

A Galaxy Interactive Environment for exploring the Neo4j Graph Database

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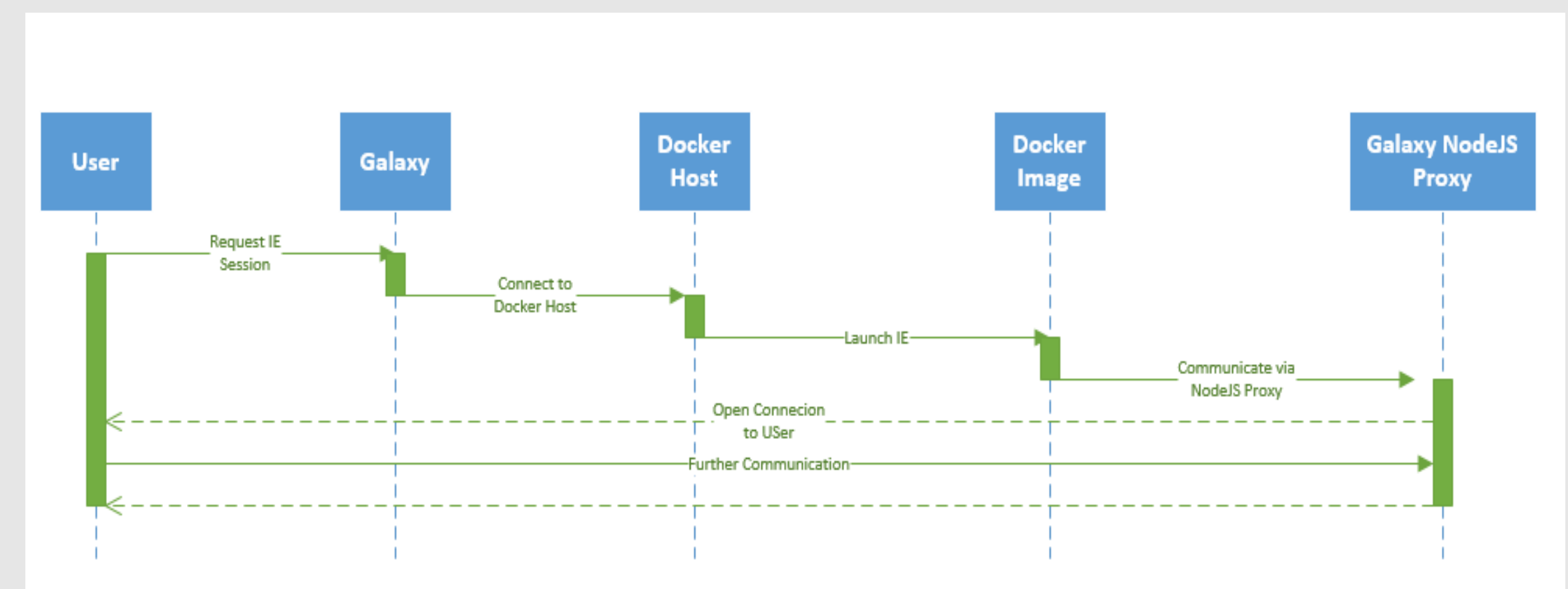
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Introduction

- Storing biological data involves modeling and storing thousands of entities that are interrelated in complex ways.
- Relational databases meet very specific needs and are not designed to fit all scenarios.
- Graph databases, which focus on connections between entities, are seen as a natural fit to these complex relationships.
- Neo4j is a highly scalable graph database with a declarative query language called Cypher.

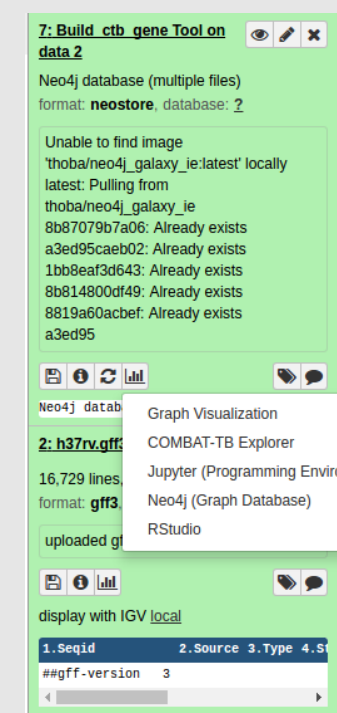
Implementation

- We implemented a Galaxy Interactive Environment (GIE) to explore a Neo4j database that is stored as a Galaxy dataset.
- The GIE was built by linking a Galaxy plugin to a Docker container based on a modified version of the Neo4j:2.3 Docker image.
- The Interactive Environment is launched from the visualisation menu and is only available for Neo4j database (neostore) datatypes.

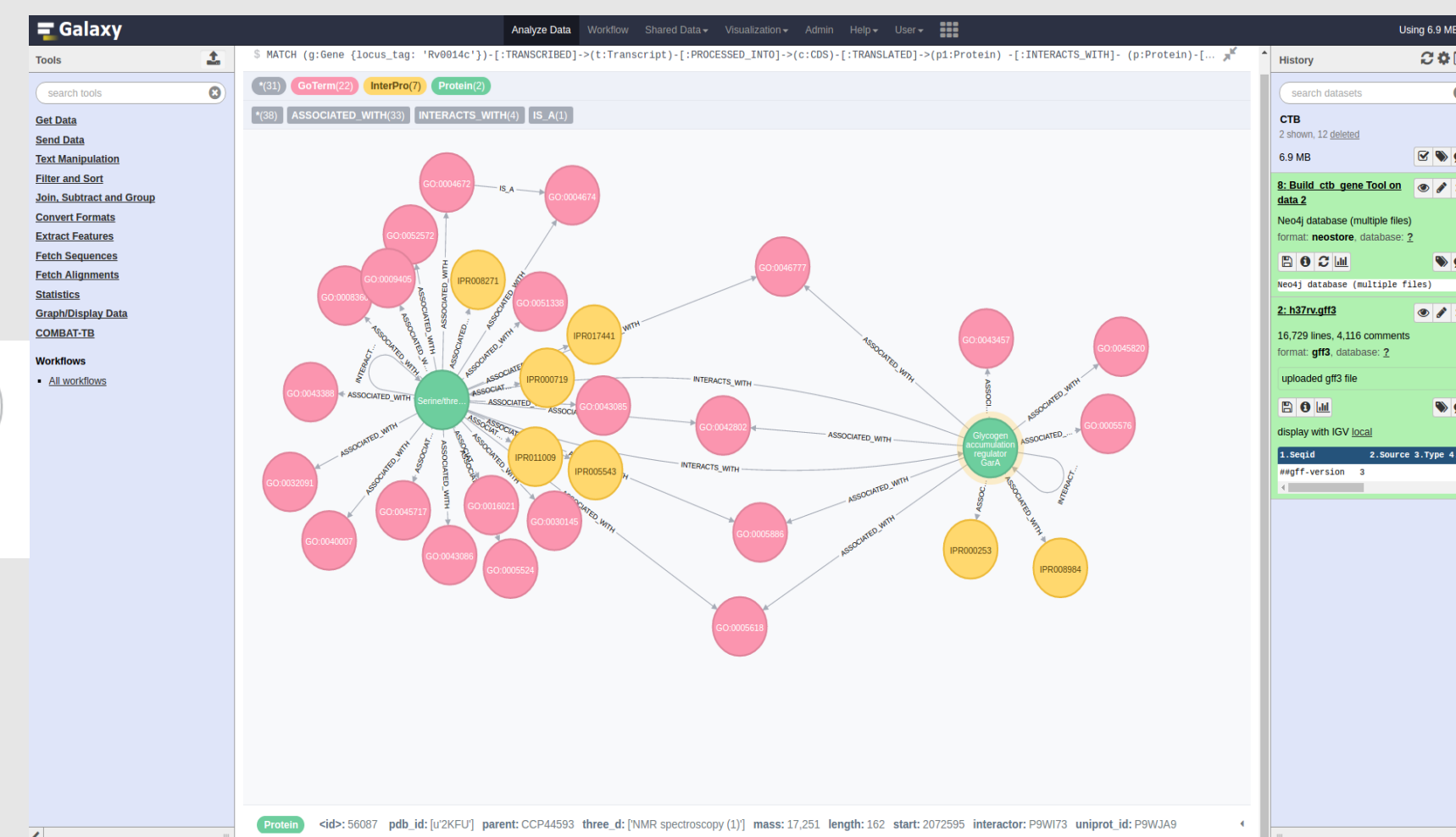


Results

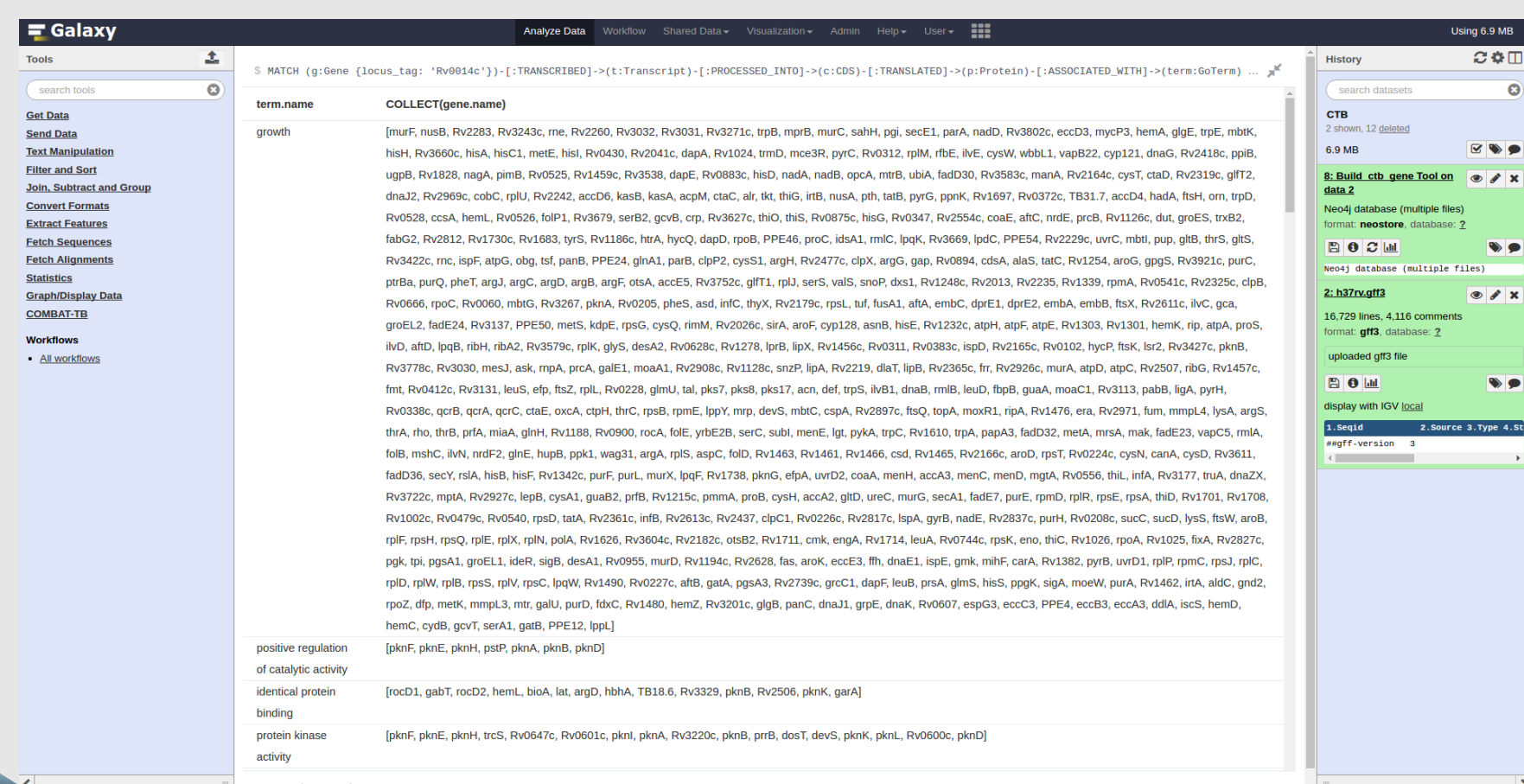
A Neo4j GIE to explore the graph database using Cypher, a declarative query language designed for expressing graph queries. Thus, allowing logical querying across entities.



```
$ MATCH (g:Gene {locus_tag: 'Rv0814c'})-[:TRANSCRIBED]->(t:Transcript)-[:PROCESSED_INT0]->(c:CDS)-[:TRANSLATED]->(p1:Protein)-[:INTERACTS_WITH]- (p:Protein)-[:ASSOCIATED_WITH]->(term) RETURN p, term
```



```
1 MATCH (g:Gene {locus_tag: 'Rv0814c'})-[:TRANSCRIBED]->(t:Transcript)-[:PROCESSED_INT0]->(c:CDS)-[:TRANSLATED]->(p1:Protein)-[:ASSOCIATED_WITH]->(term:GoTerm) WITH COLLECT(term) AS terms UNWIND terms AS term
2 MATCH (gene:Gene)-[:TRANSCRIBED]->(c:CDS)-[:TRANSCRIBED]->(p:Protein)-[:INTERACTS_WITH]->(term:GoTerm) RETURN term.name, COLLECT(term.name)
```



Conclusion

The development of the Neo4j GIE bridges the gap between Galaxy and graph databases by allowing users to interrogate, using the Cypher declarative language, data without leaving Galaxy.

References



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