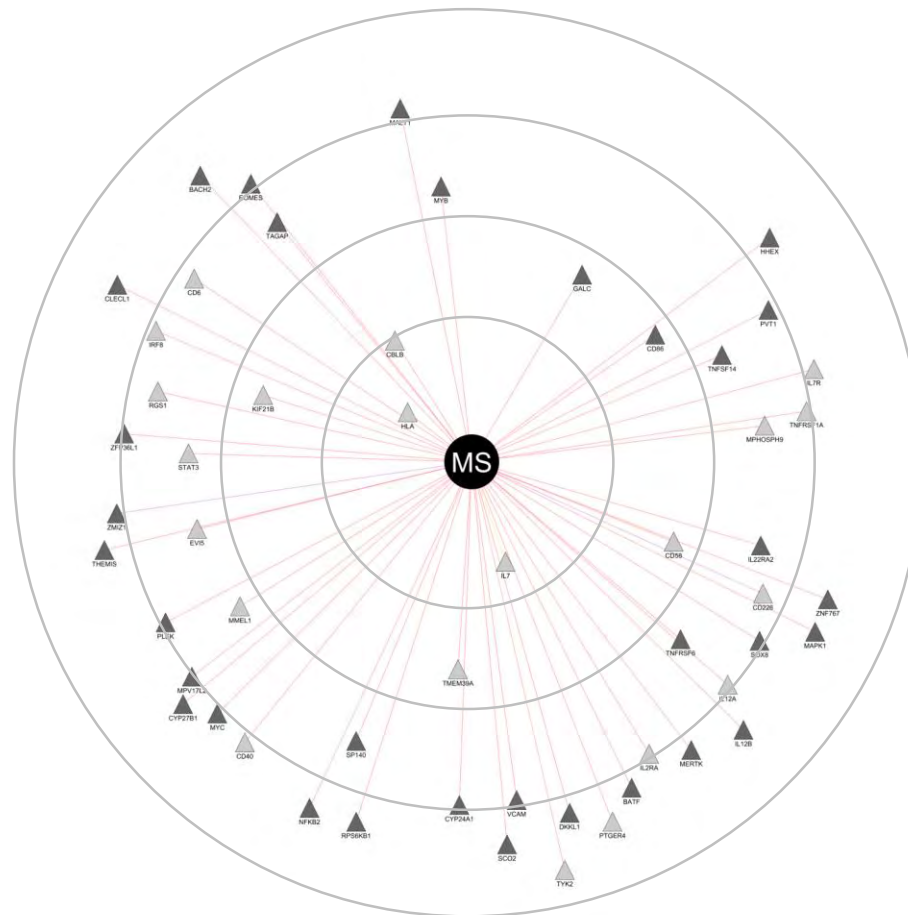
IMSGC. *Genes Immun* 10:11, 2009  
DeJager et al. *Nat Genet* 41: 776 , 2009



## Whole genome sequencing of MS twins

Ban et al. *Genes Immun* 11:660, 2010  
Zuvich et al. *Hum Genet* 127:525, 2010  
IMSGC. *Hum Mol Genet* 19:953, 2010  
IMSGC. *Nat Genet* 42:469, 2010

# The Genetic Landscape of MS



**2011**  
**Second generation**  
**GWAS**  
**(10,000 patients),**  
**GWAS Follow-ups**

LETTER

Genetic risk and a primary role for cell-mediated immune mechanisms in multiple sclerosis

The International Multiple Sclerosis Genetics Consortium\* & the Wellcome Trust Case Control Consortium 2\*

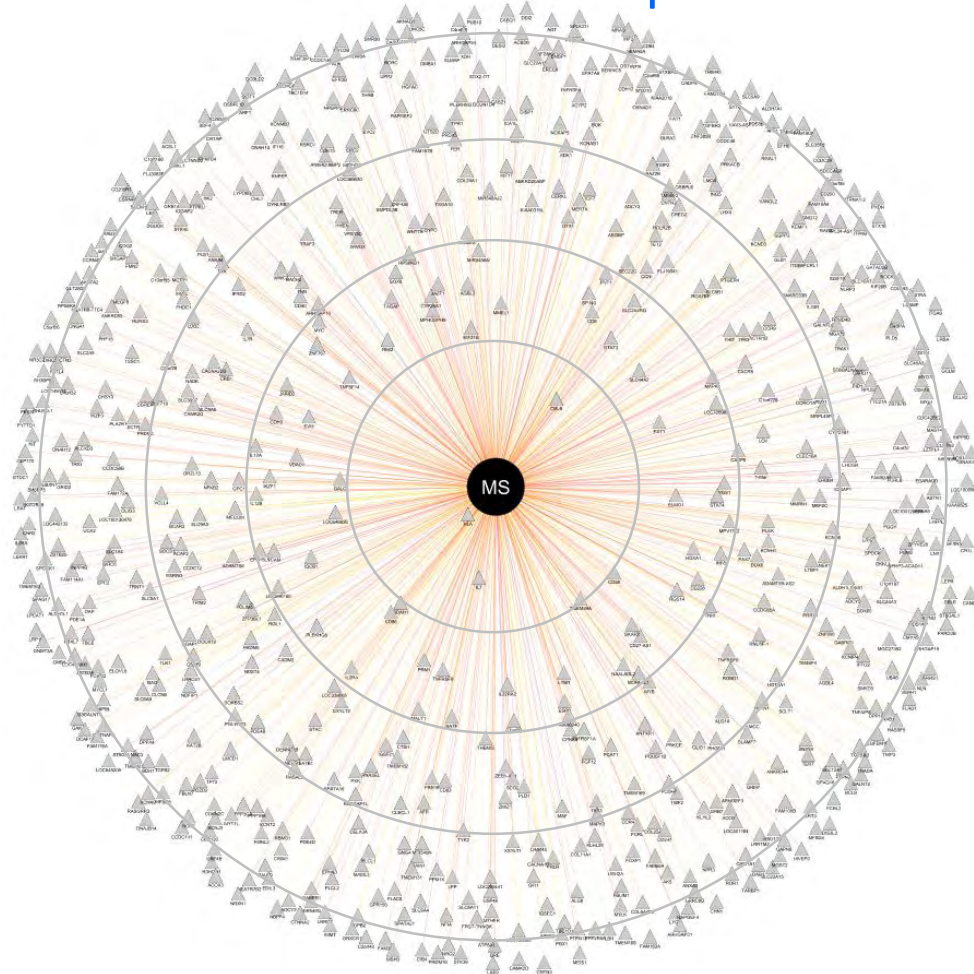
doi:10.1038/nature10251

ImmunoChip  
(>15,000 patients)

### Analysis of immune-related loci identifies 48 new susceptibility variants for multiple sclerosis

A full list of authors and affiliations appears at the end of the article.

# The Genetic Landscape of MS



**2019**

Meta-analysis 3.0/MSchip  
(>40,000 patients)

IMSGC. Science. In revision, 2018

# Public databases

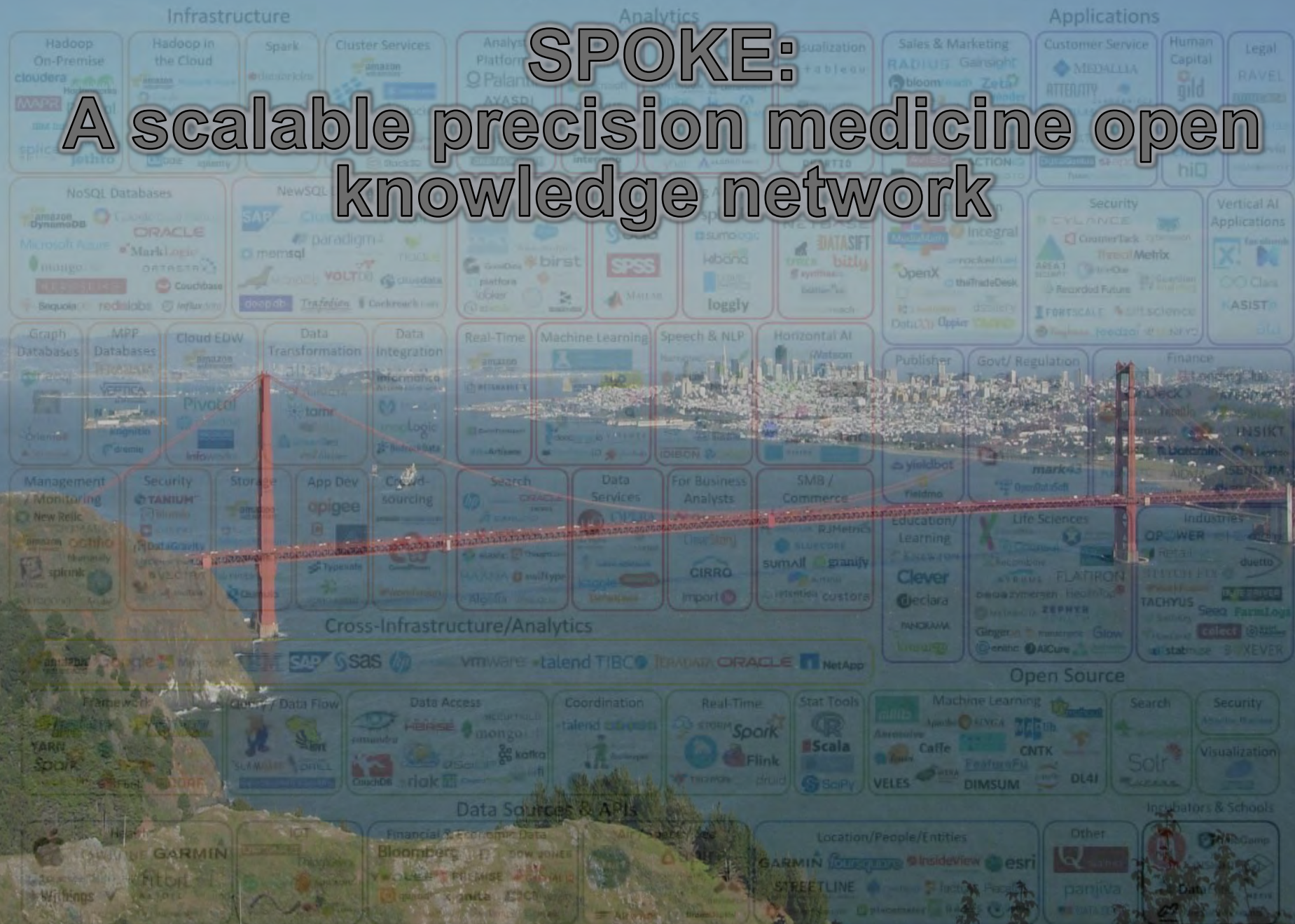
## ...Really?

- SwissProt
- GWAS Catalog
- ChEMBL
- Diseases
- Drugbank
- KEGG
- STRING
- SIDER



# SPOKE:

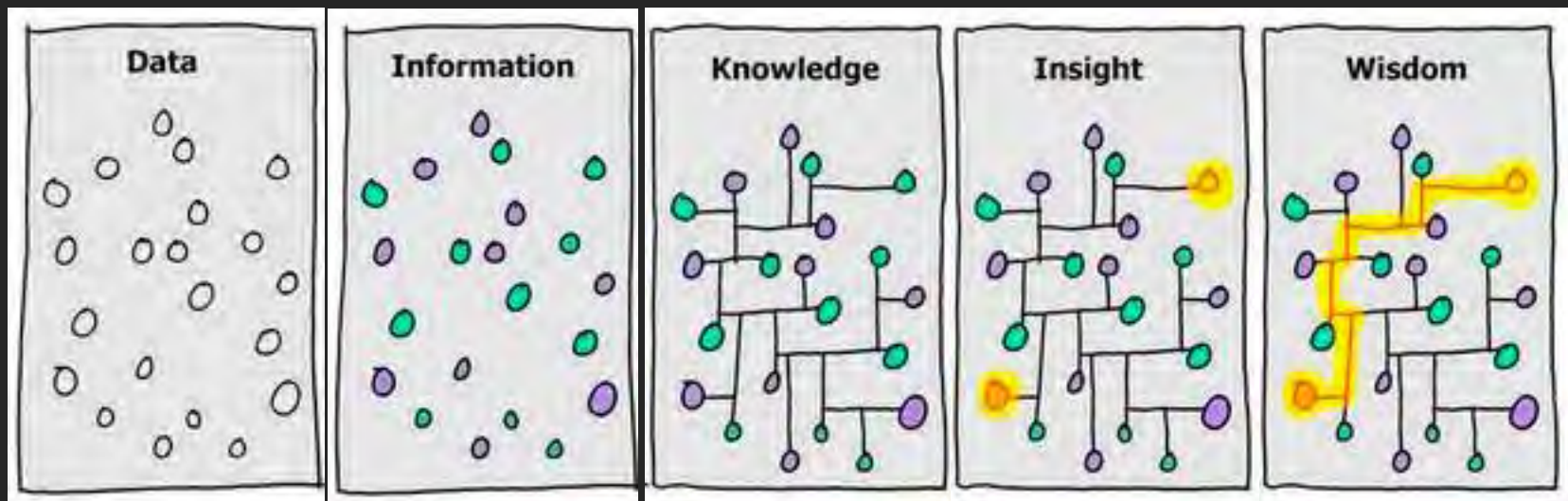
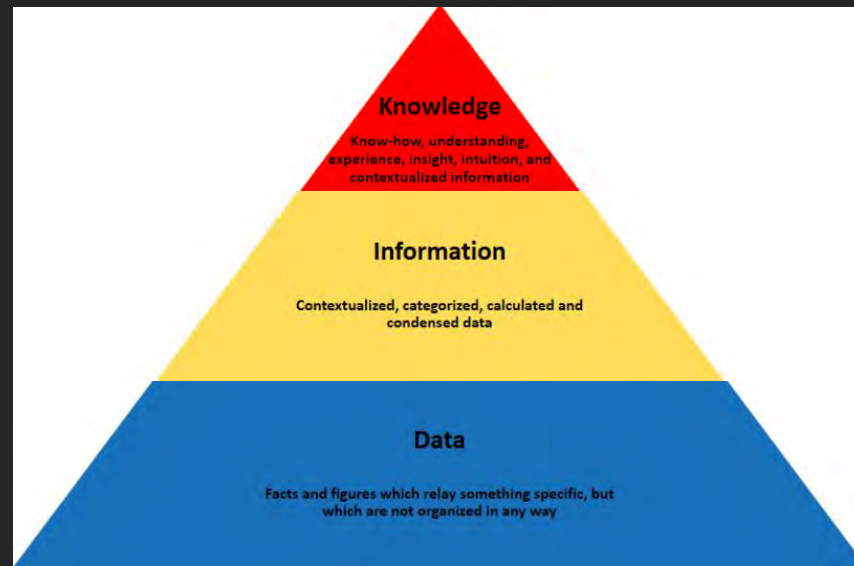
# A scalable precision medicine open knowledge network



# Building “Google maps” for health

- To develop a first-in-class AI system to track and predict disease outcomes
- System is based on a multi-scale, data-driven knowledge graph

# Data -> information -> Knowledge

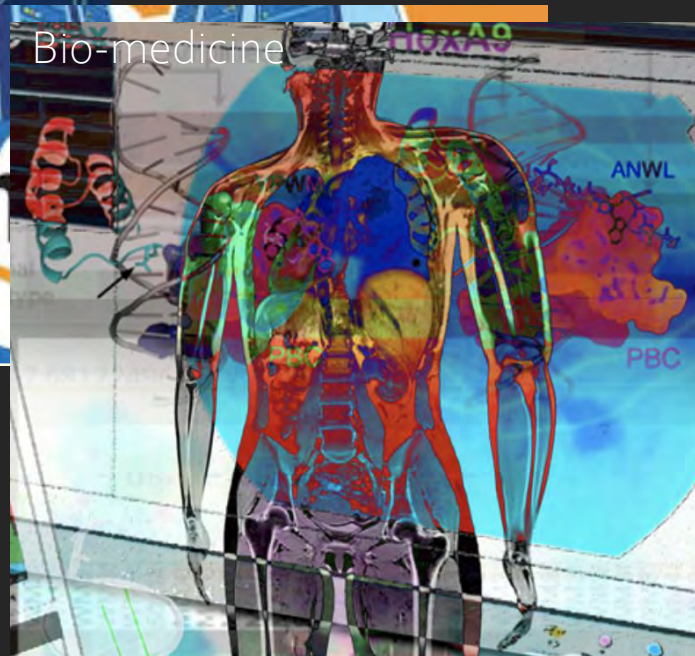
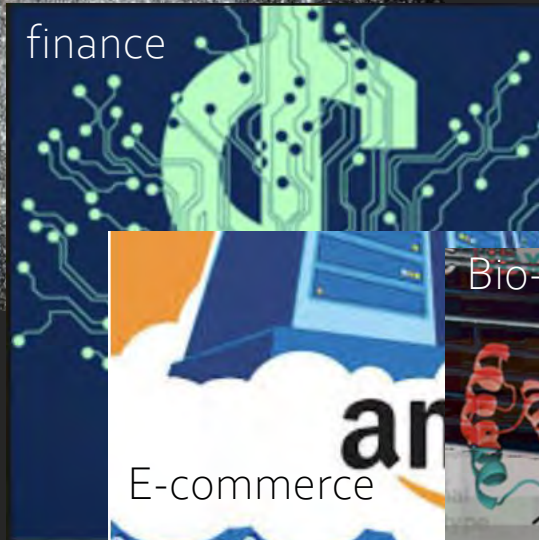




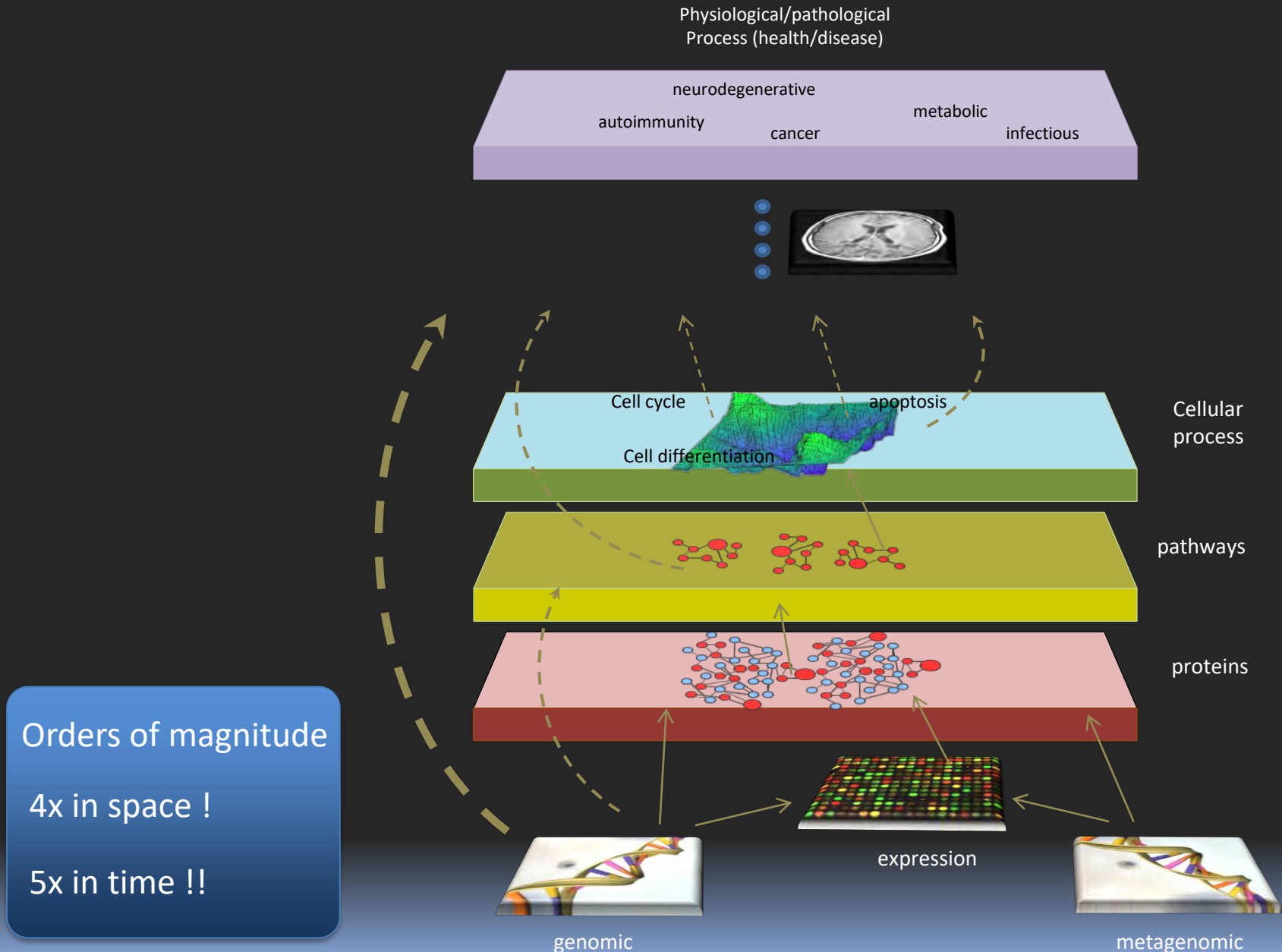
# How much data is out there?

- ~33 Zettabytes in 2018 (~175 by 2025)
- more data was generated in 2017 than in the history of humanity
- 90% of the world's data was created in last 2 years

# Who has the biggest problem to solve with Big data?



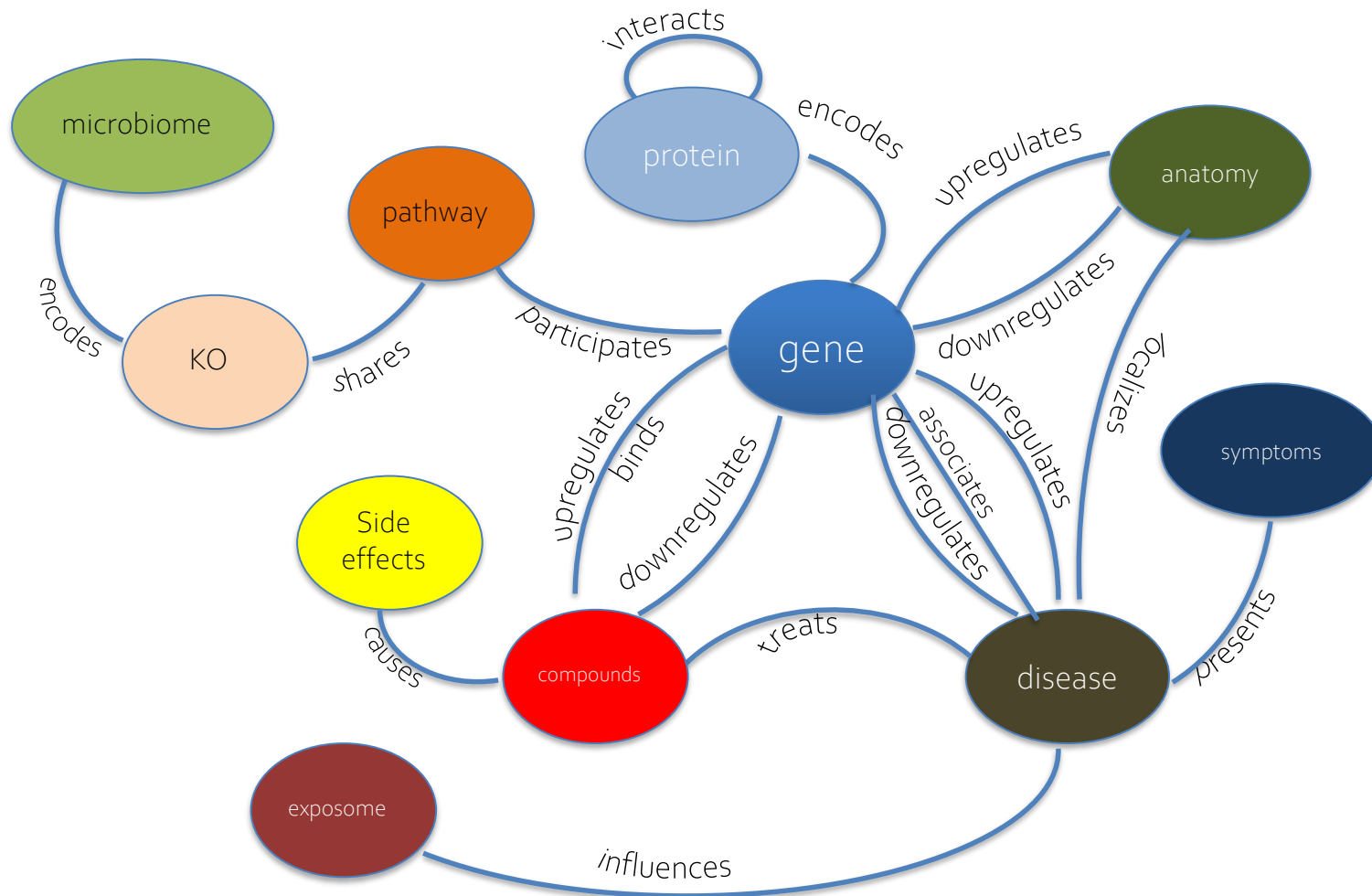
# Hierarchical organization of biological complex



# SPOKE

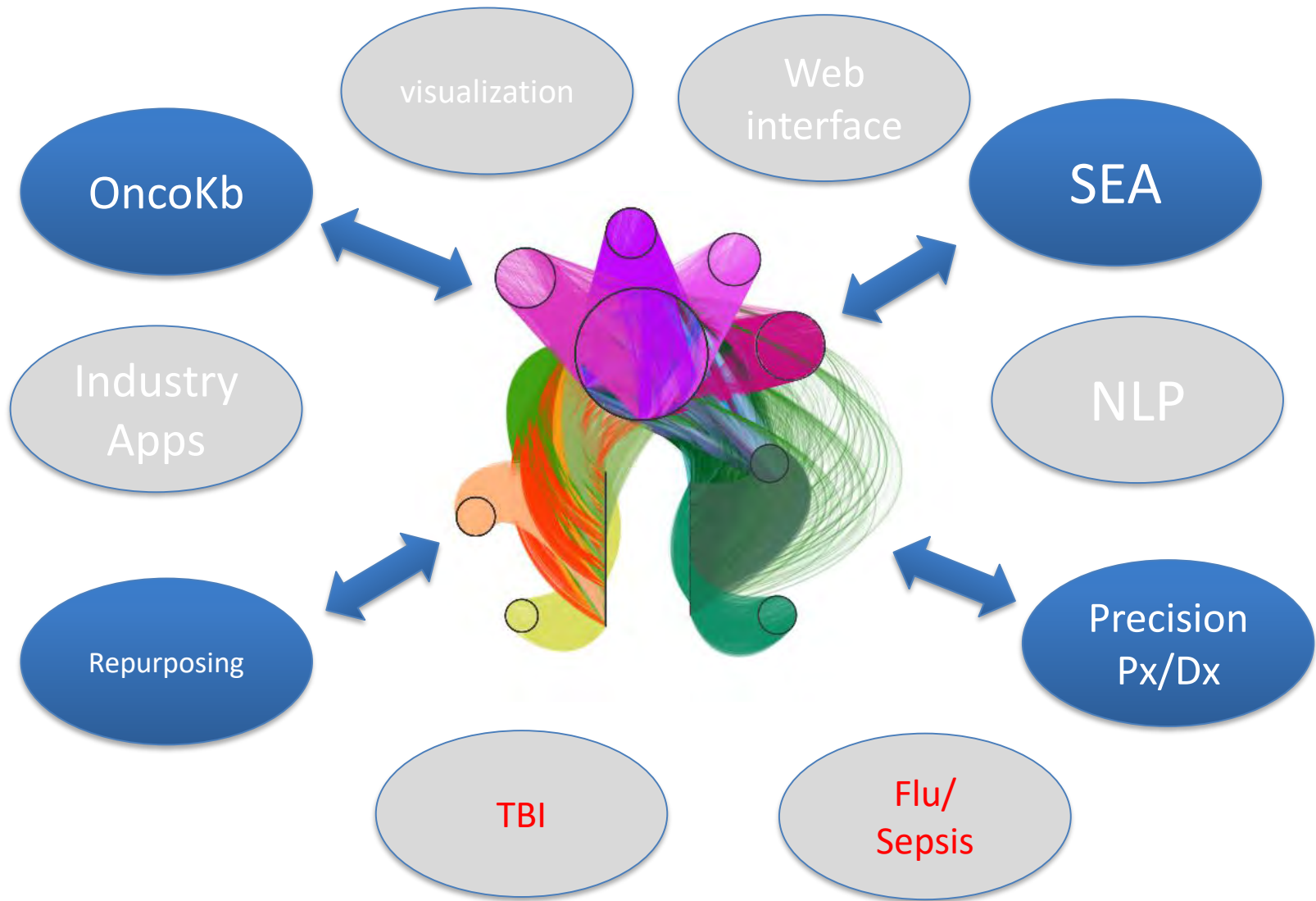
(Scalable Precision Medicine Open Knowledge Engine)

A database of databases



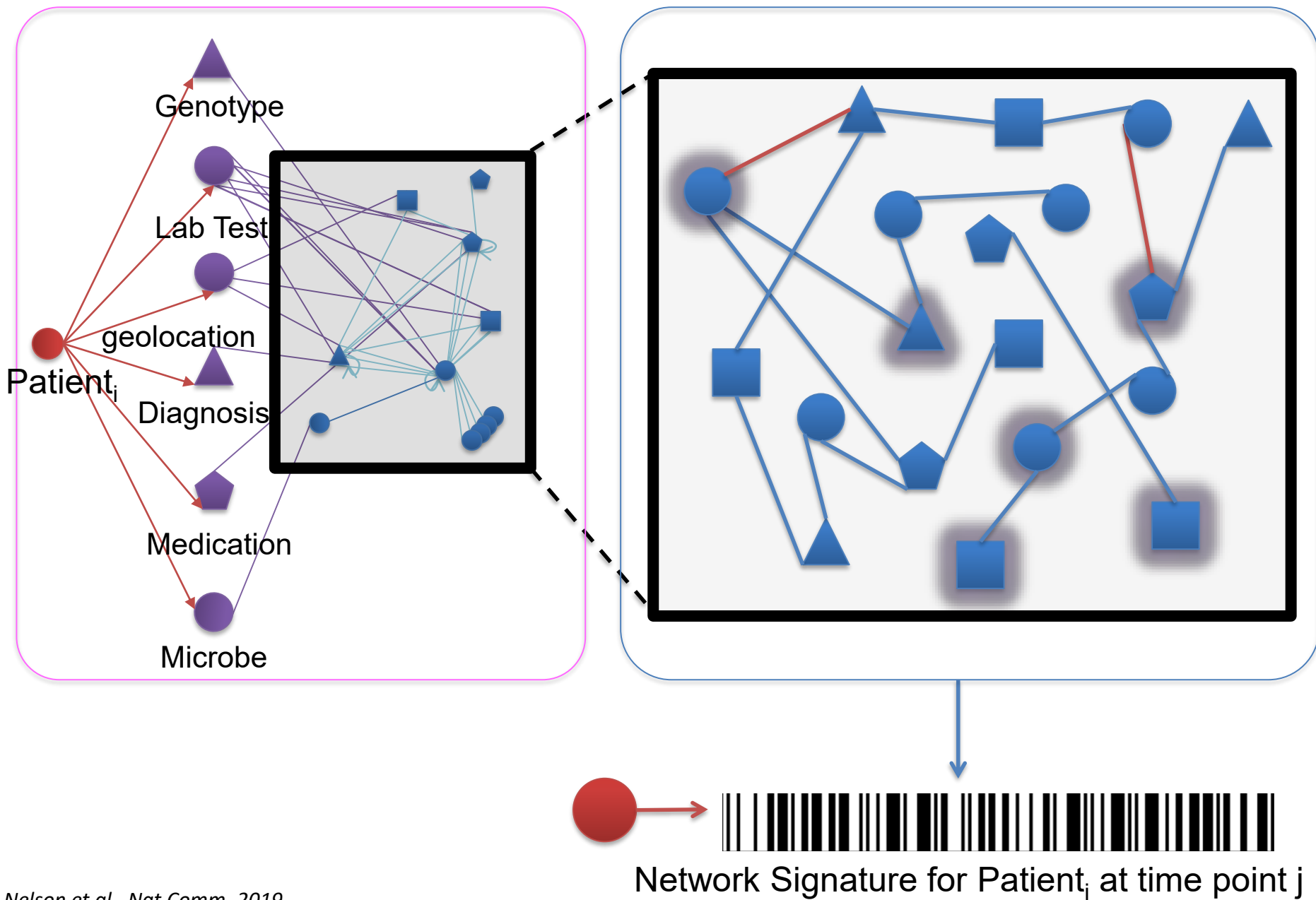
25+ databases  
> 3 M nodes  
> 5 M edges

# SPOKE Apps model



## EHR

## SPOKE



UCSF Patient  
Cohort

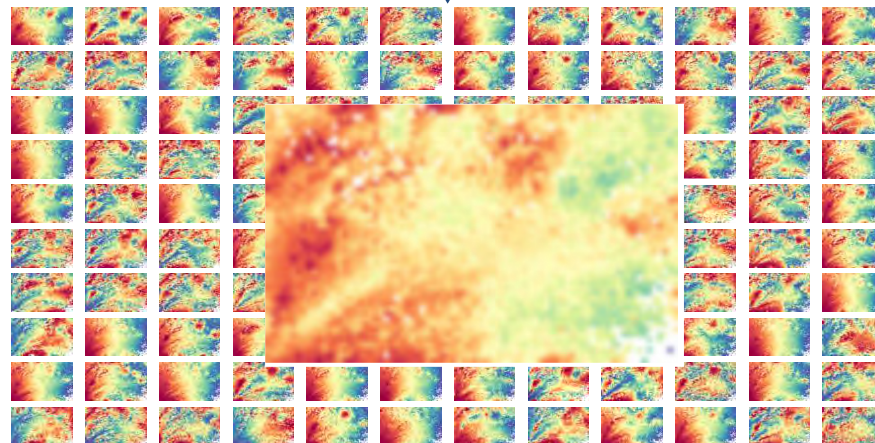
4,357 PTSD  
Patients



Add New Patient Record

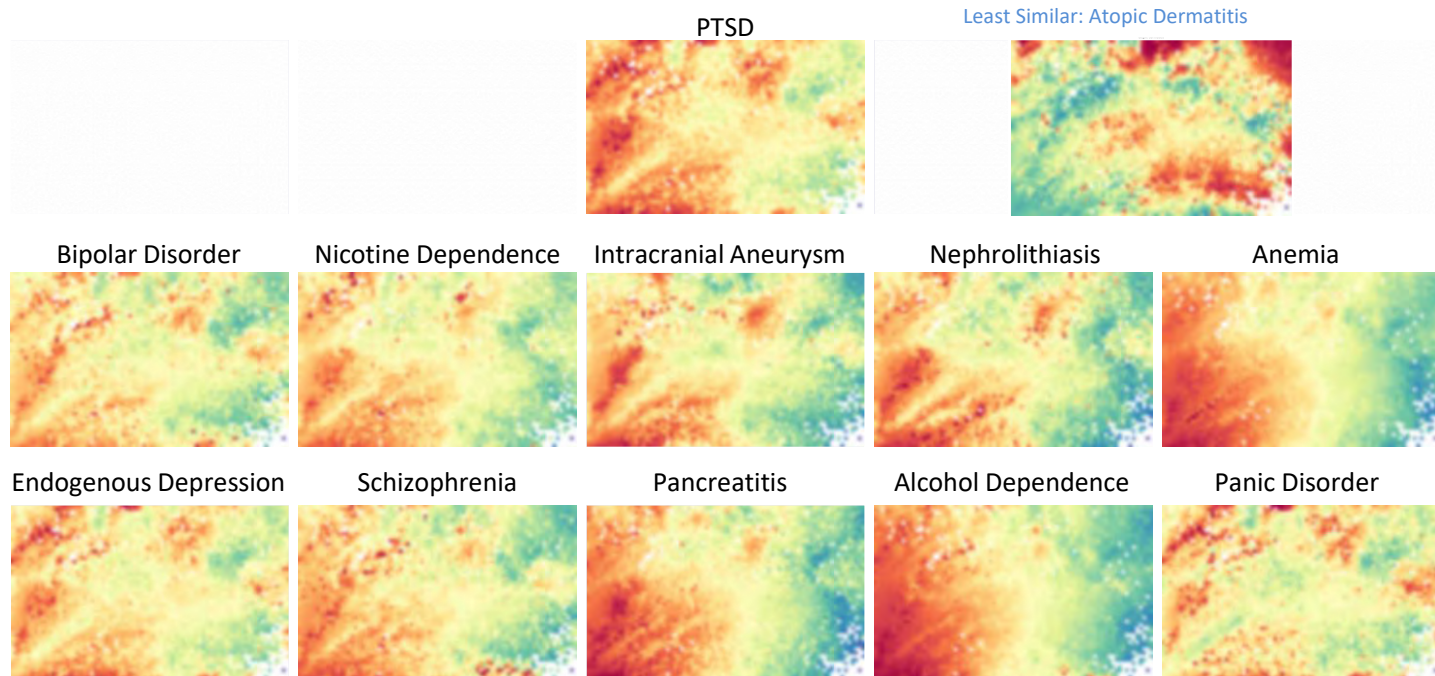
Browse...

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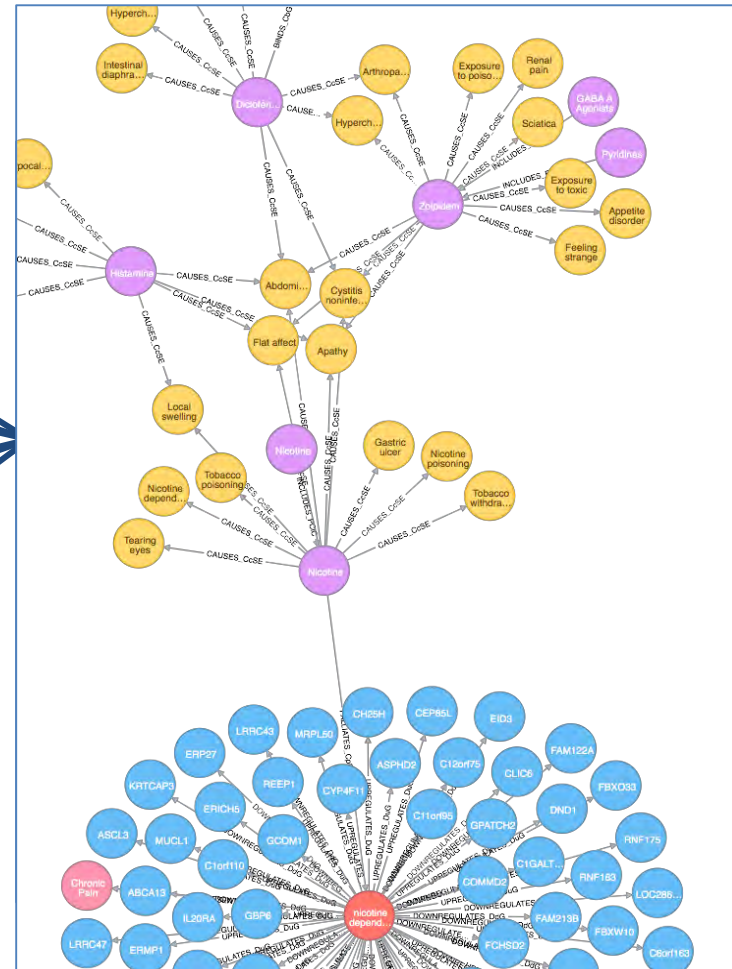




# Most Similar Diseases to the Average PTSD Patient at UCSF









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## NSF Convergence Accelerator (C-Accel)

