

In-depth Analysis of Densest Subgraph Discovery in a Unified Framework

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ABSTRACT

As a fundamental topic in graph mining, Densest Subgraph Discovery (DSD) has found a wide spectrum of real applications. Several DSD algorithms, including exact and approximation algorithms, have been proposed in the literature. However, these algorithms have not been systematically and comprehensively compared under the same experimental settings. In this paper, we first summarize a unified framework to incorporate all DSD algorithms from a highlevel perspective. We then extensively compare representative DSD algorithms over a range of graphs - from small to billion-scale and examine the effectiveness of all methods, providing a thorough analysis of DSD algorithms. As a byproduct of our experimental analysis, we are also able to identify new variants of the DSD algorithms over undirected graphs, by combining existing techniques, which are up to 10× faster than the state-of-the-art algorithm with the same accuracy guarantee. Finally, based on the findings, we offer promising research opportunities. We believe that a deeper understanding of the behavior of existing algorithms can provide new valuable insights for future research.

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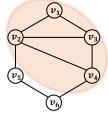
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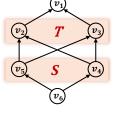
The source code, data, and/or other artifacts have been made available at https://github.com/JayLZhou/DensestSubgraph.

1 INTRODUCTION

Graph data are often used to model relationships between objects in various real-world applications [2, 19, 45, 48, 61, 96, 97]. For example, the Facebook friendship network can be modelled as an undirected graph by mapping users to vertices and friendships to edges [19]; In X (formerly known as Twitter), a directed edge can represent the "following" relationship between two users [45]; The

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(a) An undirected graph G

(b) A directed graph D

Figure 1: Examples of undirected and directed graphs.

Web network itself can also be modelled as a vast directed graph [2]. Figures 1 (a) and (b) depict an undirected graph and a directed graph respectively.

As a fundamental problem in graph mining, the densest subgraph discovery (DSD) problem aims to discover a very "dense" subgraph from a given graph [52, 58]. More precisely, given an undirected graph, the DSD problem [38] asks for a subgraph with the highest density, defined as the number of edges over the number of vertices in the subgraph, and it is often termed the densest subgraph (DS). The DSD problem lies at the core of graph mining [7, 37], and is widely used in many areas. For instance, in social networks, the DS discovered can be used to detect communities [18, 88], reveal fake followers [10], and identify echo chambers and groups of actors engaged in spreading misinformation [51, 58]. In e-commerce networks [10], the DS is useful for detecting fake accounts. In graph databases, the DSD is a building block for solving many graph problems, such as reachability queries [20] and motif detection [33, 80]. In biological data analysis, DSD solutions have been shown to be useful in identifying regulatory motifs in genomic DNA [33] and gene annotation graphs [80]. Besides, the DSD problem is closely related to other fundamental graph problems, such as network flow and bipartite matching [82]. Due to the theoretical and practical importance, researchers from the database, data mining, computer science theory, and network communities have designed efficient and effective solutions to the DSD problem.

In Table 1, we categorize representative DSD algorithms by their computation models, main methods, and approximation ratio guarantee. The exact algorithms include maximum flow and convex programming-based approaches; the approximation algorithms include peeling-based, convex programming-based, and network flow-based approaches. After a careful literature review, we make the following observations. First, no prior work has proposed a unified framework to abstract the DSD solutions and identify key

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Table 1: Classification of existing DSD works.

Casala			Complexit	y	1		Oı	ptimization	
Graph type	Algorithm	Key technique	Time complexity	Space complexity	Approx. ratio	# of Iteration	Early termination	Graph reduction	Parallel friendly
	FlowExact [38]	Network flow	$O(\log n \cdot t_{\text{Flow}})$	O(m)	1	N/A	X	Х	Х
	CoreExact [31]	Network flow	$O(\log n \cdot t_{\mathrm{Flow}})$	O(m)	1	N/A	1	1	Х
	FWExact [24]	Convex programming	$O(T \cdot m + \log T \cdot t_{\text{Flow}})$	O(m)	1	$\Omega(T)$	1	×	1
	MWUExact [40]	Convex programming	$O(T \cdot m + \log T \cdot t_{\text{Flow}})$	O(m)	1	$\Omega(T)$	1	×	1
	FISTAExact [40]	Convex programming	$O(T \cdot m + \log T \cdot t_{\text{Flow}})$	O(m)	1	$\Omega(T)$	1	Х	1
Undirected	Greedy [16]	Peeling	O(m+n)	O(m)	2	N/A	X	Х	X
graphs	CoreApp [31]	Peeling	O(m+n)	O(m)	2	N/A	X	X	X
grapiis	Greedy++ [14]	Peeling	$O(T \cdot m \log n)$	O(m)	$(1+\epsilon)$	$\Omega\left(\frac{\Delta(G)\cdot\log m}{\rho_G^*\cdot\epsilon^2}\right)$	×	×	×
	FWApp [24]	Convex programming	$O(T \cdot m)$	O(m)	$(1 + \epsilon)$	$\Omega\left(\frac{m\Delta(G)}{\epsilon^2}\right)$	1	×	1
	MWUApp [40]	Convex programming	$O(T \cdot m)$	O(m)	$(1+\epsilon)$	$\Omega\left(\frac{m\Delta(G)}{\epsilon^2}\right)$	1	×	1
	FISTAApp [40]	Convex programming	$O(T \cdot m)$	O(m)	$(1+\epsilon)$	$\Omega\left(\frac{\sqrt{m\Delta(G)}}{\epsilon}\right)$	1	×	1
	FlowApp*[92]	Network flow	$O(T \cdot m \log m)$	O(m)	$(1+\epsilon)$	$\Omega(\frac{\log m}{\epsilon})$	1	1	Х
	DFlowExact [50]	Network flow	$O(n^2 \cdot \log n \cdot t_{\text{Flow}})$	O(m)	1	N/A	X	Х	Х
	DCExact [65]	Network flow	$O(h \cdot \log n \cdot t_{\text{Flow}})$	O(m)	1	N/A	1	1	Х
Directed	DFWExact [66]	Convex programming	$O(T \cdot t_{\mathrm{Flow}})$	O(m)	1	$\Omega(T)$	1	1	1
graphs	DGreedy [16]	Peeling	$O(n^2 \cdot (n+m))$	O(m)	2	N/A	X	X	X
grapiis	XYCoreApp [65]	Peeling	$O(\sqrt{m}\cdot(n+m))$	O(m)	2	N/A	X	Х	Х
	WCoreApp [59]	Peeling	$O(\Delta(D)\cdot(n+m))$	O(m)	2	N/A	Х	Х	/
	DFWApp [66]	Convex Programming	$O(T \cdot m)$	O(m)	$(1+\epsilon)$	$\Omega\left(\frac{m \cdot \kappa}{\epsilon^2}\right)$	1	1	1

^{*} Note: n and m denote the numbers of vertices and edges in the graph respectively; $\Delta(D)$ is the maximum value of highest out-degree and in-degree of directed graph D.

performance factors. Second, existing works focus on evaluating the overall performance, but not individual components. Third, there is no existing comprehensive comparison between all these algorithms.

Our work. To address the above issues, in this paper we conduct an in-depth study on sequential DSD algorithms. We first propose a unified framework with three modules, namely graph reduction, vertex weight update (VWU), and candidate subgraph extract and verify (CSV), which capture the core ideas of all existing algorithms. Given a graph G and an error threshold ϵ , graph reduction aims to locate the DS in a small subgraph; VWU aims to update vertex weights over T iterations; and CSV extracts a candidate subgraph based on vertex weights and verifies if it satisfies the ϵ error requirement. Under this framework, we systematically compare 12 (resp. 7) representative algorithms for undirected (resp. directed) graphs, respectively. We conduct comprehensive experiments on both real-world and synthetic datasets and provide an in-depth analysis.

In summary, our principal contributions are as follows.

- Summarize a unified framework for DSD solutions from a high-level perspective (Section 3).
- Comprehensively examines DSD algorithms for both undirected and directed graphs respectively (Sections 4 and 5).
- Conduct extensive experiments from different angles using various datasets, providing a thorough analysis of DSD algorithms. Based on our analysis, we identify new variants of DSD algorithms over undirected graphs, by combining existing techniques, which significantly outperform the state-of-the-art (Section 6).
- Summarize lessons learned and propose practical research opportunities that can facilitate future studies (Section 7).

In Section 2, we present the preliminaries and introduce a unified DSD framework in Section 3. Section 8 reviews related work while Section 9 summarizes the paper.

PRELIMINARIES

In this section, we provide the definitions of DSD problems over both undirected and directed graphs, i.e., UDS and DDS problems respectively.

2.1 Problem definitions

Table 2: Notations and meanings.

Notation	Meaning
G = (V, E)	An undirected graph with vertex set V and edge set E
D = (V, E)	A directed graph with vertex set V and edge set E
N(v,G)	The set of neighbors of a vertex v in G
$d_G(v)$	The degree of v in G , i.e., $d_G(v) = N(v, G) $
$d_{D}^{+}(v), d_{D}^{-}(v)$	The out-degree and in-degree of v in D , respectively
G[S]	The subgraph of <i>G</i> induced by vertices in <i>S</i>
D[S,T]	The subgraph of D induced by vertices in S and T
$\mathcal{D}(G)$	The densest subgraph of <i>G</i>
$\rho(S,T)$	The density of subgraph $D[S, T]$
k^*	The largest k such that the k -core in G exists.
$\Delta(G)$	The highest degree of <i>G</i> .

We denote an undirected graph by G = (V, E), where |V| = n and |E| = m are the numbers of vertices and edges of G, respectively. The set of neighbors of a vertex u in G is denoted by N(u, G), and the degree of u is $d_G(u) = |N(u,G)|$. Given a vertex set S, we use G[S] = (S, E(S)) to denote the subgraph of G induced by S, where $E(S) = \{(u, v) \in E \mid u, v \in S\}$ denotes the set of edges in G contained

We refer readers to [84] for a comparison of parallel DSD algorithms.

^{*} Note: $\Delta(G)$ denotes the highest degree of undirected graph G; κ is an integer proportional to the maximum value of highest out-degree and in-degree of directed graph D.

in S. For a given undirected graph H, we denote its sets of vertices and edges by V(H) and E(H), respectively.

Definition 2.1 (**Density of undirected graph** [38]). Given an undirected graph G = (V, E), its density $\rho(G)$ is defined as the number of edges over the number of vertices, i.e., $\rho(G) = \frac{|E|}{|V|}$.

PROBLEM 1 (UDS PROBLEM [31, 38, 87]). Given an undirected graph G, find the subgraph $\mathcal{D}(G)$ whose density is the highest among all the possible subgraphs, which is also called the undirected densest subgraph (UDS).

Let D=(V,E) be a directed graph. For each vertex $v\in V$, denote by $N_D^+(v)$ (resp. $N_D^-(v)$) the out-neighbors (resp. in-neighbors) of v, and correspondingly denote by $d_D^+(v):=|N_D^+(v)|$ (resp. $d_D^-(v):=|N_D^-(v)|$) the out-degree (resp. in-degree) of v. Given two vertex subsets $S,T\subseteq V$ that are not necessarily disjoint, $E(S,T)=E\cap (S\times T)$ denotes the set of all edges from S to T in the graph D. The (S,T)-induced subgraph of D contains the vertex sets S,T and the edge set E(S,T).

Definition 2.2 (**Directed graph density** [47, 50, 65, 67]). Given a directed graph D=(V,E) and two vertex sets S and T, the density of an (S,T)-induced subgraph is defined as $\rho(S,T)=\frac{|E(S,T)|}{\sqrt{|S||T|}}$.

PROBLEM 2 (DDS PROBLEM [7, 16, 37, 47, 50]). Given a directed graph D, find the subgraph $\mathcal{D}(D) = D[S^*, T^*]$ whose corresponding density is the highest among all the possible (S, T)-induced subgraphs, also called the directed densest subgraph (DDS).

Denote the density of $\mathcal{D}(G)$ and $\mathcal{D}(D)$ by ρ_G^* and ρ_D^* respectively. E.g., in Figures 1 (a) and (b), the subgraphs in the dashed ellipses are the UDS and DDS respectively, with ρ_G^* =5/4 and ρ_D^* =2.

3 A UNIFIED FRAMEWORK

In this section, we develop a unified framework, consisting of three stages: *Graph reduction, Vertex Weight Update* (VWU), and *Candidate subgraph Extract and Verify* (CSV), which can cover all existing UDS and DDS algorithms, as shown in Algorithm 1.

Algorithm 1: A unified framework of DSD

```
input : G = (V, E), \epsilon, T
output: An exact/approximate densest subgraph \mathcal{D}(G)

1 f \leftarrow \text{False}; \underline{\rho} \leftarrow k^*/2; \overline{\rho} \leftarrow k^*; \mathbf{w} \leftarrow \emptyset;

2 repeat

| // (1) The graph reduction method.

3 G \leftarrow \text{ReduceGraph}(G, \underline{\rho});

| // (2) The vertex weight update method.

4 \mathbf{w} \leftarrow \text{VWU}(G, \mathbf{w}, T, \underline{\rho}, \overline{\rho});

| // (3) The candidate subgraph extract and verify.

5 (f, \mathcal{D}(G), \underline{\rho}, \overline{\rho}) \leftarrow \text{CSV}(G, \underline{\rho}, \overline{\rho}, \mathbf{w}, \epsilon);

6 until f = \text{True};

7 return \mathcal{D}(G);
```

Specifically, given a graph G, an approximate ratio ϵ , and the number of iterations T, we initialize the upper and lower bounds of ρ_G^* (line 1), where k^* is the maximum core number which will be introduced later. We then iteratively execute operations in the following three stages (lines 2-6):

(1) Locate the graph into a smaller subgraph (i.e., $\lceil \underline{\rho} \rceil$ -core) utilizing the lower bound $\underline{\rho}$ (Section 4.1);

- (2) Update the vertex weight vector **w** for each vertex over *T* iterations (Section 4.2);
- (3) Extract the candidate subgraph using vertex weight vector w, update upper and lower bounds of ρ^{*}_G, and verify if the candidate subgraph meets the requirements (Section 4.3). Terminate the process if it does; otherwise, update the parameters and repeat the above steps.

In Table 3, we illustrate the details of these three stages for each category of DSD algorithms for undirected graphs. For example, for CoreExact, it discovers UDS via binary search, for each search process: **Stage (1)** locate graph into a smaller *k*-core; **Stage (2)** compute the maximum flow on this smaller graph, and **Stage (3)** verify the result optimality via minimum cut. The vertex weight vector **w** represents the weights assigned to different vertices. It holds different meanings and serves different purposes in different algorithms:

- In the network flow-based algorithms, **w** denotes the flows from the vertices to the target node;
- In the CP-based algorithms, w represents the weight sum received by each vertex;
- In the peeling-based algorithms, w is used to select which vertex should be removed.

Hence, computing the maximum flow, optimizing the CP formulation, and peeling vertices are exactly the vertex weight update process. Our abstraction unifies the different uses of weights in different algorithms. Due to space limits, we present an overview of the DDS problem in our technical report [78].

Algorithm 2: FWExact under our unified framework

```
input : G = (V, E), \epsilon = 0, T output: An exact densest subgraph \mathcal{D}(G)

1 f \leftarrow \mathsf{False}; \underline{\rho} \leftarrow k^*/2; \overline{\rho} \leftarrow k^*;

2 repeat

3 G \leftarrow \mathsf{ReduceGraph}(G, \underline{\rho});

4 \mathbf{w} \leftarrow \mathsf{FWExactVWU}(G, \mathbf{w}, T, \underline{\rho}, \overline{\rho}) (Algorithm 4);

5 (f, \mathcal{D}(G), \underline{\rho}, \overline{\rho}) \leftarrow \mathsf{FWExactCSV}(G, \underline{\rho}, \overline{\rho}, \mathbf{w}, \epsilon) (Algorithm 6);

6 until f = \mathsf{True};

7 return \mathcal{D}(G);
```

Here, we provide an example to illustrate how to run FWExact using the three steps of our unified framework, as shown in Algorithm 2. The other CP-based algorithms are almost the same as FWExact, due to the space limit, we only present FWExact here, and we also show an example about FlowExact in our technical report [78]. In the VWU stage of FWExact, it repeats T iterations to optimize the CP formulation of the UDS problem (see Algorithm 4). In the CSV stage of FWExact, the vertices with higher weights are more likely to appear in the densest subgraph $\mathcal D$, since they are linked by more edges (see Algorithm 6).

4 COMPARISON AND ANALYSIS OF DSD ALGORITHMS FOR UNDIRECTED GRAPHS

In this section, we systematically compare and analyze all the UDS algorithms in terms of the three stages of our unified framework.

4.1 Graph reduction

We first review the notion of *k-core*.

Table 3: Overview of the three stages of the existing UDS algorithm

Method	Stage (1): ReduceGraph	Stage (2): VWU	Stage (3): CSV
FlowExact	no reduction	compute the maximum flow	0 extract the minimum cut;
CoreExact	locate graph into $\lceil \underline{\rho} \rceil$ -core	compute the maximum now	2 verify if it is optimal;
FlowApp	lacata granh into [a] cara	perform the blocking flow	extract the residual graph;
FlowApp	locate graph into $\lceil \underline{\rho} \rceil$ -core	perform the blocking now	2 verify the approximation ratio;
CP-based	no reduction	optimize CP(<i>G</i>)	extract the maximum prefix sum set;
Cr-baseu	no reduction	optimize CF(G)	2 verify if it is exact or satisfies the approximation ratio criteria;
Peeling-based	no reduction	iteratively remove vertices	0 extract the subgraph with the highest density during the peeling process;

Definition 4.1 (k-core [31, 38, 87]). Given an undirected graph G and an integer k ($k \ge 0$), its k-core, denoted \mathcal{H}_k , is the largest subgraph of G, such that $\forall v \in \mathcal{H}_k$, $deg_{\mathcal{H}_k}(v) \ge k$.

The *core number* of a vertex $v \in V$ is the largest k for which a k-core contains v; the maximum core number among all vertices is denoted k^* . A k-core has an interesting "nested" property [9]: for any two non-negative integers i and j s.t. i < j, $\mathcal{H}_i \subseteq \mathcal{H}_i$.

CoreExact [31] locates the UDS into some k-cores. Specifically, the $\mathcal{D}(G)$ is contained in the $\lceil \rho_G^* \rceil$ -core, however, since ρ_G^* is unknown in advance, we cannot use it directly.

Instead, we locate the UDS into some k-cores using a lower bound on ρ_G^* , thanks to the nested property of k-cores.

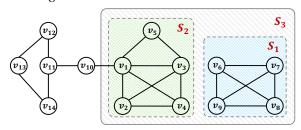


Figure 2: An example of the core-based graph reduction.

Example 1. For example, for the undirected graph in Figure 2, suppose the lower bound of ρ_G^* is 3. Then, the UDS can be located in the 3-core, i.e., S_1 and S_2 , which is smaller than the entire graph.

Due to the nested property of k-core, vertices with smaller core numbers in the remaining graph can be successively removed in the search process thus gradually reducing the size of the graph, since as we shall see, the lower bound of ρ_G^* is progressively increasing. Besides, as shown in [31], the ρ_G^* cannot be larger than k^* and cannot be smaller than $k^*/2$.

4.2 Vertex weight updating

We show that the key components in various UDS algorithms can be considered as the process of vertex weight updating.

Algorithm 3: The template of VWU

- 1 **1** initialize **w** and auxiliary variables;
 - // Algorithms have varied stop conditions.
- 2 while stop condition is not met do
- 3 update the auxiliary variables;

We outline the VWU template in Algorithm 3, which begins by initializing the vertex weight vector **w** along with auxiliary variables facilitating the update process of **w**. We update the variables until the stop condition is met, where in each iteration, the auxiliary variables and **w** are updated (lines 2-3). The different algorithms have various stop conditions in the while-loop, such as for FlowExact, the stop condition is the maximum flow is reached. We summarize the stopping conditions of the while-loop for different DSD algorithms in our technical report [78].

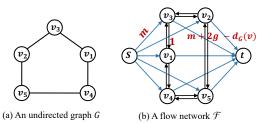


Figure 3: An undirected Graph G and its flow network \mathcal{F} .

- Network flow-based algorithms. For the exact algorithms, FlowExact and CoreExact need to 1 build a flow network based on the guessed maximum density g (i.e., $g = (\rho + \overline{\rho})/2$). For lack of space, we omit the detailed steps of building the flow network [38]. For example, Figure 3 shows a flow network of an undirected graph. We use auxiliary variables to denote the capacity and flow of each edge, and $\mathbf{w}(v)$ represents the value of flow from vertex v to the sink node t. After constructing the flow network, they try to update the flows of some edges in \mathcal{F} , and 2 increase \mathbf{w} . The above process is repeated until the maximum flow is reached. CoreExact follows the same steps as FlowExact, but it utilizes k-core for graph reduction (refer Section 4.1). Unlike exact algorithms, FlowApp does not need to calculate the exact maximum flow, it only needs to perform partial maximum flow computations.
- **CP-based algorithms.** We first introduce the well-known CP formulation of the UDS problem [24] as follows:

$$CP(G) \quad \min \sum_{u \in V} \mathbf{w}^{2}(u)$$
s.t.
$$\mathbf{w}(u) = \sum_{(u,v) \in E} \alpha_{u,v}, \quad \forall u \in V$$

$$\alpha_{u,v} + \alpha_{v,u} = 1, \quad \forall (u,v) \in E$$

$$\alpha_{u,v} \ge 0, \alpha_{v,u} \ge 0 \quad \forall (u,v) \in E$$

$$(1)$$

This CP(G) can be visualized as follows. Each edge $(u, v) \in E$ has a weight of 1, which it wants to assign to its endpoints: u and v such that the weight sum received by the vertices is as even as possible.

Indeed, after a sufficient number of weight update iterations, we can derive an optimal solution to formulation 1, by inducing the subgraph with vertices of the highest weights based on \mathbf{w} , and the weight of any vertex in this subgraph equals the density of UDS. Following this intuition, $\alpha_{u,v}$ in $\mathsf{CP}(G)$ indicates the weight assigned to u from edge (u,v), and $\mathbf{w}(u)$ is the weight sum received by u from its adjacent edges.

All CP-based algorithms aim to solve CP(G) in Eq. (1). There are three widely used types of CP solvers for the DSD problem: Frank-wolfe, MWU, and FISTA-based algorithms. All these algorithms, no matter whether they are designed for approximation or exact solutions, share the same VWU procedure.

Algorithm 4: VWU of Frank-wolfe and MWU

```
1 ① initialize w and auxiliary variables;

2 foreach (u, v) \in E do \alpha_{u,v}^{(0)} \leftarrow 1/2; \alpha_{v,u}^{(0)} \leftarrow 1/2; t \leftarrow 1;

3 foreach u \in V do \mathbf{w}^{(0)}(u) \leftarrow \sum_{(u,v) \in E} \alpha_{u,v}^{(0)};

4 while t \leq T do

5 ② update the auxiliary variables;

6 foreach (u, v) \in E do update \widehat{\alpha}_{u,v} and \widehat{\alpha}_{v,u};

7 \alpha^{(t)} \leftarrow (1 - \gamma_t) \cdot \alpha^{(t-1)} + \gamma_t \cdot \widehat{\alpha}, with \gamma_t = \frac{2}{t+2} or \frac{1}{t+1};

8 ③ update w via \alpha;

9 foreach v \in V do \mathbf{w}^{(t)}(u) \leftarrow \sum_{(u,v) \in E} \alpha_{u,v}^{(t)};

10 t \leftarrow t+1:
```

The Frank-wolfe and MWU-based algorithms solve $\operatorname{CP}(G)$ in an iterative manner, where $\gamma_t = \frac{2}{t+2}$ for Frank-Wolfe algorithms and $\gamma_t = \frac{1}{t+1}$ for MWU-based algorithms. In each iteration, algorithms linearize the objective function at the current point and move towards minimizing it [17, 24, 44]. Algorithm 4 shows the process, \bullet starting with initializing α and \bullet (lines 2-4). Next, in the t-th iteration, \bullet each edge $(u,v) \in E$ attempts to distribute its weight, i.e., 1, to the endpoint with a smaller \bullet wolve, and the \bullet values of all vertices are computed as a convex combination by \bullet (lines 5-7). Then, \bullet \bullet wolve, is updated by the weight sum received by each vertex in V (line 9).

FISTA-based algorithms [40] also adopt the iterative paradigm, but leverage the *projection* operation [73] to speed up the optimization process. The projection operation in each iteration initially proceeds with a gradient descent step to move towards a local minimum. Subsequently, it adjusts the point to ensure it falls within the feasible region by projecting back if necessary. Harb et al. [40] further proved that they converge faster in theory, but experimentally there is a gap between the theoretical conclusion and practical results, i.e., they are slower than other CP-based algorithms as the *projection* is very time-consuming.

• **Peeling-based algorithms.** The key idea of peeling-based algorithms is to find a vertex with the minimum weight, remove it from the graph and update the weights of the remaining vertices. Algorithm 5 presents the details of Greedy and Greedy++. Specifically, when T=1, this algorithm is Greedy, removing the vertex with the minimum degree one by one. Greedy++ algorithm is based on Greedy and iteratively removes the vertices via T iterations. In each iteration, it iteratively removes the vertex with the smallest weight, where the weight of vertex v in each iteration is the sum of its induced degree (w.r.t. the remaining vertices) and the weight of v in the previous iteration (line 6). CoreApp reveals that k^* -core

Algorithm 5: VWU of Greedy and Greedy++

```
1 1 initialize w and auxiliary variables;
2 foreach v \in V do \mathbf{w}^{(0)}(v) \leftarrow 0, H \leftarrow G; t \leftarrow 1;
^{3} while t ≤ T do
         H \leftarrow G;
         repeat
              select vertex v minimizing \mathbf{w}^{(t-1)}(v) + d_H(v);
              2 update the auxiliary variables;
              foreach u \in N(v, H) do d_H(u) \leftarrow d_H(u) - 1;
              \mathbf{3} update \mathbf{w} via d_H;
              \mathbf{w}^{(t)}(v) \leftarrow \mathbf{w}^{(t-1)}(v) + d_H(v);
10
              remove v and all its adjacent edges (u, v) from H;
11
12
         until V(H) \neq \emptyset;
         t \leftarrow t + 1;
13
```

can serve as a 2-approximation solution, where the k^* -core can be computed by following a peeling-based process.

4.3 Candidate subgraph extraction and verification

In this section, we explore the CSV stage across various UDS algorithms and examine how to utilize upper and lower bounds of the optimal density ρ_G^* for verifying results. Let g represent the guessed density, and set $g = (\rho + \overline{\rho})/2$.

- FlowExact and CoreExact. FlowExact and CoreExact need to compute the minimum cut (S, \mathcal{T}) via the maximum flow. If S contains only the source node $\{s\}$, $\overline{\rho}$ is updated to g; otherwise, $\underline{\rho}$ is set to g and $S\setminus \{s\}$ is returned as the candidate subgraph. To verify the quality of the candidate subgraph, FlowExact checks if the difference between $\overline{\rho}$ and $\underline{\rho}$ is less than $\frac{1}{n\cdot(n-1)}$, as the density difference between any two subgraphs must be larger than $\frac{1}{n\cdot(n-1)}$. When the condition is met, the exact solution is found. In contrast, CoreExact uses a less stringent stopping criterion, checking if the density difference is $<\frac{1}{|V_C|\cdot(|V_C|-1)}$, where V_C is the largest connected component in the graph.
- **FlowApp.** FlowApp searches for an augmenting path in \mathcal{F} after h blocking flows. If such a path exists, $\underline{\rho}$ is updated to g and the residual graph of \mathcal{F} is returned as the candidate subgraph; otherwise, $\overline{\rho}$ is updated to g and no candidate subgraph is obtained.

```
Algorithm 6: CSV of CP-based algorithms
```

```
1 foreach 1 \le i \le |V(G)| do
2 u_i \leftarrow the vertex with the i-th highest weight in V(G);
3 G_i \leftarrow the induced subgraph of top-i weight vertices;
4 u_i \leftarrow d_{G_i}(u_i);
5 s^* \leftarrow \arg\max_{1 \le s \le n} \frac{1}{s} \sum_{i=1}^s y_i;
6 S \leftarrow the subgraph induced by the first s^* vertices;
7 \underline{\rho} = \max(\rho(S), \rho); \overline{\rho} = \max_{1 \le i \le n} \min\left\{\frac{1}{i} \binom{i}{2}, \frac{1}{i} \sum_{j=1}^i \mathbf{w}(u_j)\right\};
8 f \leftarrow verify via maximum flow;
9 return f, S, \underline{\rho}, \overline{\rho};
```

FlowApp verifies whether a candidate subgraph is a (1 + ϵ)-approximation solution by checking if $\frac{g-\rho}{2q}<\frac{\epsilon}{3-2\epsilon}$ [92].

• CP-based algorithms. All CP-based algorithms use the vertex weight vector w and the PAVA algorithm [24] to extract the candidate subgraph.

Algorithm 6 presents the details, where the (lines 1-6) are the process of using PAVA to extract the candidate subgraph. Intuitively, the vertices with higher weights are more likely to appear in the densest subgraph $\mathcal{D}(G)$, since they are linked by more edges. Thus, the subgraph induced by the first s* vertices with the largest weights is returned as the candidate subgraph. Here, ρ is updated to $\rho(\mathcal{S})$, and

 $\overline{\rho}$ is updated to $\max_{1 \le i \le n} \min \left\{ \frac{1}{i} {i \choose 2}, \frac{1}{i} \sum_{j=1}^{i} \mathbf{w}(u_j) \right\}$ [24, 85]. After a sufficient number of iterations, CP-based algorithms converge to the optimal solution, which can be verified using maximum flow [24]. For the approximation algorithms, we can use the ratio of $\overline{\rho}$ over $\rho(S)$ to estimate the empirical approximation ratio, as the optimal density is not known in advance [24, 85].

• Peeling-based algorithms. Greedy and Greedy++ extract the highest density subgraph during the process of vertex peeling. In addition, CoreApp utilizes the k^* -core as their returned candidate subgraph. These algorithms do not require updating ρ and $\overline{\rho}$ as they do not need to verify results.

COMPARISON AND ANALYSIS OF DSD ALGORITHMS FOR DIRECTED GRAPHS

The DDS problem is more complicated than the UDS problem because it is an induced subgraph of two vertex sets S and T, which leads to a search space of n^2 possible values of the ratio between the size of the two vertex sets (i.e., c = |S|/|T|) which must be examined when computing the maximum density. Next, we show how to adapt our framework (Algorithm 1) for the DDS problem.

• Exact algorithms. The exact algorithms need to enumerate all possible values of c and compute the DS for each c. For each fixed c, the exact algorithms share the same paradigm with UDS algorithms, indicating that they can be easily incorporated into our framework. Similar to CoreExact, DC-Exact [65] reduces the size of the graph by locating the DDS into some [x, y]-cores [65].

Definition 5.1 ([x, y]-core [65]). Given a directed graph D=(V, y)*E*), the [x, y]-core is the largest (S, T)-induced subgraph D[S, T], which satisfies:

- $\begin{array}{ll} (1) \ \, \forall u \in S, d^+_{D[S,T]}(u) \geq x \text{ and } \forall v \in T, d^-_{D[S,T]}(v) \geq y; \\ (2) \ \, \nexists D[S',T'] \neq D[S,T], \text{ such that } D[S,T] \text{ is a subgraph of} \end{array}$
- D[S', T'], i.e., $S \subseteq S'$, $T \subseteq T'$, and D[S', T'] satisfies (1);

The $[x^*, y^*]$ -core is the [x, y]-core with the largest $x \cdot y$ values. Note that [x, y]-core is an extension of the k-core, as each vertex in the [x, y]-core has at least x out-neighbors and y in-neighbors.

By Theorem 5.6 in [65], we only need to discover DDS in the $\left[\frac{\rho}{2\sqrt{c_r}}, \frac{\sqrt{c_l}\rho}{2}\right]$ -core, where (c_l, c_r) specifies the interval of c values under consideration during a particular stage of the divide-andconquer approach and ρ is the lower bound of the optimal density.

• Approximation algorithms. There are two groups of approximation algorithms, i.e., peeling-based [7, 16, 50, 59, 65] and CP-based [63] algorithms. DGreedy needs to enumerate all c values to obtain the approximation solution, and when a specific c is fixed, it is analogous to Greedy. XYCoreApp focuses on identifying the $[x^*, y^*]$ -core. Each time it fixes one dimension and optimizes

the other dimension via iteratively peeling vertices in the other dimension to find the $[x^*, y^*]$ -core, where the peeling process is similar to Greedy. WCoreApp sequentially eliminates vertices based on the lowest weight, where a vertex's weight is determined by the product of its out-degree and in-degree. The key difference between WCoreApp and Greedy is in how they calculate a vertex's weight. The CP-based algorithms need to update \mathbf{w}_{α} and \mathbf{w}_{β} via two auxiliary variables α and β , whose process is similar to Algorithm 4.

Moreover, DCExact [65] first introduced a divide and conquer method to reduce the number of $\frac{|S|}{|T|}$ values examined from n^2 to h, where theoretically $h \leq \mathit{n}^2,$ but practically $h \ll \mathit{n}^2.$ DFWExact [63] introduced a new divide-and-conquer method, utilizing the relationship between the DDS and the *c*-biased DS to skip searches for certain *c* values in the DSD process.

In addition, Sawlani and Wang [82] showed that the DDS problem can be transformed into $O(\log_{1+\epsilon} n)$ vertex-weighted UDS problems, where the vertex weights are assigned according to $O(\log_{1+\epsilon} n)$ different guesses of $\frac{|S|}{|T|}$. In our study, we test its performance by adapting the SOTA algorithm for the UDS problem.

6 EXPERIMENTS

We now present the experimental results. Section 6.1 discusses the setup. The experimental results of the UDS algorithms and DDS algorithms are reported in Sections 6.2 and 6.3, respectively.

6.1 Setup

Table 4: Undirected graphs used in our experiments.

Dataset	Category		E	ρ
bio-SC-GT (BG)	Biological	1,716	31,564	18.4
econ-beacxc (EB)	Economic	507	42,176	83.2
DBLP (DP)	Collaboration	317,080	1,049,866	3.3
Youtube (YT)	Multimedia	3,223,589	9,375,374	2.9
LiveJournal (LJ)	Social	4,036,538	34,681,189	8.6
UK-2002 (UK)	Road	18,483,186	261,787,258	14.2
WebBase (WB)	Web	118,142,155	881,868,060	7.5
Friendster (FS)	Social	124,836,180	1,806,067,135	14.5

Table 5: Directed graphs used in our experiments.

Dataset	Category		E	ρ
maayan-lake (ML)	Foodweb	183	2,494	13.6
maayan-figeys (MF)	Metabolic	2,239	6,452	2.9
Openflights (OF)	Infrastructure	2,939	30,501	10.4
Advogato (AD)	Social	6,541	51,127	7.8
Amazon (AM)	E-commerce	403,394	3,387,388	8.4
Baidu-zhishi (BA)	Hyperlink	2,141,300	17,794,839	8.3
Wiki-en (WE)	Hyperlink	13,593,032	437,217,424	32.2
SK-2005 (SK)	Web	50,636,154	1,949,412,601	38.5

We use sixteen real datasets from different domains including 8 undirected graphs and 8 directed graphs, which are available on the famous online platforms [75, 76, 79]. Tables 4 and 5 report the statistics of these graphs, where ρ denotes the density (i.e., |E|/|V|) of each graph. Due to space limitations, we present results on twelve datasets here and include additional datasets and results (e.g., the sizes of UDS and DDS obtained by various DSD algorithms) in our technical report [78]. We implement all the algorithms in C++ and run experiments on a machine having an Intel(R) Xeon(R) Gold

Table 6: Comparison of 2-approx. UDS algorithms (Red denotes the most efficient algorithm, and Blue denotes the best accuracy).

Algorithm	EB		DP		YT		LJ		WB		FS	
	time	ratio	time	ratio	time	ratio	time	ratio	time	ratio	time	ratio
FlowApp*	43.8 ms	1.556	126.3 ms	1.001	9.1 s	1.761	2.4 s	1.075	41.0 s	1.085	2095.6s	1.799
CoreApp	1.6 ms	1.182	98.0 ms	1.001	0.7 s	1.139	2.0 s	1.033	41.1 s	1.085	156.0 s	1.065
PKMC	4.3 ms	1.182	104.5 ms	1.001	11.2 s	1.139	8.5 s	1.033	37.8 s	1.085	5546.4 s	1.065
FWApp	11.8 ms	1.072	371.6 ms	1.111	9.1 s	1.004	16.4 s	1.078	383.6 s	1.214	1468.2 s	1.200
MWUApp	2.7 ms	1.001	213.6 ms	1.420	24.8 s	1.001	9.8 s	1.115	281.6 s	1.197	723.7 s	1.029
FISTAApp	3.4 ms	1.000	390.3 ms	1.152	49.1 s	1.016	28.7 s	1.480	2377.0 s	1.105	2053.8 s	1.026
Greedy	0.8 ms	1.000	72.8 ms	1.001	0.6 s	1.000	1.8 s	1.013	37.3 s	1.016	130.6 s	1.000

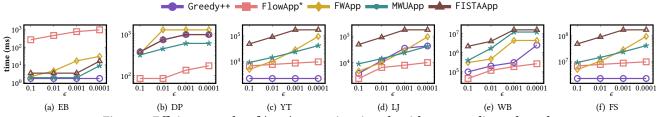


Figure 4: Efficiency results of $(1 + \epsilon)$ -approximation algorithms on undirected graph.

6338R 2.0GHz CPU and 512GB of memory, with Ubuntu installed. If an exact algorithm cannot finish in three days or an approximation algorithm cannot finish in one day, we mark its running time as **INF** in the figures and "—" in the tables.

6.2 Evaluation of UDS algorithms

1. Overall performance. We first report the running time of all 2-approximation algorithms and the actual approximation ratios of approximation solutions returned by each algorithm in Table 6. Here, we set ϵ =1 for $(1+\epsilon)$ -approximation algorithms. We observe that Greedy always achieves the best performance when ϵ =1 in terms of accuracy and efficiency.

We then examine the efficiency of $(1 + \epsilon)$ -approximation algorithms by varying ϵ from 0.1 to 0.0001, and report their running time in Figure 4. Specifically, we report the running time needed by different $(1 + \epsilon)$ -approximation algorithms to find an approximation solution whose density is at least $(1+\epsilon) \times \rho_G^*$. For a fair comparison, we stop algorithms when they achieve a density of $(1 + \epsilon) \times \rho_G^*$, rather than the theoretical number of iterations, since it often takes far fewer iterations in practice than the theoretical value. We observe that FlowApp* and Greedy++ always perform faster than the others, since FlowApp* utilizes the *k*-core-based graph reduction to reduce the search space; Greedy++ uses fewer iterations to obtain higher accuracy, and for each iteration, Greedy++ requires just a straightforward operation: iteratively remove the vertex with the lowest weight. As for CP-based algorithms, FISTAApp often takes more time to obtain solutions with the same accuracy as FWApp and MWUApp, despite theoretically needing fewer iterations. The is because it requires an extra projection operation in each iteration, which is very time-consuming, especially for large graphs. Besides, FWApp and MWUApp achieve the comparable performance. Finally, we present the running time of all exact algorithms in Figure 5. To be specific, CoreExact performs best for the most part, since it can compute the minimum cut on the smaller flow network, and these CP-based algorithms achieve comparable performance.

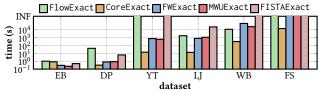


Figure 5: Efficiency results of exact algorithms for UDS.

2. Evaluation of graph reduction. We evaluate the effectiveness of two graph reduction strategies on all algorithms, except 2-approximation algorithms: 1) Single-round reduction, which involves reducing the entire graph to a $k^*/2$ -core once and then executing all algorithms on this smaller graph, instead of the original large graph; and 2) Multi-round reduction, which is applied whenever a tighter lower bound of ρ_G* is identified, as it locates the UDS within a smaller subgraph with a higher core number, resulting in a smaller graph. We present the efficiency of all algorithms and their variants in Figure 6 on two datasets, where the original algorithm names appended with "S" and "M" indicate the adoption of a single-round and multi-round reduction, respectively. We also present the number of remaining edges after each of the first five rounds of graph reductions for all CP-based exact algorithms, on two datasets (Figure 7), wherein the x-axis, "0" denotes the original graph before any reduction, and we record the speedup ratio of graph reduction over different algorithms in our technical report [78].

We make the following analysis: 1) Graph reduction significantly enhances the efficiency of all algorithms. 2) Using multi-round reduction can result in a much greater performance increase than just using single-round reduction in most cases, yet in some cases, the multi-round reduction might bring extra time consumption due to the time needed to perform the reduction itself. For example, applying multi-round reduction can speed up FISTA 1097 times, more than 98 × speedup achieved with a single-round reduction on the YT dataset with $\epsilon = 0.0001$. 3) FISTA, employing multi-round graph reduction outperforms MWU and FW-based algorithms in terms of efficiency while using similar reduction strategies, with

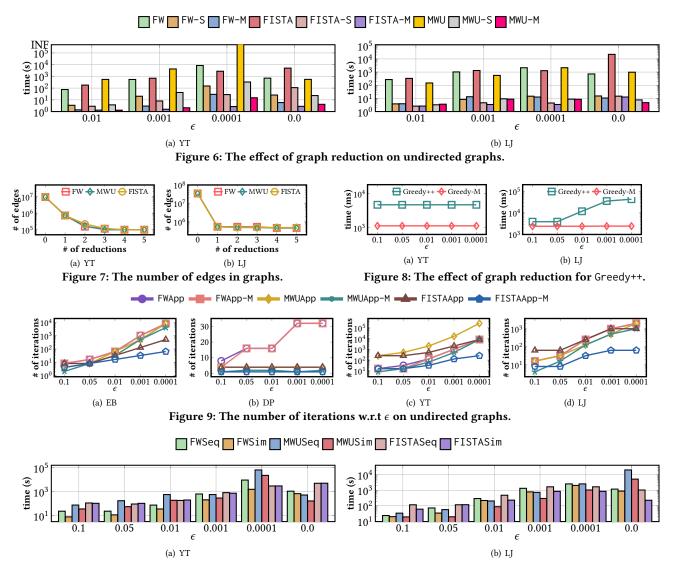


Figure 10: The effect of update strategies on undirected graphs.

the same accuracy. This efficiency boost stems from less projection time on smaller graphs and a reduction in the number of iterations needed. 4) The first graph reduction drastically reduces the graph size, pruning over 98% edges after five rounds of reductions. 5) As shown in Figure 8, graph reduction significantly enhances the performance of Greedy++, achieving an improvement of up to one order of magnitude faster than Greedy++ without graph reduction. 3. Accuracy of CP-based algorithms. In this experiment, we report the number of iterations each algorithm needed to achieve different levels of accuracy in Figure 9. We make the following observations: 1) The actual numbers of iterations needed for all CP-based algorithms are much less than their theoretical numbers. For example, on the YT dataset with $\epsilon = 0.1$, FWApp, MWUApp, and FISTAApp theoretically require at least 10^{16} , 10^{16} , and 10^{8} iterations on the original graph to obtain the approximated solution, while in practice they only need 16, 256, and 256 iterations, respectively; similarly, for the reduced graph, FWAPP-M, MWUApp-M, and FISTAApp-M theoretically require 10¹⁴, 10¹⁴, and 10⁶ iterations, but practically

they also only need 16, 8, and 16 iterations. 2) Graph reduction techniques significantly decrease the number of iterations required by CP-based algorithms, and this phenomenon is especially marked in FISTA-based algorithms because it uses Nesterov-like momentum [40] in its projection phase and adopts $\frac{1}{2 \cdot \Delta(G)}$ as the learning rate, since graph reduction can reduce $\Delta(G)$.

4. Evaluation of weight update strategies. Recall that Danisch [24] employs a simultaneous weight update strategy for FWExact and FWApp. Specifically, within each iteration, if a vertex v's weight $\mathbf{w}(v)$ changes, then the updated vertex weight is promptly visible to subsequent updates of other vertices in the same iteration. The simultaneous weight update strategy enables a more balanced weight distribution among vertices, making the algorithm converge faster, as all the vertices in the UDS have the same weight upon convergence. In this experiment, we test the effect of the vertex weight update strategies, i.e., sequential and simultaneous update strategies, in all CP-based algorithms. As shown in Figure 10, we observe that the algorithms using the simultaneous update strategy

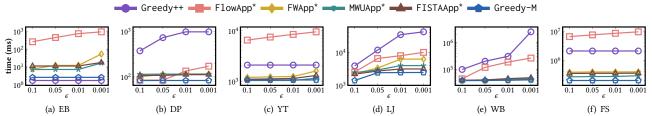


Figure 11: Efficiency results of new $(1 + \epsilon)$ -approximation algorithms on undirected graphs.

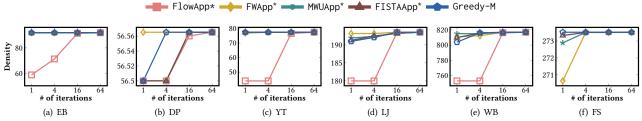


Figure 12: The densities of UDS obtained by the $(1+\epsilon)$ -approximation algorithms.

almost always take less time to achieve the same approximation ratios than the algorithms using the sequential weight update strategy. This is because the simultaneous update strategy makes a more balanced weight distribution.

Based on the above results and discussions, we obtain two major conclusions: 1) Graph reduction is highly beneficial for all algorithms, and employing multi-round reduction often yields better outcomes than single-round reduction. 2) For CP-based algorithms, the simultaneous update strategy typically outperforms the sequential one. By combining existing techniques, we introduce new algorithms which we present and evaluate in the next section.

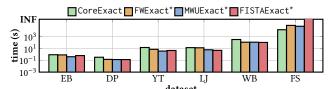


Figure 13: Efficiency of exact algorithms for UDS.

5. New algorithms. Specifically, we apply a simultaneous update strategy and multi-round graph reductions for all CP-based algorithms. We denote modified versions of original algorithms with an asterisk (*). For Greedy++, we enhance efficiency through graph reduction, and denote this variant by Greedy-M. We first evaluate the efficiency of various $(1 + \epsilon)$ -approximation algorithms with the optimal density used in the early stop check. Figure 11 indicates that for smaller datasets, graph reduction is not very useful. However, for large graphs, Greedy-M is always the best one, and those new CP-based algorithms achieve comparable performance. This is because that Greedy-M is a much simpler algorithm than the CPbased ones. Besides, we report how the density of $\mathcal{D}(G)$ changes as the number of iterations increases for the UDS algorithms, as shown in Figure 12. We can make the following observations: (1) All the iteration-based algorithms converge to solutions with densities very close to the optimal in relatively few iterations, and the number of iterations required is significantly lower than the theoretical estimates. (2) The CP-based algorithms and Greedy-M achieve higher densities with fewer iterations compared to FlowApp*. However, as

the number of iterations increases, the solution densities obtained by these algorithms become nearly identical.

We test all CP-based and FlowApp* algorithms by reporting their running time on the four largest datasets. In real situations, the optimal density is not known in advance, so we terminate an algorithm when it meets the early stop conditions that are set according to the specified ϵ , and report the running time in our technical report [78]. Our findings lead to two main insights: firstly, all CP-based algorithms perform similarly well. Secondly, CP-based algorithms are up to one order of magnitude faster than FlowApp*.

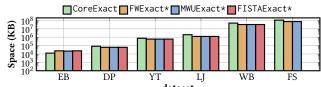
In addition, we compare the efficiency of the newly designed CP-based exact algorithms and CoreExact. As shown in Figure 13, the new algorithms generally require less time than CoreExact except on the FS dataset. After we deeply delve into the FS dataset, we discover a subgraph with a density of 273.518, nearly matching the optimal density of 273.519. We conjecture that if some subgraphs have densities very close to the optimal density, the exact CP-based algorithms often require a larger number of iterations to obtain the optimal solution. To verify it, we synthesize four datasets as follows: 1) we generate two cliques, each with 1,000 vertices, and connect these two cliques by a single edge; 2) we then randomly remove 0.001%, 0.01%, 0.1%, and 1% of the edges from the second clique to simulate if there exists a subgraph with a density close to that of the densest subgraph (i.e., the first clique).

Table 7: The number of iterations.

Method	1%	0.1%	0.01 %	0.001%
FW	128	2,048	32,768	131,072
MWU	128	1,024	16,384	65,536
FISTA	1	128	512	1,024

Table 7 reports the number of iterations required by these algorithms on these datasets. We observe that as the density of the second graph approaches that of a clique, CP-based exact algorithms require a significantly higher number of iterations.

6. Memory usage. The memory costs of all algorithms are almost at the same scale because all algorithms incur linear memory usage



dataset Figure 14: Memory usage on undirected graphs.

w.r.t. the graph size (i.e., O(m)). For approximation algorithms, their theoretical space costs are also the same.

6.3 Evaluation of DDS algorithms

1. Overall performance. Similar to the UDS part, we first evaluate the overall performance of various DDS algorithms. To be specific, (1) we compare the efficiency and accuracy of all approximation algorithms in Table 8. Here, we also set $\epsilon = 1$ for DFWApp and VWApp; (2) we test the performance of DFWApp and VWApp over different ϵ values, from 0.1 to 0.0001 in Figure 15; and (3) we record the running time of DFlowExact, DCExact and DFWExact in Figure 16. Based on the above results, we make the following observations: (1) WCoreApp is the most efficient one for larger graphs, but DFWApp usually yields a more accurate solution. (2) Although we have incorporated the SOTA UDS algorithm into VWApp, it is still slower than DFWApp. The main reason is that, although VWApp reduces the number of different trials of c from $O(n^2)$ to $O(\log_{1+\epsilon} n)$, this number is still too many compared to that of DFWApp. (3) For exact algorithms, DFWExact always outperforms the rest, because it significantly reduces the time cost incurred for computing maximum flow, and it can compute the minimum cut on the smaller flow network.

2. Effect of the lower bound of binary search. Recall that for DFlowExact, it utilizes binary search to find the maximum density for each c. Intuitively, a tighter binary search space can speed up the algorithm, which suggests we should use ρ as the search lower bound. However, as shown in our experiment, we find that a higher lower bound may affect the effectiveness of the divide-and-conquer strategy. DCExact sets the lower bound of the binary search as 0 for each c, instead of directly using ρ . To investigate the effect of the lower bound, we introduce a hyper-parameter y, which controls the lower bound of binary search by setting $\rho = \gamma \cdot \rho$. We evaluate DCExact by varying γ from 0 to 1 on four datasets, and report their running time and the actual visited c values in Figure 18. We can see that a higher lower bound may increase the number of *c* values examined and take more time. Although we cannot predict the best *y* in advance, smaller values of *y* usually perform better. We conjecture that this is because smaller y leads to more edges remaining, more feasible values of c for the reduced graph, and then a larger interval of c to be pruned. Hence, we follow [65] to set $\rho = 0$ for each c to keep the effectiveness of the divide and conquer strategy and improve the overall performance.

3. Evaluation of graph reduction. DFWApp [63] already adopts multi-reduction to improve its efficiency. To evaluate its usefulness, we conduct an ablation study by presenting the running time, the average numbers of vertices, and edges before and after the graph reduction for different values of ϵ , across all datasets. The [x,y]-core graph reduction method can substantially reduce the size of the original graph and speed up the overall efficiency. For lack of space, we present the detailed results in our technical report [78].

Given a lower bound $\underline{\rho}$ for the optimal density, and the required interval of c values $(c_l, \overline{c_r})$ to be examined, DFWExact and DFWApp

do not use $\left[\frac{\underline{\rho}}{2\sqrt{c_r}}, \frac{\sqrt{c_l}\rho}{2}\right]$ -core to reduce the search space. The main

reason is that for a wide interval of c, (i.e., smaller c_1 , and larger c_r), only a small fraction of vertices and edges will be removed due to the small values of x and y. To further harness the power of graph reduction, the CP-based algorithms adjust the interval length of c values to enhance the effectiveness of [x, y]-core reduction. Specifically, a larger c_1 and a smaller c_r lead to higher x and y values, which significantly reduce the size of the graph. To test the effectiveness of this adjustment strategy, we evaluate the CP-based algorithms by employing and not employing this strategy, which are denoted by DFW and DFWReN, across different ϵ values on the two largest datasets in Table 9. Specifically, we record the number of c values to be checked (marked as "#c"), the average number of edges remaining after applying the [x, y]-core reduction (marked as "#edges"), the average number of iterations processed by the Frank-Wolfe algorithm (marked as "#iterations"), the product of these three items (marked as "product"), and the running time. We can observe that: (1) the running time is generally proportional to the value of the "product". (2) While this strategy may increase the number of *c* values that need to be examined, it significantly reduces the scale of the graph and the value of the "product", indicating a better performance. We find that this adjustment strategy is not useful for DCExact, since for each c, it sets ρ = 0 which means that it needs to calculate the maximum flow on a bigger graph.

- **4. Evaluation of weight update strategies.** We also compare the simultaneous and sequential update strategies on directed graphs in Figure 17. Unlike the UDS problem, we discover that the former strategy is not always more effective than the latter one. The main reason is that these two strategies have different effects on the divide-and-conquer strategy, with the performance gap largely due to the different number of *c* values that need to be examined.
- **5. Memory usage.** In this experiment, we evaluate the memory usage of all exact algorithms. As shown in Figure 19, we can observe that DCExact uses less memory than DFWExact due to the effectiveness of its graph reduction. For those approximation algorithms, they are around the same scale (i.e., O(m)), so we omit the details.

7 LESSONS AND OPPORTUNITIES

We summarize the lessons (L) for practitioners and propose practical research opportunities (O) based on our observations.

Lessons:

 $\underline{\mathbf{L1}}$. In Figure 20, we depict a roadmap of the recommended DSD algorithms, highlighting which algorithms are best suited for different scenarios.

<u>L2.</u> Graph reduction is very useful for all algorithms and using graph reduction multiple times is better than using it only once.

L3. Furthermore, no single CP-based UDS algorithm dominates other algorithms in all tests, as their performance highly depends on the structure of the graph.

<u>L4.</u> For all iteration-based algorithms, the number of iterations required in practice is much less than the theoretical estimate.

<u>L5.</u> It is not a good idea to transfer the DDS problem to the UDS problem, since it cannot utilize the divide and conquer strategy to reduce the number of *c* values that need to be examined.

<u>L6.</u> For the DDS problem, it is essential to reduce the number of c values examined. Besides, when attempting to speed up the efficiency for a specific c, it is important to evaluate if it affects the number of c values that need to be examined subsequently.

Table 8: Comparison of 2-approx. DDS algorithms (Red denotes the most efficient algorithm, and Blue denotes the best accuracy).

Algorithm	OF		AD		AM		BA		WE		SK	
	time	ratio	time	ratio	time	ratio	time	ratio	time	ratio	time	ratio
DGreedy	_	_	_	_	_	_	_	_	_	_	_	
DFWApp	87.8 ms	1.001	175.3 ms	1.009	822.2 ms	1.004	26.8 s	1	645.9 s	1	1473.6 s	1.000
FWVW	139.0 ms	1.135	391.1 ms	1.122	6623.3 s	1.004	557.1 s	1.226	7855.1 s	1	19585.2 s	1.010
MWUVW	204.7 ms	1.024	600.0 ms	1.122	_	_	480.0 s	1.226	7739.7 s	1	18417.0 s	1.000
FISTAVW	251.7 ms	1.135	480.1 ms	1.122	846.6 s	1.004	408.1 s	1.000	7349.1 s	1	22076.3 s	1.000
XYCoreApp	27.9 ms	1.422	17.2 ms	1.130	751.7 ms	1.004	4.6 s	1.000	131.0 s	1.007	1423.9 s	1.000
WCoreApp	333.7 ms	1.396	6.0 ms	1.130	253.9 ms	1.004	1.4 s	1.000	12.6 s	1.007	100.3 s	1.000

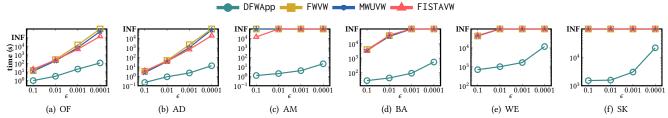


Figure 15: Efficiency results of (1 + ϵ)-approximation algorithms on directed graphs.

Table 9: Effectiveness of graph reduction technique on directed graphs.

	Dataset	$\epsilon = 0.01$		$\epsilon = 0$	0.001	$\epsilon = 0$.0001	$\epsilon = 0$		
Dataset		DFW	DFWReN	DFW	DFWReN	DFW	DFWReN	DFW	DFWReN	
	# of Ratios	25	21	27	32	54	35	20	19	
	# of Edges	9,458,420	41,014,857	9,597,565	30,606,451	9,221,802	28,859,994	10,767,492	44,912,585	
WE	# of Iterations	174	152	486	603	2,156	2,580	100	105	
	Product	3.78×10^{10}	1.31×10^{11}	1.35×10^{11}	5.91×10^{11}	6.36×10^{11}	2.61×10^{12}	2.15×10^{10}	8.98×10^{10}	
	Time (s)	1,033.3	3,723.7	2,266.8	13,658.4	7,869.2	23,916.9	643.0	2,865.3	
	#Ratios	17	15	24	18	28	_	21	15	
	# of Edges	44,552,779	375,979,205	44,837,242	313,662,857	44,362,182	_	45,844,129	365,743,496	
SK	#Iterations	124	120	258	344	1,914	_	124	147	
	Product	9.36×10^{10}	6.77×10^{11}	2.78×10^{11}	1.94×10^{12}	2.38×10^{12}	_	1.19×10^{11}	8.05×10^{11}	
	Time (s)	1,419.7	15,856.6	3,425.8	48,106.6	24,922.1	_	2,425.7	15,909.8	

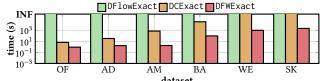


Figure 16: Efficiency of exact algorithms on DDS.

Opportunities:

<u>O1.</u> For all UDS algorithms, simultaneous weight updates tend to result in better efficiency than sequential weight updates. Providing a solid theoretical explanation of this phenomenon is an important open problem.

O2. All existing DSD algorithms (both UDS and DDS) assume the setting of a single machine. What if the graph is too large to fit on a single machine? For example, the Facebook social network contains 1.32 billion nodes and 140 billion edges (http://newsroom.fb. com/companyinfo). Can we design I/O-efficient, distributed, or sublinear time algorithms for the DSD problem? Given the advantages of GPUs in executing multiple tasks concurrently due to their highbandwidth parallel processing capabilities, designing GPU-friendly DSD algorithms [64] is another interesting opportunity.

<u>O3.</u> In many domains, the network is private, and returning the DSD can reveal information about the network. Some existing UDS

algorithms [25, 27] have considered the framework of local differential privacy. Designing a similar extension for DDS is important.

O4. Heterogeneous graphs are prevalent in various domains such as knowledge graphs, bibliographic networks, and biological networks. A promising future research direction is to study DSD for heterogeneous graphs. Besides, studying DSD for attributed graphs is also an interesting future research problem. More lessons and opportunities are reported in our technical report [78].

O5. None of the well-known graph systems, such as Neo4j [72], Nebula [71], and TigerGraph [86], has incorporated DSD algorithms using only their APIs. Therefore, enabling these popular graph systems to support DSD algorithms presents an exciting opportunity.

8 RELATED WORKS

In this section, we review the related works, including the variants of the UDS and DDS solutions and dense subgraph discovery.

• Other variants of DSD. Many variants of DSD have been studied [5, 22, 32, 62, 77, 82, 92]. The clique densest subgraph (CDS) problem is proposed to better detect "near-clique" subgraphs [29, 31, 41, 58, 68, 87, 98]. Note that when k = 2, it reduces to the UDS problem. The network flow-based [31, 68, 87] and convex programming-based [41, 85] algorithms are developed to solve this

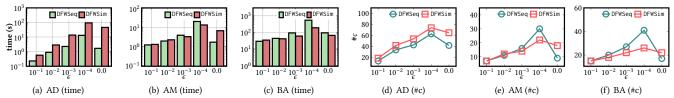


Figure 17: The effect of update strategies on directed graphs.

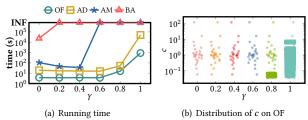


Figure 18: The effect of setting ρ for DCExact.

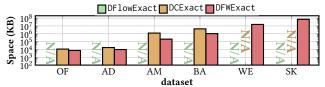


Figure 19: Memory usage on directed graphs.

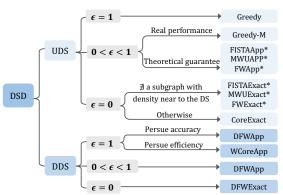


Figure 20: The taxonomy tree of DSD algorithms.

problem. The size-constrained DSD problems are studied in undirected and directed graphs [6, 11–13, 15, 39, 49, 74]. Another version of DSD called optimal quasi-clique [88, 89] extracts a subgraph that is more compact, with a smaller diameter than the DSD. To identify locally dense regions, Qin et al. [77] and Ma et al. [62] studied the top-k locally DS problem. Veldt et al. [90] studied the P-mean DSD problem and proposed a generalized peeling algorithm. To personalize results, the anchored DSD problem [22] aims to maximize R-subgraph density of the subgraphs containing an anchored node set. Besides, the DSD problems in bipartite, multilayer, and uncertain graphs were studied [4, 34, 35, 42, 46, 68, 70]. Recently, the fair DS problem and diverse DS problems [3, 69] have been explored to achieve equitable outcomes and overcome algorithmic bias.

• **Dense subgraph discovery.** Another group of works highly related to DSD are about dense subgraph discovery. Many cohesive subgraph models like k-core [9, 83], k-truss [21, 81, 95], k-ECC [43, 94], k-clique [23], quasi-clique [1], and k-plex [8, 99] have

been studied. k-core is one of the most widely used dense subgraphs, in which all vertices have a degree of at least k. k-truss is a dense subgraph based on the constraint of the number of triangles, in which each edge is contained by at least k-2 triangles. k-ECC is a subgraph based on edge connectivity, and the edge connectivity of two vertices u and v is the minimum value of the number of edges that need to be removed to make u and v disconnected. k-clique is a complete graph of size k, while k-plex allows for up to k missing connections in a clique, reflecting a quasi-clique model. These models are often used in community search as they are cornerstone of the community [28]. Besides, these models are extended to bipartite graphs, such as (α, β) -core [26, 54], bitruss [91, 100], biclique [60], and biplex [57, 93]. The directed dense subgraph models such as D-core [30, 36, 53, 56] and D-truss [55] have also been studied. Nevertheless, these works are different from DSD since they do not use the density definition as a key metric.

To the best of our knowledge, our work is the first study that provides a unified framework for all existing DSD algorithms and compares existing solutions comprehensively via empirical results.

9 CONCLUSIONS

In this paper, we provide an in-depth experimental evaluation and comparison of existing Densest Subgraph Discovery (DSD) algorithms. We first provide a unified framework, which can cover all the existing DSD algorithms, including exact and approximation algorithms, using an abstraction of a few key operations. We then thoroughly analyze and compare different DSD algorithms under our framework for both undirected and directed graphs, respectively. We further systematically evaluate these algorithms from different angles using various datasets, and also develop variations by combining existing techniques, which often outperform state-of-the-art methods. From extensive empirical results and analysis, we have identified several important findings and analyzed the critical components that affect the performance. In addition, we have summarized the lessons learned and proposed practical research opportunities that can facilitate future studies.

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