

# Cytokine Graph Database

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# Overview

1 Introduction

2 What?

3 How?

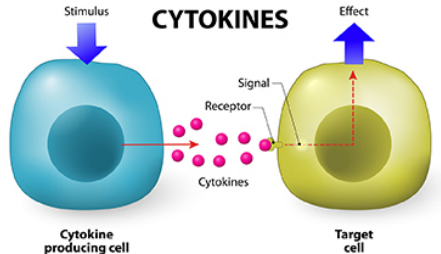
4 Why?

5 Demo

6 Next steps

# Cytokines

- Molecular messengers between cells
- Interact with cells of the immune system
- Regulate the body's response to disease and infection
- Mediate normal cellular processes in the body



# Graph databases

- Nodes
- **Directed** edges / relationships
- Properties

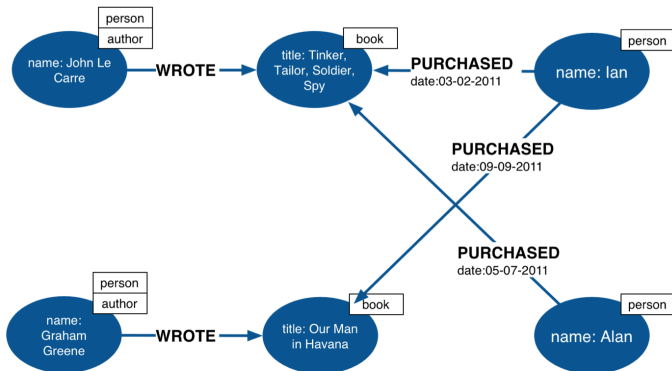




Figure: Neo4j - a graph database management system

## Example of a Cypher query

```
MATCH (:Person { name: "Alan" })-[p:PURCHASED]->(b:book)
RETURN p.date, b
```

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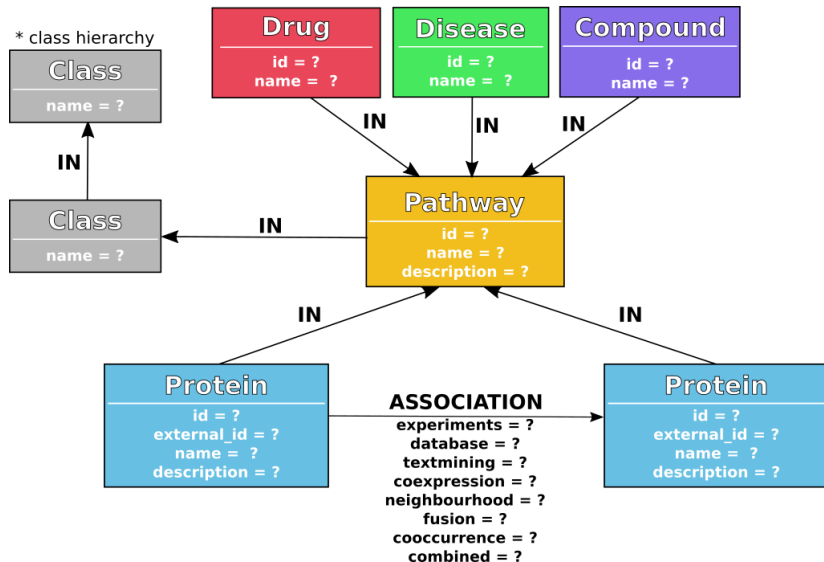
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# Database scheme



- Nodes

- Protein:
- Pathway:
- Drug:
- Disease:
- Compound:
- Class:

- Relationships

- **ASSOCIATION:**
- **IN:**



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- **STRING**

- proteins
- protein - protein associations



- **KEGG PATHWAY**

- pathways
  - classes
  - compounds
  - drugs
  - diseases

# STRING score channels

- 7 score channels → combined score

- 1 Experiments
- 2 Database
- 3 Textmining
- 4 Co-expression
- 5 Neighbourhood
- 6 Fusion
- 7 Co-occurrence

## Evidence suggesting a functional link:

Neighborhood in the Genome:	none / insignificant.
Gene Fusions:	none / insignificant.
Cooccurrence Across Genomes:	none / insignificant.
Co-Expression:	yes (score 0.152). In addition, putative homologs are coexpressed in other species (score 0.043).
Experimental/Biochemical Data:	yes (score 0.360).
Association in Curated Databases:	yes (score 0.900).
Co-Mentioned in PubMed Abstracts:	yes (score 0.962). In addition, putative homologs are mentioned together in other species (score 0.127).

Combined Score: 0.997

Figure: Scores for CCR5 and CCL5

- ① Collect the data
  - STRING: SQL dumps
  - KEGG PATHWAY: flat text files
- ② Extract the useful information
  - Query the STRING SQL database
  - Parse the KEGG flat files
- ③ Merge the databases
  - Map KEGG identifiers to STRING identifiers
- ④ Build a graph database
  - Translate the associative data into a graph

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# What is the point?

- Drug discovery
- ...

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(demo)



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# Next steps

- Extend the database
  - Protein - protein actions
  - [KEGG DRUG](#)
  - [KEGG DISEASE](#)
  - [KEGG COMPOUND](#)
- Other species
- All proteins
- Web server
- Machine learning

# Thanks :)

<https://backofenlab.github.io/cytokine-graph-db/>  
<https://github.com/BackofenLab/cytokine-graph-db>