

neoGERM: Mining Graph Evolution Rules in Neo4j

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Motivation

With the increasing popularity of Web large network data can be retrieved to study the evolution of social networks and information flow at microscopic level. The evolution of such network data can be represented in one single structure (*time evolving graph*) since



the data in these networks does not disappear with time but rather grows.

Subgraph Isomorphism

Support calculation requires solving subgraph isomorphism problem.

Current implementation of GERM uses Ullmann's algorithm (Ullmann, 1976).

gSpan

- gSpan is developed for *transactional setting* (Yan and Han, 2002)
- By constructing DFS Code Tree patterns are taken from the original graph and not generated randomly
- By introducing DFS Lexicographic Order search space is efficiently pruned



DFS Code

Host Graph

(4)

2

9

- **DFS Code** $\alpha = (a_0, a_1, ..., a_m)$, where $a_k = (i_k, j_k, l_{i_k}, l_{(i_k, j_k)}, l_{j_k})$
- DFS Lexicographic Order is a lexicographic combination of DFS order over edges and full order over node labels



DFS Code encodes a pattern, but not uniquely

GERM vs neoGERM

- GERM adapts gSpan to one single graph setting
- GERM works only for undirected graphs
- GERM requires data to be stored as document
- GERM allows only integer numbers as labels over nodes

 neoGERM requires data to be stored in Neo4j and incorporates in-built methods from Neo4j
neoGERM works for directed graphs

 neoGERM works with node labels as set of integers and redefines pattern for partial matching

neoGERM further improves support calculation



Neo4j

Neo4j is a **NoSQL graph database** (www.neo4j.org). It has a traversal framework for high speed traversals on the nodes. Gephi has a plug-in to visualise databases from Neo4j.

Neo4j is an open source project in a GPLv3 Community edition with Advanced and Enterprise editions available under both AGPLv3 and commercial licenses.

Initial Results

Venues as labels



Related Work

- Global evolution of networks: (Leskovec, Kleinberg, Faloutsos, 2005) (Backstrom, Huttenlocher, Kleinberg, Lan, 2006)
- Microscopic evolution: (Lescovec, Backstrom, Kumar, Tomkins, 2008)
- Link prediction algorithms like Adamic-Adar/Liben-Nowell and Kleinberg are capable to predict links between existing nodes.

Future Work

- Experiment on different datasets with different settings.
- Further improve support calculation.
- Develop link prediction based upon discovered graph evolution rules.
- Investigate neoGERM scalability and performance on larger datasets.