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Federated Data Analytics for Genomics Data

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Master Degree Course in **Computer Engineering** – **Web Information and Data Engineering** (WIDE)

A. A. **2023/2024**

The Challenge

Biomedical data is becoming increasingly complex. The rapid evolution of data storage systems has created a landscape where:

- Diverse data models exist (relational, hierarchical, graph-based);
- Integration of these models is crucial for making data meaningful.

Understanding genetic diseases and advancing personalized treatments relies on integrating and analyzing this diverse data effectively.

Background

Methodologies:

1. Data **Federation**
2. Data **Virtualization**
3. Semantic Data **Integration** (OBDA)
4. Knowledge Graphs, Ontologies & RDF

Technologies:

1. SPARQL query language;
2. Data **Virtualization** solutions: Denodo, Dremio, Teiid
3. **OBDA** Solutions: Mastro, Ontop

	Support	Free and Open Source	Well Documented	Scalability	Solid Logging Capabilities
Denodo	✓		✓	✓	✓
Teiid		✓			
Dremio	✓	✓	✓	✓	✓

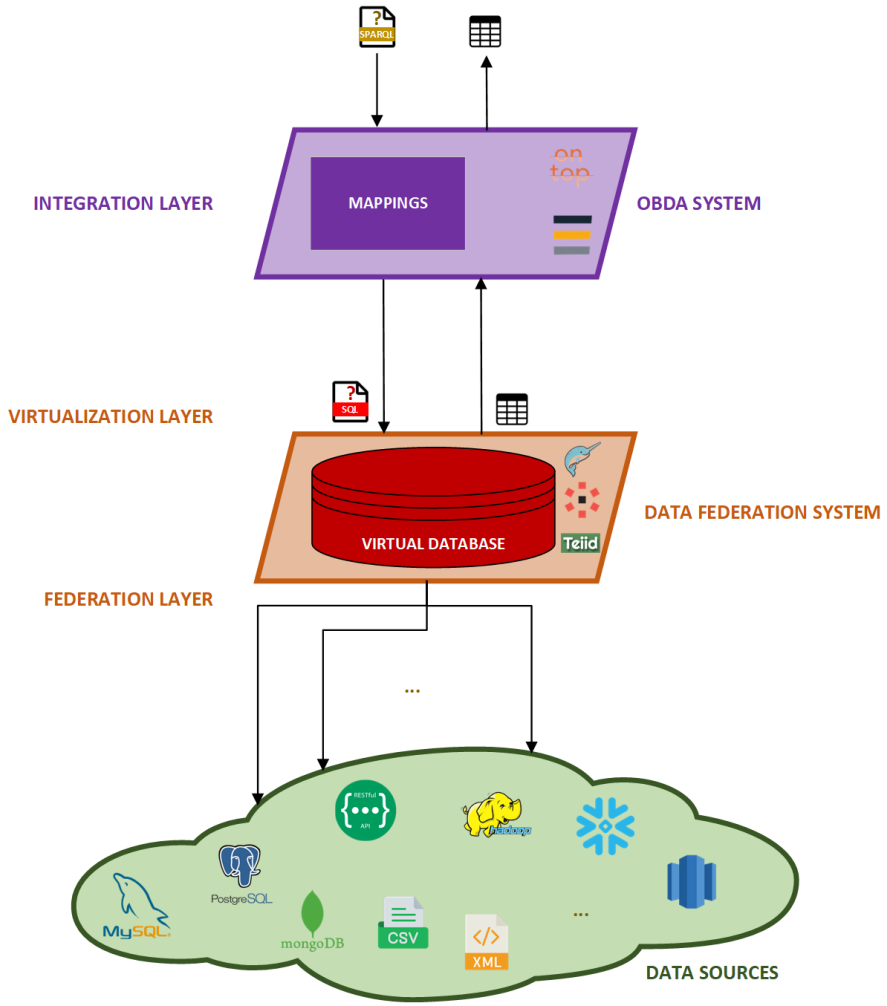
Proposed Solution

Federated Data Analytics System

A system designed to integrate and analyze clinical and genomics data seamlessly.

- Objective: Enable researchers to perform complex queries across multiple datasets without extensive preprocessing.
- Key Components: OBDA, Data Virtualization, Ontology.

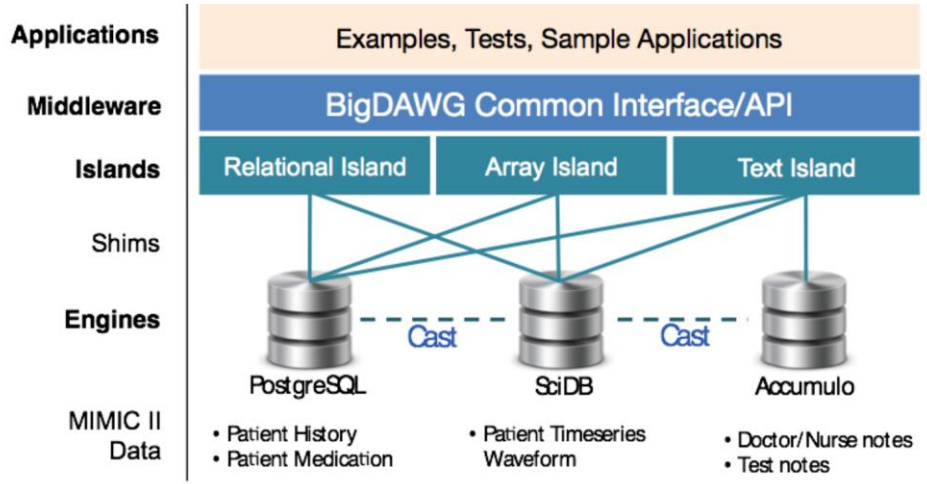
This approach reduces complexities in biomedical data management and accelerates research.



Related Polystore Solutions

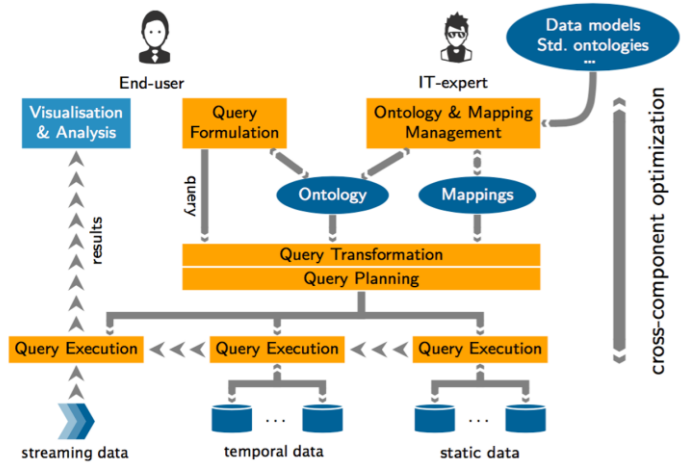
BigDAWG

- It is structured in 4 layers, from the data stores layer up to the application layer.
- A crucial role is played by the "**Island layer**", a layer of autonomous components that abstracts data sources.



Optique

- Initially conceived as an EU funded project, now implemented and maintained by Siemens.
- 3-layers architecture, **OBDA based**, optimized for streaming data. Provides a visual query tool to ease its use.



EU Project HEREDITARY

Objective:

Transforming our understanding of brain diseases through integrated multimodal data analysis.

Focus Areas:

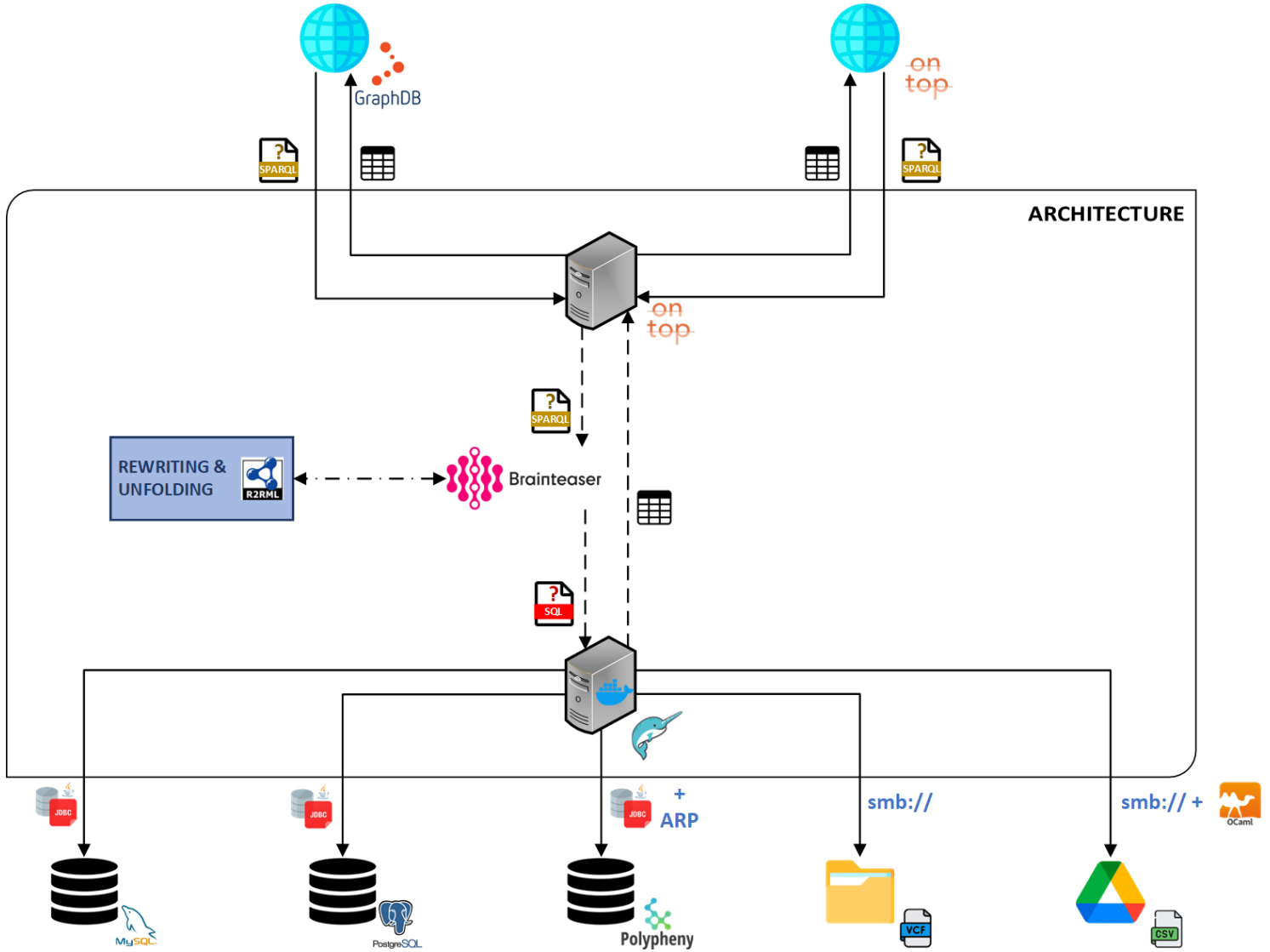
1. Integrating genomic and clinical data from various European stakeholders;
2. Addressing the challenges of data heterogeneity;
3. Ensuring privacy compliance and data security.

(one of the) Outcomes:

A unified DBMS that supports advanced querying capabilities across different data sources and that allows to gather insightful analytics.

System Architecture

- **Ontop**: Semantic Data Integration Layer.
- **Dremio**: Data Federation and Virtualization Layer.
- **BRAINTEASER Ontology**: Provides the vocabulary structure for querying.



System Architecture – Data Sources

- **MySQL**: Stores structured clinical data.
- **PostgreSQL**: Manages additional clinical datasets.
- **Polypheny**: Supports various data models, enhancing flexibility.
- **NAS Shared Folders**: Hosts genomics data in VCF files.
- **Google Drive**: Integrates a cloud-based storage use case, hosting CSV files.

Each data source is connected via Dremio's federation layer, enabling a unified view across diverse data repositories.

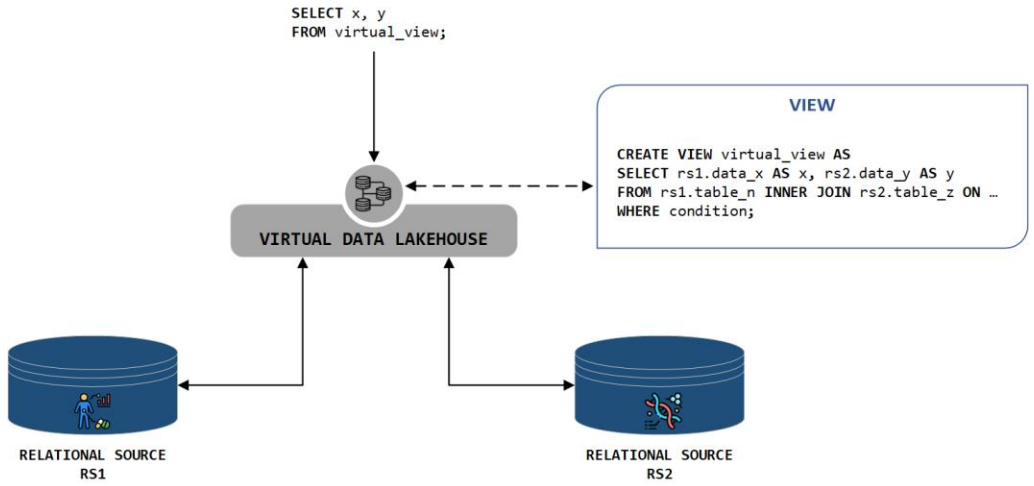
Federation and Virtualization

Virtualization Layer:

Creates virtual views without materializing data, optimizing performance and enabling complex queries across the federated system.

Federation Layer:

Integrates data across various sources, ensuring seamless access and querying.



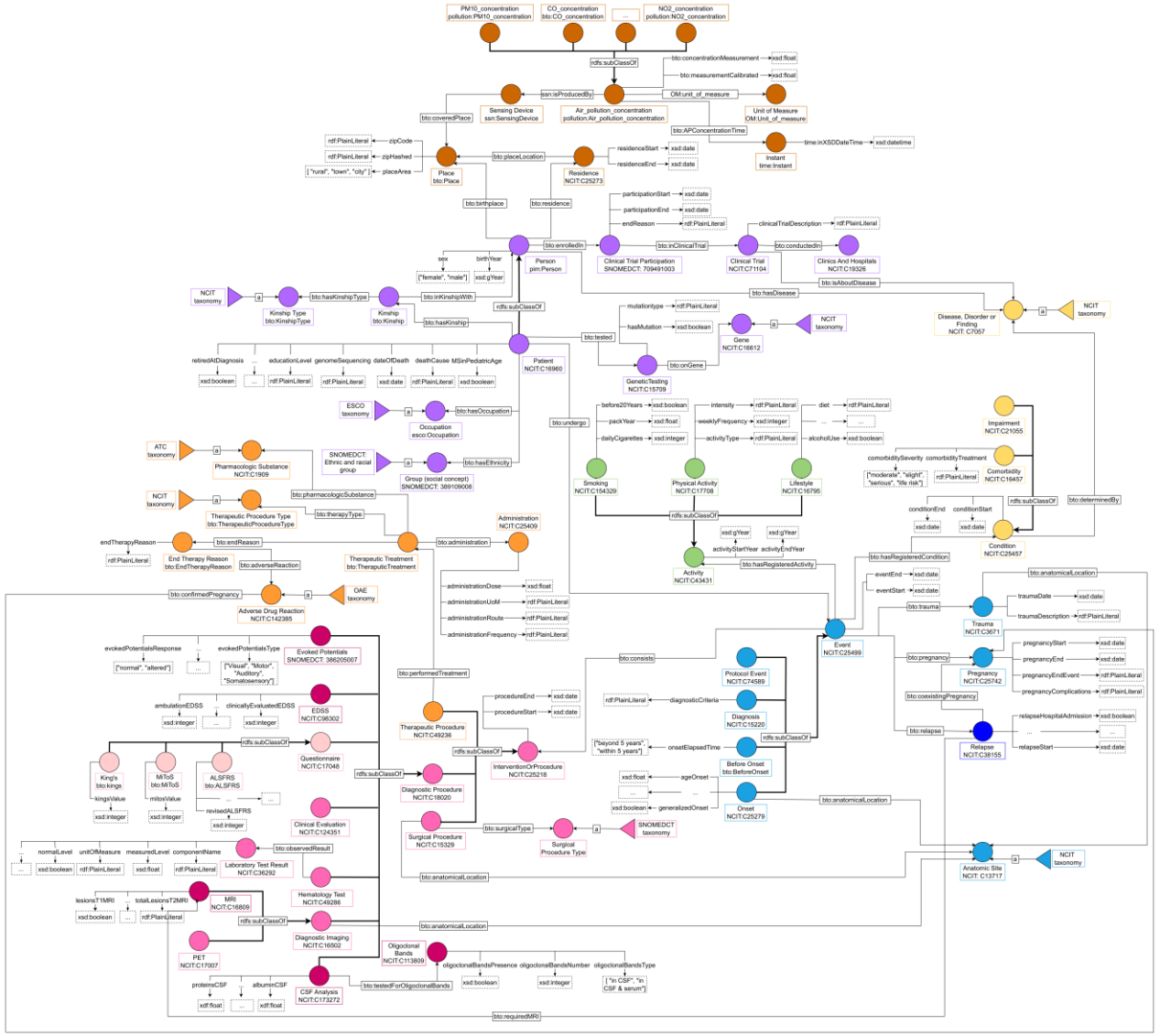
Dremio serves as the core engine for both layers, providing a robust platform for handling large-scale data integration tasks.

Ontology Integration

The **BRAINTEASER** Ontology is central to the system

Features:

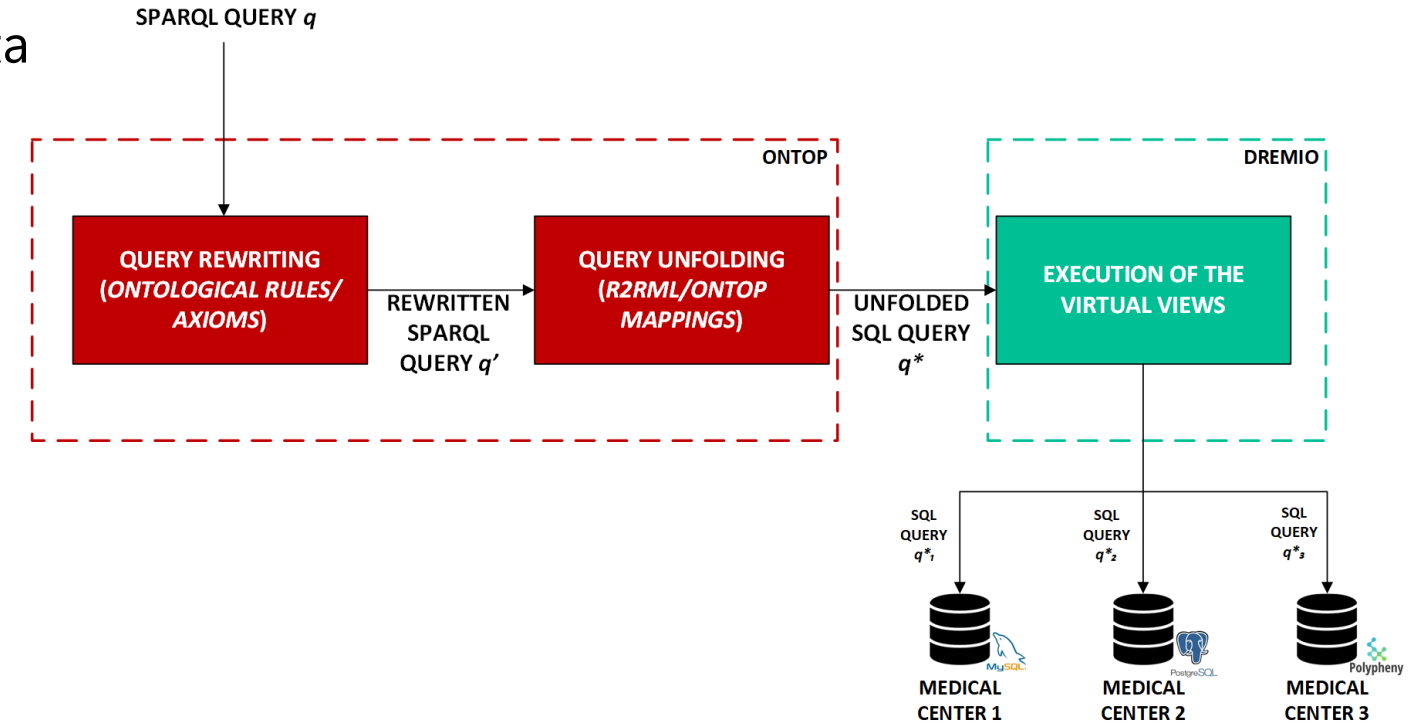
- Models relationships between clinical and genomics data;
- Enhances query capabilities through semantic enrichment;
- Ensures interoperability and scalability.



Use Case: ALSFRS Data Querying

Use Case: Querying ALSFRS (Amyotrophic Lateral Sclerosis Functional Rating Scale) data across multiple sources.

- **SPARQL Query**: Retrieves comprehensive patient data;
- **Process**:
  - Query Rewriting;
  - Unfolding into SQL;
  - Execution in Dremio.



```
PREFIX bto: <https://w3id.org/brainteaser/ontology/schema/>

SELECT ?patient ?date ?tot ?bulbar ?motor ?respiratory ?q1 ?q2 ?q3 ?q4
 ?q5 ?q6 ?q7 ?q8 ?q9 ?q10 ?q11 ?q12 WHERE {
    ?p bto:undergo ?e.
    ?e bto:consists ?alsfrs.
    ?alsfrs a bto:ALSFRS;
           bto:procedureStart ?date;
           bto:revisedALSFRS ?tot;
           bto:bulbarSubscore ?bulbar;
           bto:motorSubscore ?motor;
           bto:respiratorySubscore ?respiratory;
           bto:alsfrs1 ?q1;
           bto:alsfrs2 ?q2;
           bto:alsfrs3 ?q3;
           bto:alsfrs4 ?q4;
           bto:alsfrs5 ?q5;
           bto:alsfrs6 ?q6;
           bto:alsfrs7 ?q7;
           bto:alsfrs8 ?q8;
           bto:alsfrs9 ?q9;
           bto:alsfrs10 ?q10;
           bto:alsfrs11 ?q11;
           bto:alsfrs12 ?q12.
    BIND(SUBSTR( (STR(?p)), 48) AS ?patient)
}
ORDER BY ?patient ?date
```

Original SPARQL Query

REWRITING
+
UNFOLDING

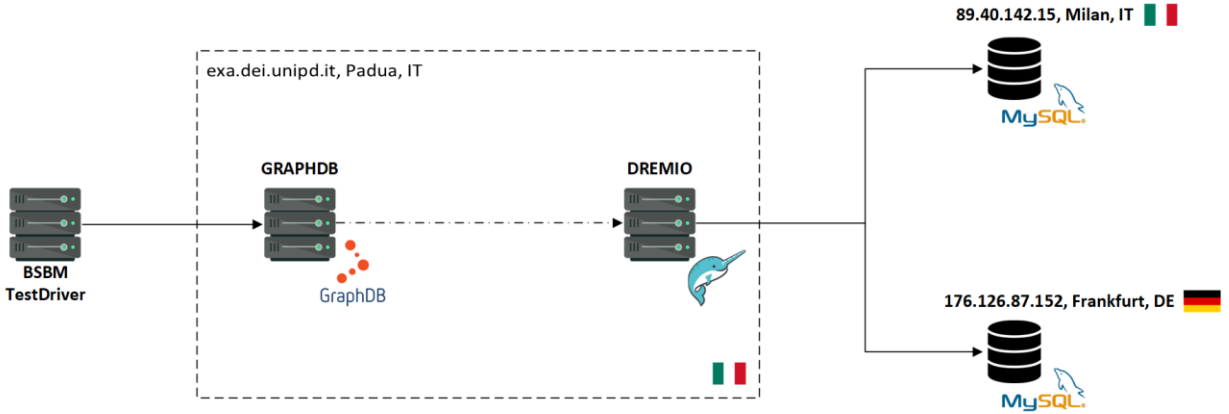
```
CONSTRUCT [patient, date, tot, bulbar, motor, respiratory, q1, q2, q3
    , q4, q5, q6, q7, q8, q9, q10, q11, q12]
   [date/RDF(CHARACTER VARYINGToVARCHAR(date2m51), xsd:datetime),
    q1/RDF(INTEGERToVARCHAR(q11m25), xsd:integer),
    motor/RDF(INTEGERToVARCHAR(motor1m55), xsd:integer),
    ...]
NATIVE
SELECT
    v23."bulbar1m17" AS "bulbar",
    v23."date2m51" AS "date",
    v23."motor1m55" AS "motor",
    v23."q101m44" AS "q10",
    v23."q111m42" AS "q11",
    v23."q11m25" AS "q1",
    v23."q121m41" AS "q12",
    v23."q21m24" AS "q2",
    v23."q31m23" AS "q3",

    --- ... more fields

    v23."patient26m9" AS "patient"
FROM (
    SELECT DISTINCT
        v7."bulbar" AS "bulbar1m17",
        v5."date2m51" AS "date2m51",
        v8."motor" AS "motor1m55",
        v1."patient" AS "patient26m9",
        v19."q10" AS "q101m44",
        v20."q11" AS "q111m42",
        v10."q1" AS "q11m25",

        --- ... more fields

        v6."tot" AS "tot1m45"
    FROM "clinical_data"."ALSFRS" v1

    --- ... recursive joins

    WHERE v1."patient" IS NOT NULL AND v1."date" IS NOT NULL
) v23

ORDER BY v23."date2m51" NULLS FIRST
```

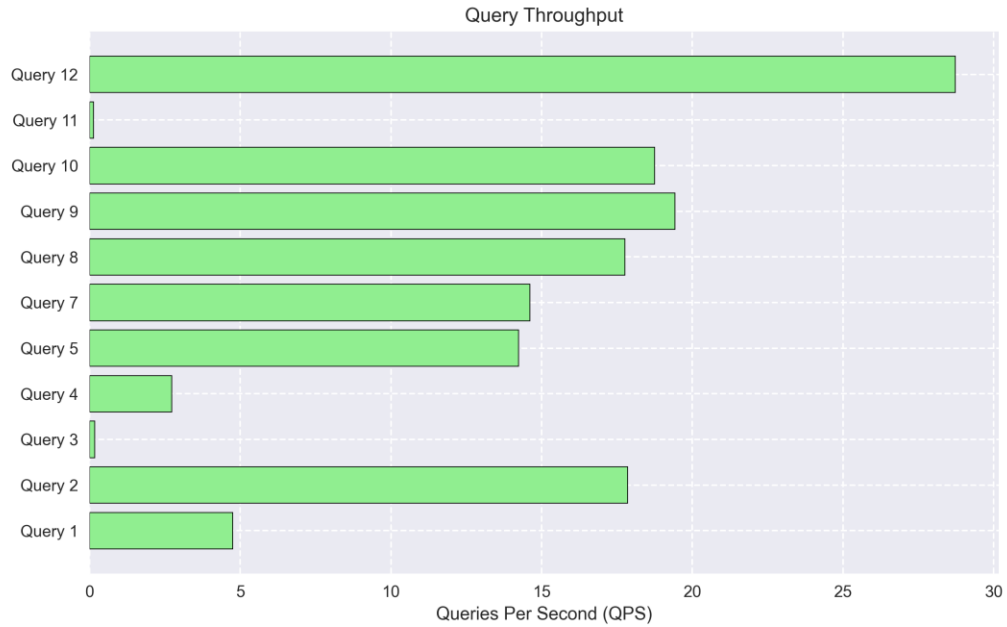
Unfolded SQL Query

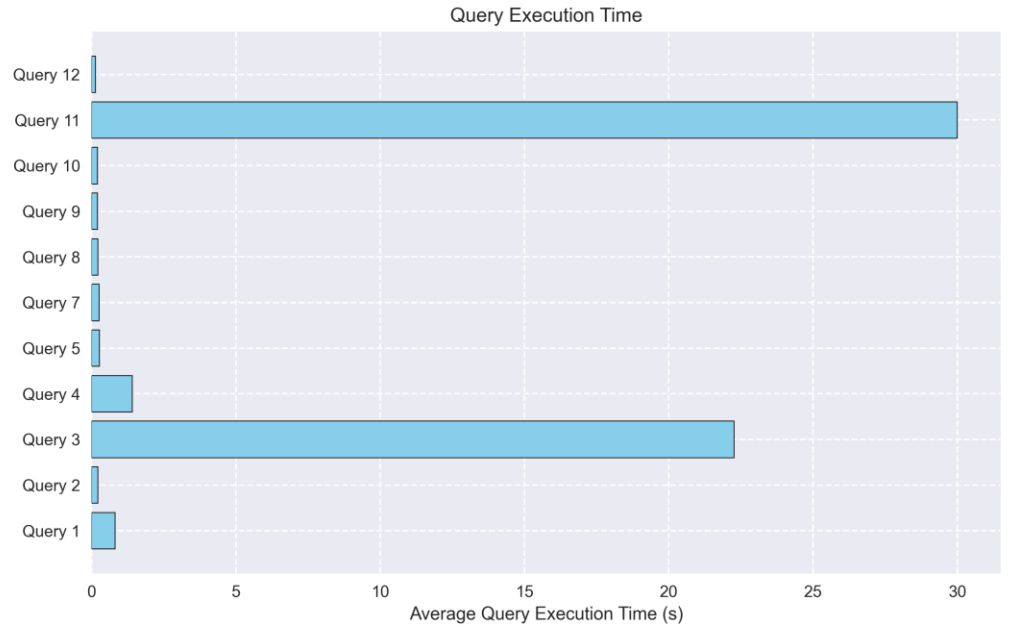
Benchmark 1: BSBM

BSBM Benchmark

Query performances evaluation over a synthetic dataset across diverse sources.



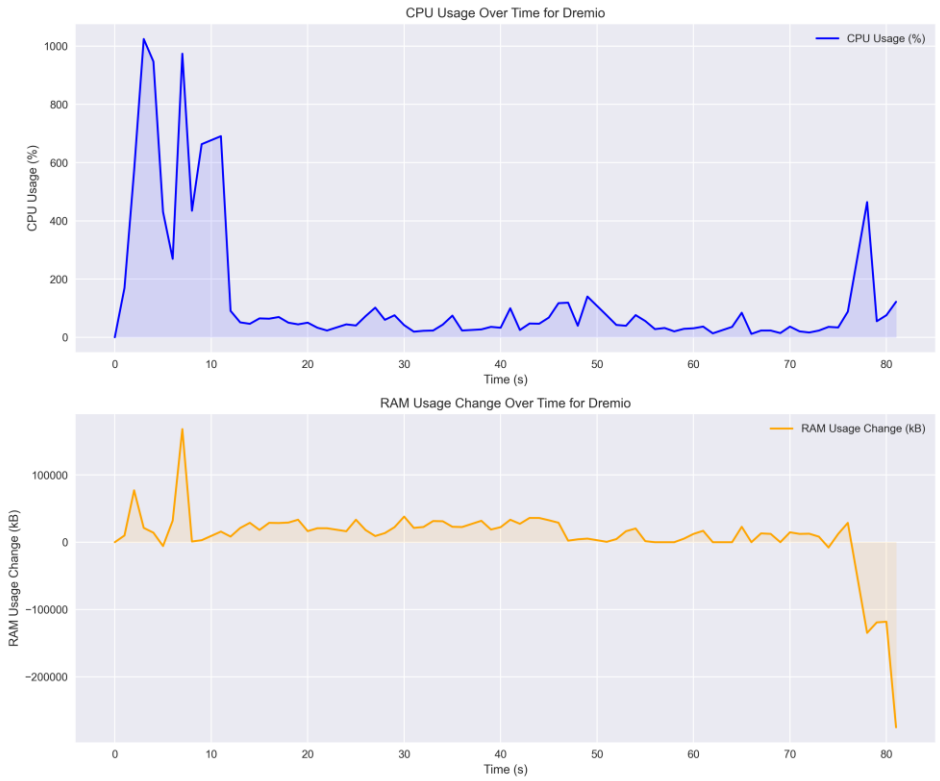




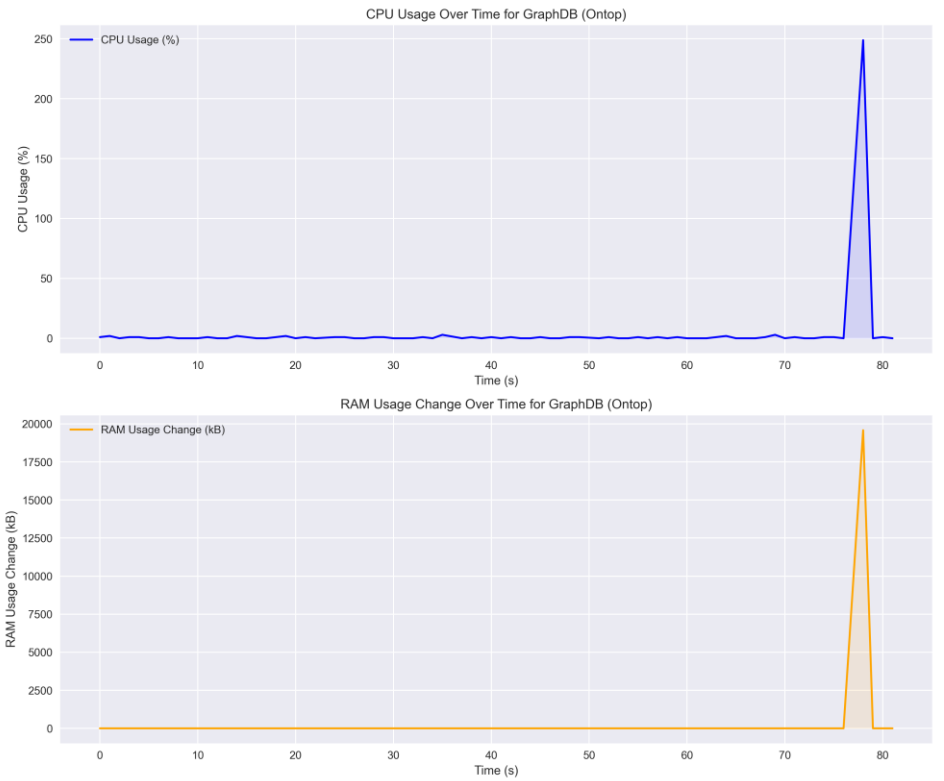
Benchmark 2: SEASHELL

SEASHELL Benchmark

Real-world clinical data highlights system efficiency in handling complex queries.



Dremio



GraphDB

Conclusions and Future Works

The federated data analytics system developed in this thesis offers a powerful tool for integrating and analyzing heterogeneous biomedical data.

Future Directions:

1. **Optimization**: Improve architecture performances, considering SEASHELL monitoring data as an entry point;
2. **Privacy**: Strengthen data security and compliance with regulations, such as GDPR;
3. **Usability**: Simplify both the architecture usage and deployment, so to reach a larger audience.

This system lays the groundwork for future advancements in biomedical data research, contributing to better healthcare outcomes.

Thank You!

Do you have any question?

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