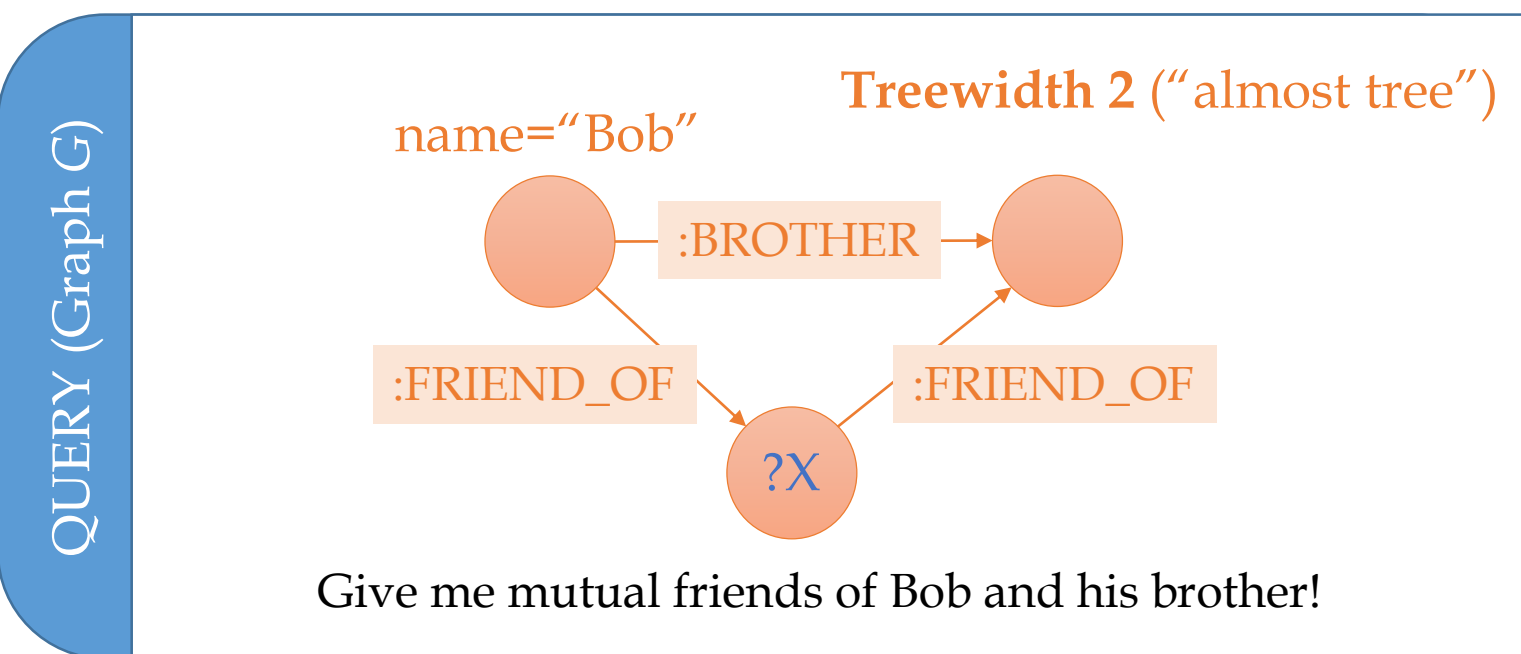
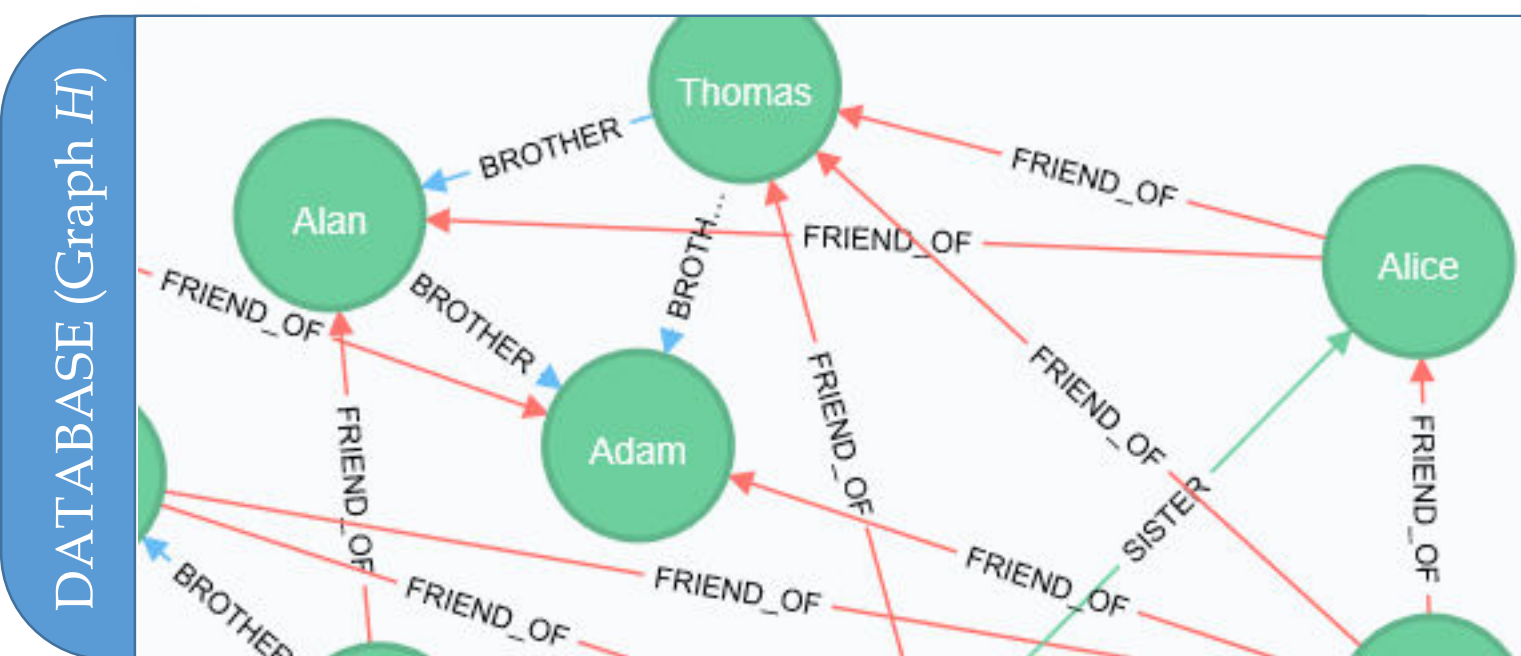


Graph databases are a promising branch of storage systems. We have proposed a new algorithm for querying such databases with the aim to ensure the best possible worst-case time complexity. Our approach is based on the recent results from the field of graph homomorphism and uses them in the context of graph databases. By a series of experiments we have shown that our algorithm is faster than the leading graph database Neo4j in some of the scenarios.

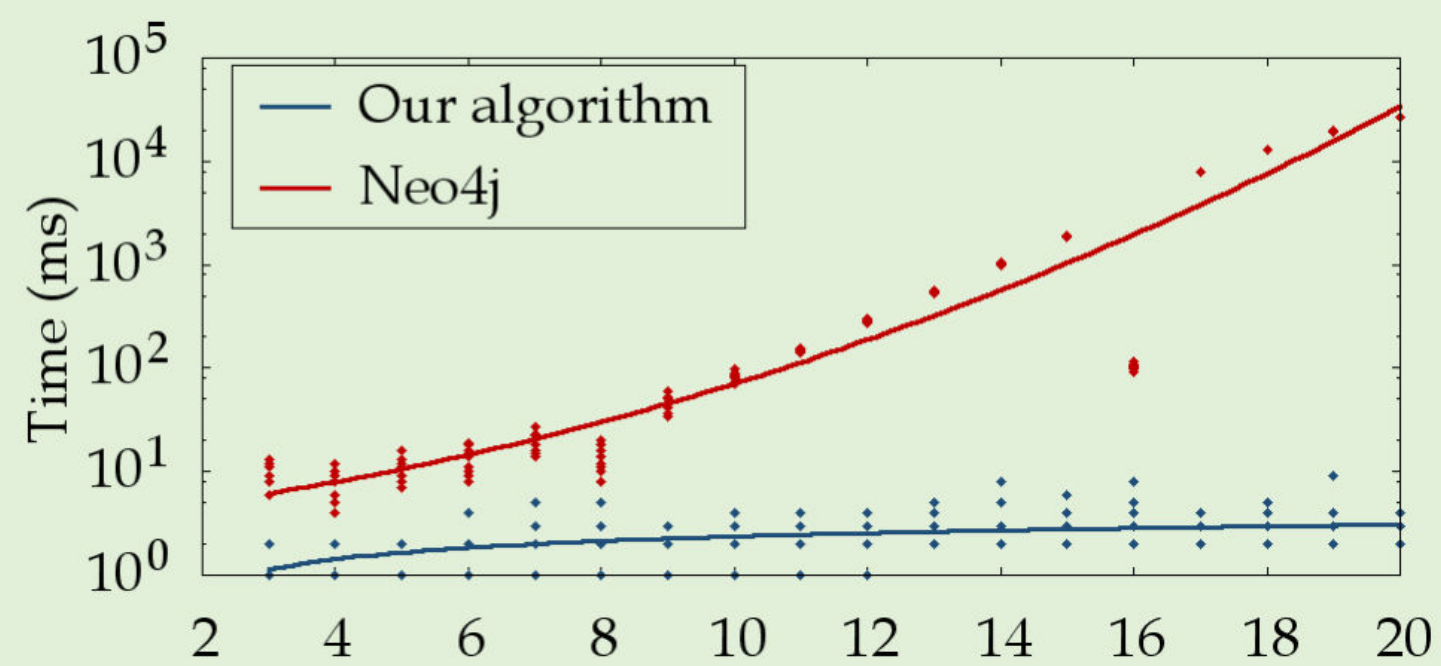
ASK THE GRAPH



Homomorphism

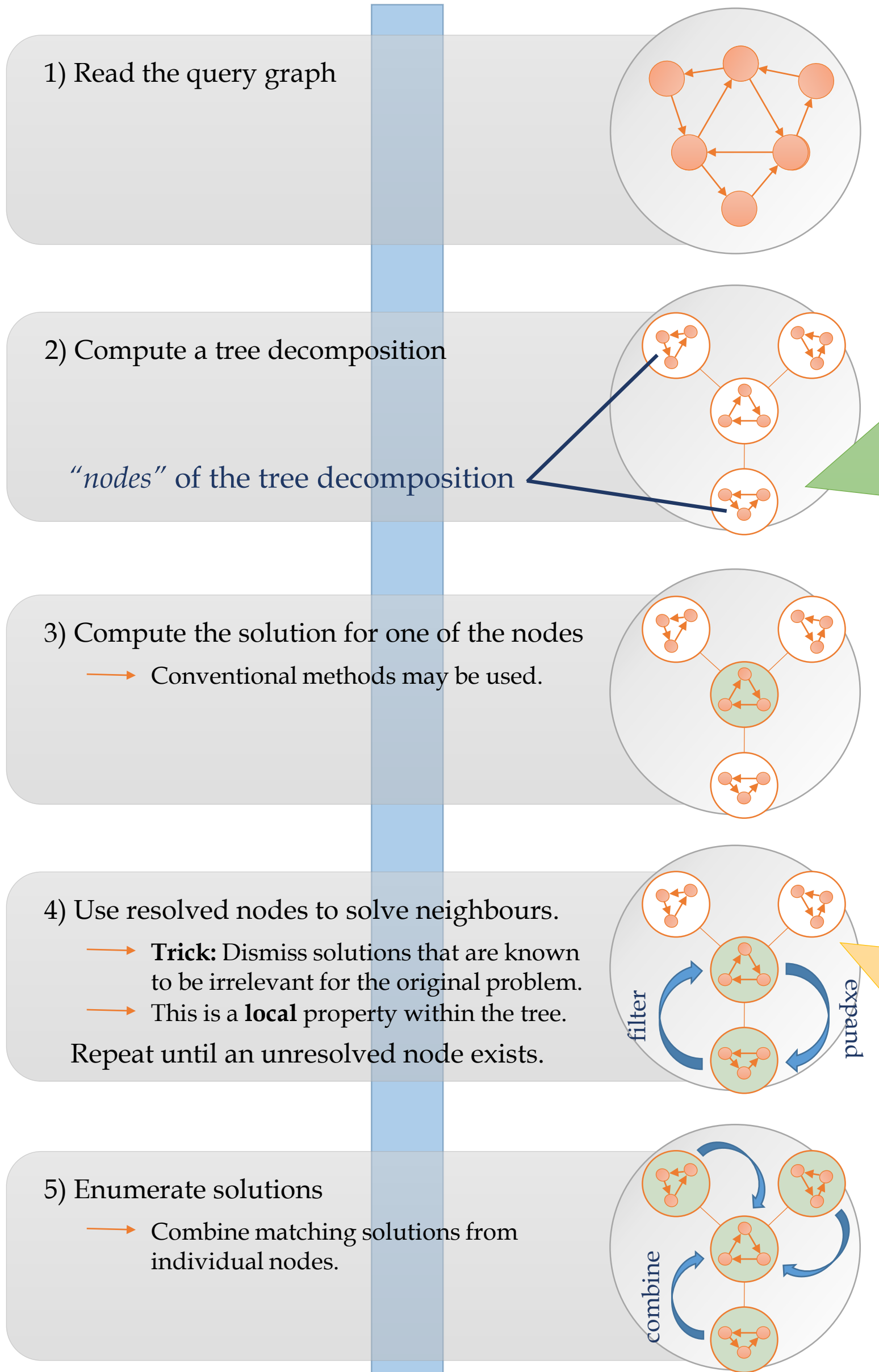


BEST RESULT



Finding directed cycles of an increasing length in a DAG

SOLUTION



TREewidth

- Measures similarity of a graph to a tree.
- The size of the largest portion of a graph, that has to be treated together

Treewidth 0	Treewidth 1	Treewidth 2
(without edges)	(trees)	(series parallel graphs)

KNOWN RESULTS

For simple undirected graphs the decision problem (i.e. decide whether a homomorphism exists) is in:

- **NP-C** for unrestricted queries (unless database is a bipartite graph) [1990, Hell and Nešetřil]
- **PTIME** for queries of bounded treewidth
- **PTIME** iff queries have bounded treewidth modulo homomorphic equivalence (decision variant only) [2002, Dalmau et al. ; 2007, Grohe]

$$O(V(T) \cdot tw(G) \cdot V(H)^{tw(G)+1} + OUT)$$

$V(T)$ - number of nodes, $tw(G)$ - treewidth of the query, $V(H)$ - number of vertices in the database, OUT - number of results

Exponential only in the treewidth