



SURAJ THYAGARAJAN PARAMASIVAM

CSE 736- DATABASE SEMINAR



Data Integration and Genomic Medicine

Brenton Louie, Peter Mork, Fernando Martin-Sanchez, Alon Halevy, Peter Tarczy-Hornoch

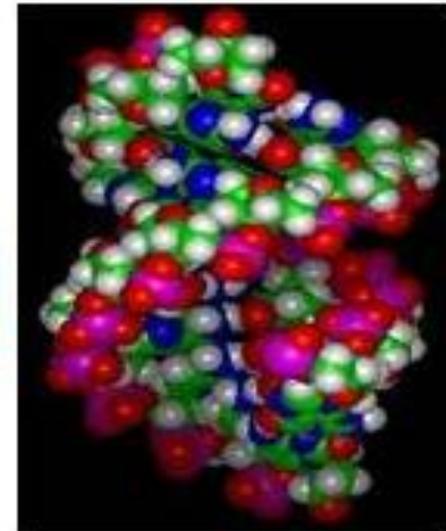
Bio2RDF: Towards a mashup to build bioinformatics knowledge systems

Francois Belleau, Marc-Alexandre Nolin, Nicole Tourigny, Philippe Rigault, Jean Morissette

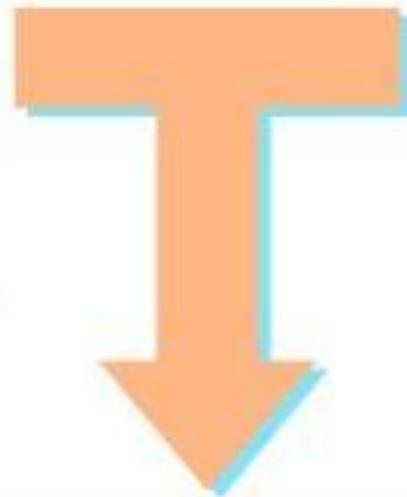
What is Bioinformatics ???



Computer systems



Biological systems



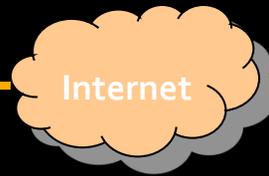
BIOINFORMATICS

Why Bioinformatics ???



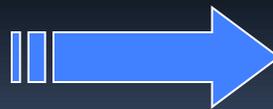
Descriptive, observational
science
→ Hypothesis driven

RESEARCH



Predictive information
Science
→ "Discovery" driven

SEARCH

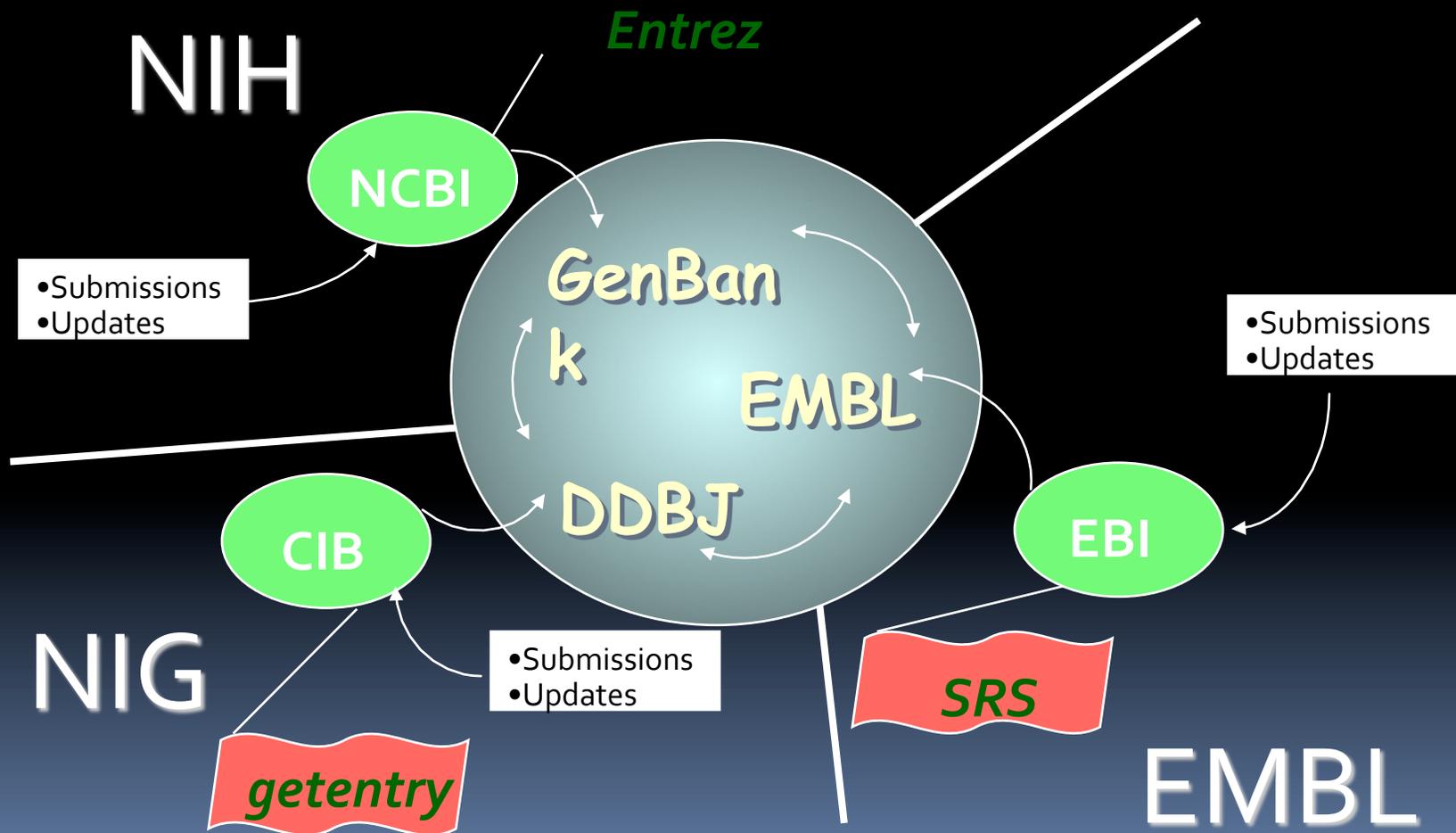


If not for Bioinformatics !!

- Structural Plasticity of the Human Genome
(Copy number variants)
- Individual Human Variation (when a mutation is not a mutation!)
- Alternate Splicing
- Non-Coding RNAs (genes?)

None of these most important genetic discoveries would have been possible !!!

Data Integration.. Why is that important ??



Common Data Integration Architectures

Data Warehouses

-  Fast queries and clean data
-  Stale Data, Complex Schema

Database Federation

-  Current Data, Flexible architecture
-  Slower queries, Complex Schema, unclean Data

Database federation with mediated schema

-  Current Data, Flexible architecture, schema tailored to users
-  Slower queries, complex schema, unclean data, mapping from source schema to mediated schema required

Peer data management systems

-  Current Data, Flexible Architecture, Schema Tailored to users, Mapping between schemas distributed across peers
-  Experimental, slower queries, unclean data



Two Dimensions of Data Integration

The Integration Axis

(Where the data resides)

Data and Knowledge Representation



Integration Architecture

- Data Warehouse
 - Faster Queries – non trivial for biologists since performance is often the key
 - Handling Volumes: The volume of data in this field is simply too high to handle. Updates suffer and Maintenance becomes an issue
 - Schema Restrictions: The restriction of inability to create a global schema is a deterrent since data is extremely rich
 - Best suited for specific and narrow areas of research .
Eg. UCSC Genome Browser, BioMolQuest..



■ Database Federations

- Common Data Model – maintains a common data model and relies on schema mapping for integration
- Federations relieve the temporal problems of a data warehouse since they reside at the source and are updated constantly
- Some of the extremely difficult queries could be solved using database federations

- 
- Database Federations with Mediated Schema
 - Dealing with Various Source Schema- This drawback of database federations is dealt by having a database federation with mediated schema
 - Federations as Middleware – The federations with mediated schema act as middleware, where data sources are mapped to mediated schema
 - Best suited to situations when researchers need to ask complex questions spanning disparate knowledge resources.



- Peer Data Management Systems

- Tailored and Focused Mediated Schema – Developing such schemas and integrating is PDMS.
- Each Data source provides a semantic mapping to one or more peers
- Addresses the problem of creating a global mediated schema
- Technology still in evolutionary stage



Data and Knowledge Representation

- Relational Schemas
 - Traditional model of table with tuples and attributes
 - Well understood and robust, but is modeling complex
 - Hierarchically structured biological data is difficult to model
 - Most common and ubiquitous



■ Semi Structured Data

- Free from rigid structures
 - Data with a series of labels and associated values
 - More natural modeling of Biological data due to features like nesting
 - Complex relationships are still difficult to model
 - XML, RDF are examples
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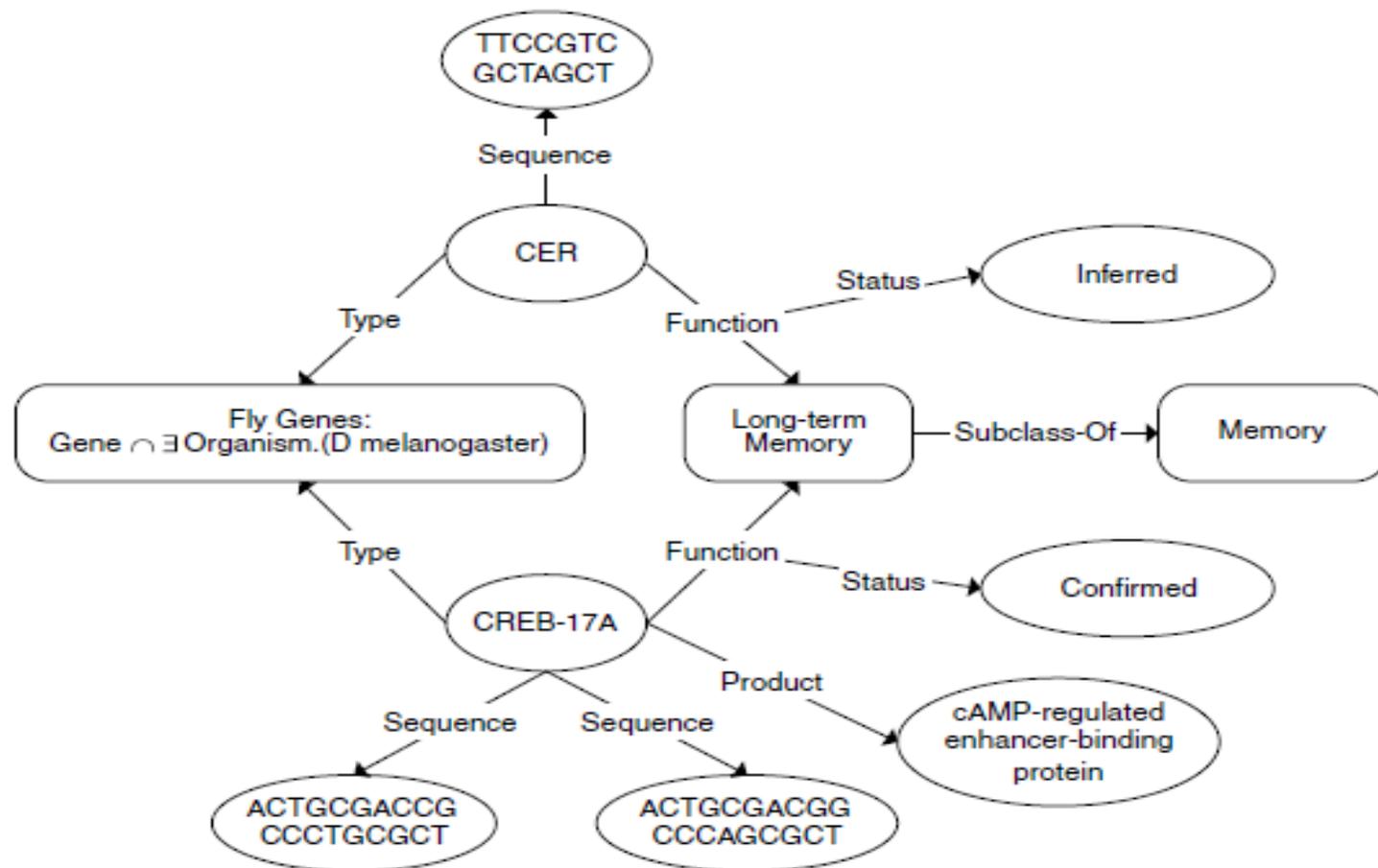
■ Ontology

- Defined as a “specification of a conceptualization”
- Best suited to represent semantic web
- Specify objects classes, relationships and functions
- Well suited for representing biological data

```

<?xml version="1.0"?>
<GeneList>
  <Gene symbol="CREB-17A" organism="D. melanogaster">
    <Sequence>ACTGCGACCGCCCTGCGCT</Sequence>
    <Sequence>ACTGCGACGGCCAGCGCT</Sequence>
    <Product>cAMP-regulated enhancer-binding protein</Product>
    <Function id="0007616" status="confirmed"><Term>long-term memory</Term></Function>
  </Gene>
  <Gene symbol="CER" organism="D. melanogaster">
    <Sequence>TTCCGTCGCTAGCT</Sequence>
    <Function id="0007616" status="inferred"><Term>long-term memory</Term></Function>
  </Gene>
</GeneList>

```



Genomic Medicine with relevance to Data Integration

- Modern Human Genetics
 - Researchers “Swim a sea of data” to study diseases and their links to genes
 - Lack of Standards, Presence of huge number of data sources makes it even more difficult
 - Queries often vague and highly complex, require join of multiple databases
 - Difficulties in combining clinical and genetic information



- Microarray Studies

- Genes represented as spots on microarrays
 - For each experiment, external annotation needed which often come from public databases
 - Need integrated information to perform studies effectively
- 

Application of Data Integration concepts to genomic medicine

Warehoused
Data

Data Storage
(Tractability of Queries & Data Cleansing ->)

Biobanks

Human Genetics

Federated
Data

Pharmacogenomics

Genomics in Clinical
Practice

Microarrays

Rational Drug Design

Relational

Semi-Structured

Mediated Schema

Ontologies

Knowledge Representation
(Expressiveness ->)

BioBanks

- Also known as a biorepository
- A place that collects, stores, processes and distributes biological materials and the data associated with those materials
- Stored as Relational Tables
- <http://www.ukbiobank.ac.uk> – a public biobank

MicroArrays

- A multiplex technology used in molecular biology and in medicine
- It consists of an arrayed series of thousands of microscopic spots of DNA oligonucleotides, called features
- A repository containing microarray gene expression data is the Microarray database

Genomics in Clinical Practice and Rational Drug Design

- Technologies of Future !!! Not yet completely developed
- Some breakthrough achieved.. Drugs like Relenza to treat influenza
- Rational drug design is the creation of drugs based on the structure of the drug receptor
- Drug Design is based on the structure of the protien

Gaps in DI research to facilitate genomic medicine

- Data Availability
 - Clinical data still scarce in comparison to bioinformatics data
- Privacy
 - Issues of “De-Identification” still an issue
 - Every DNA is a Unique fingerprint
- Data issues
 - Most data available as Natural Text, More mining required



- Lack of Standards

- Too much data, Too little standards
 - Integration of diverse complex data types including genomic, proteomic, clinical, pharmacological and chemical requires standards for proper semantic integration of heterogeneous data
- 



Questions ??????