Improving Distributed Subgraph Matching Algorithm on Timely Dataflow*

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Abstract—The subgraph isomorphism problem is a well-known NP-hard problem: given a query graph q and data graph G, we need to find all subgraphs matching q in G. Subgraph isomorphism plays a vital role in the fields of e-commerce, social media and biological science. One of the most efficient and scalable distributed subgraph matching algorithm is SEED [1]. While most of the graphs in real applications are attributed graphs, SEED does not propose a cost evaluation strategy for them, which limits its application in practice. In the meanwhile, SEED is based on Map-Reduce, by which the performance of the algorithm can be greatly affected because of the frequent I/O operations when handling intermediate results in the multi-round join process. Discovering the limitations of SEED, this paper concerns with designing a cost evaluation strategy and optimized join plan for attributed graphs in a distributed context. Whats more, this paper implements SEED on Timely Dataflow instead of Map-Reduce to avoid frequent I/O operations. The generality and performance of SEED can be significantly improved after solving these two limitations. This paper also carries out large scale experiments, whose results show that the algorithm has high performance and excellent scalability.

Index Terms—subgraph matching, distributed algorithm, dataflow

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- The word "data" is plural, not singular.
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An excellent style manual for science writers is [7].

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TABLE I TABLE TYPE STYLES

Table	Table Column Head		
Head	Table column subhead	Subhead	Subhead
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^aSample of a Table footnote.

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Fig. 1. Example of a figure caption.

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