THE CDS ALGORITHM FOR ALL k VALUES

Algorithm 4: CCAS-A

```
input: A graph G, two integers T, and \omega.
     output: Approximate CDS for all k values.
 1 S_{\omega} \leftarrow an approximate \omega-clique CDS of G;
 _{2} \gamma \leftarrow \max\{x \mid {x \choose \omega} \leq \rho_{\omega}(S_{\omega}) \cdot |S_{\omega}|\};
 4 foreach k \leftarrow 3, 4, \cdots, \omega - 1 do
           SCT \leftarrow build\_SCT(G_k); // build the SCT for G_k
            S_k(G) \leftarrow run lines 3-19 in Algorithm 3;
           \mu_{k+1} \leftarrow \max\{\mu \mid \binom{\mu}{k} \leq \frac{\binom{\gamma}{k+1}}{|S_{\omega}|}\};
V_i \leftarrow \text{the first } i\text{-th vertices being removed in the first iteration;}
           \lambda \leftarrow \text{the largest } i \text{ such that } \max_{v \in V_i} l^1(v) \leq {\mu_{k+1} \choose k};
           G_{k+1} \leftarrow remove all vertices in V_{\lambda} from G_k;
11 return S_3(G), S_4(G), \cdots, S_{\omega}(G)
```

Algorithm 4 illustrates the CCAS-A algorithm for finding CDS over all k values. It first computes an approximate CDS of ω -cliques, denoted as S_{ω} , where ω represents the maximum clique size (line 1). Next, based on Lemma 6.1, it calculates the γ value, which is used for the subsequent graph reduction process (line 2). CCAS-A process the k value from 3 to ω one by one (lines 4-10). Specifically, for each k, it invokes the CCAS to compute the approximate k-CDS (lines 5-6). Afterwards, it updates the μ value and further reduces the search space based on Theorem 6.2 (lines 7-10). Finally, it returns the approximate CDS for all k values (line 11).

В ADDITIONAL PROOFS OF LEMMAS AND **THEOREMS**

LEMMA 4.2. Given a graph G = (V, E), if $|\Psi_k(G)| \ge {\mu \choose k}$ where μ is an arbitrary integer no less than k, then $|\Psi_{k-1}(G)| \ge {\mu \choose k-1}$.

PROOF. We prove Lemma 4.2 holds by proving its contrapositive, that is, given a graph G=(V,E), if $|\Psi_{k-1}(G)|<\binom{\mu}{k-1}$, then $|\Psi_k(G)|<\binom{\mu}{k}$. Since $|\Psi_{k-1}(G)|<\binom{\mu}{k-1}$, the maximum clique size in G is less than μ . Suppose $\mathcal{R}(G)$ is the set of all maximal cliques in G. For each k-clique in G, it must be included in at least one maximal clique of G, so we have:

$$|\Psi_k(G)| = \left| \bigcup_{R \in \mathcal{R}(G)} \Psi_k(R) \right| \tag{34}$$

That is, $\forall \Phi \subseteq \mathcal{R}(G)$, we denote $g(\Phi) = \left| \bigcap_{R \in \Phi} \Psi_k(R) \right|$, then:

$$g(\Phi) = \left| \bigcap_{R \in \Phi} \Psi_k(R) \right| \tag{35}$$

$$= \left| \Psi_k(\bigcap_{R \in \Phi} R) \right|$$

$$= \left(\bigcap_{R \in \Phi} R \right)$$
(36)
$$= \left(\bigcap_{R \in \Phi} R \right)$$
(37)

$$= \begin{pmatrix} | \bigcap_{R \in \Phi} R| \\ |_L \end{pmatrix} \tag{37}$$

This is because a k-clique is contained in all maximal cliques in S, if and only if it is included in the subgraph formed by the intersection of those maximal cliques. Besides, the intersection of some cliques must either be a clique or an empty set, and the number of k-cliques in this subgraph is $\binom{|\bigcap R|}{R \in S}$.

By applying the inclusion-exclusion principle on equation (34),

$$|\Psi_k(G)| = \left| \bigcup_{R \in \mathcal{R}(G)} \Psi_k(R) \right|$$

$$= \sum_{S \subseteq \mathcal{R}(G)} (-1)^{|S|} g(S)$$
(38)

$$=\sum_{S \subseteq \mathcal{R}(G)} (-1)^{|S|} g(S) \tag{39}$$

$$= \sum_{S \subseteq \mathcal{R}(G)} (-1)^{|S|} {\binom{|\bigcap R|}{R \in S}}$$
 (40)

Similarly.

$$|\Psi_{k-1}(G)| = \sum_{S \subseteq \mathcal{R}(G)} (-1)^{|S|} {|\bigcap_{\substack{R \in S \\ k-1}}}$$
(41)

Therefore, there exist a vector $\beta = \{\beta_{k-1}, \beta_k, \dots, \beta_{\mu-1}\}$, such that,

$$|\Psi_{k-1}(G)| = \sum_{i=k-1}^{\mu-1} \beta_i \binom{i}{k-1}$$
 (42)

$$|\Psi_k(G)| = \sum_{i=k-1}^{\mu-1} \beta_i \binom{i}{k}$$
 (43)

Since, $\forall i \in [k-1, \mu-1]$, we have $\frac{\binom{k}{i}}{\binom{i}{i}} = \frac{i-k+1}{k} < \frac{\mu-k+1}{k}$, that is:

$$\binom{i}{k} < \frac{\mu - k + 1}{k} \cdot \binom{i}{k - 1} \tag{44}$$

Therefore, we can conclude that

$$\frac{|\Psi_k(G)|}{|\Psi_{k-1}(G)|} = \frac{\sum_{i=k-1}^{\mu-1} \beta_i \binom{i}{k}}{\sum_{i=k-1}^{\mu-1} \beta_i \binom{i}{k-1}}$$
(45)

$$\sum_{i=k-1}^{\mu-1} \beta_{i} {i \choose k-1} \\
\leq \frac{\sum_{i=k-1}^{\mu-1} \beta_{i} \frac{\mu-k+1}{k} {i \choose k-1}}{\sum_{i=k-1}^{\mu-1} \beta_{i} {i \choose k-1}} \\
\leq \frac{\mu-k+1}{k}$$
(46)
$$\frac{\mu-k+1}{k} \\
\frac{(G)|}{(G)|} \leq \frac{\mu-k+1}{k}, \text{ therefore:}$$

$$<\frac{\mu-k+1}{k}\tag{47}$$

Hence, we have $\frac{|\Psi_k(G)|}{|\Psi_{k-1}(G)|} < \frac{\mu - k + 1}{k}$, therefore:

$$|\Psi_{k}(G)| < \frac{\mu - k + 1}{k} |\Psi_{k-1}(G)|$$

$$< \frac{\mu - k + 1}{k} {\mu \choose k-1} = {\mu \choose k}$$
(48)

$$<\frac{\mu-k+1}{k}\binom{\mu}{k-1}=\binom{\mu}{k}$$
 (49)

Lemma 4.3. Given a graph $G = (V, E), \forall v \in V, \text{ if } |\Psi_k(v, G)| \ge \rho$, then for any $r \in \mathbb{N}$, with $3 \le r \le k$, we have $|\Psi_r(v,G)| \ge {\mu \choose r-1}$, where μ denotes the maximum integer such that $\binom{\mu}{k-1} \leq \rho$.

PROOF. We use mathematical induction to prove this lemma. (1) we first prove it holds when r = k:

$$\Psi_r(v,G) = \Psi_k(v,G) \ge \rho \ge {\mu \choose k-1} \ge {\mu \choose r-1} \tag{50}$$

(2) Next, we show it holds for arbitrary $r \le k$, and we suppose that the lemma holds for r = s, so $|\Psi_s(v,G)| \ge {\mu \choose s-1}$. Let $\mathcal{Y}_s =$ $\{u \mid u \in V \land \exists C \in \Psi_s(v, G), u \in C\}$, Let $G[\mathcal{Y}_s]$ denotes the induced subgraph of \mathcal{Y}_s . Since $|\Psi_s(v)| \ge \binom{\mu}{s-1}$, we have $|\Psi_{s-1}(G[\mathcal{Y}_s])| \ge$ $\binom{\mu}{s-1}$. By Lemma 4.2, we have $|\Psi_{s-2}(G[\mathcal{Y}_s])| \ge \binom{\mu}{s-2}$. On the other hand, since all vertices in \mathcal{Y}_s are connected with

v, so we have $|\Psi_{s-1}(v,G)| = |\Psi_{s-2}(G[\mathcal{Y}_s])| \ge {\mu \choose s-2}$. Hence, the

Lemma 5.2. Given a graph G with n vertices and degeneracy of δ , CCAS costs $O(n \cdot 3^{\delta/3})$ space, and $O(n \cdot 3^{\delta/3} \cdot \delta^2)$ time for each iteration.

PROOF. Consider iteratively removing all vertices in the graph. For each root-to-leaf path Γ , we need to process it each time a node it contains is removed. Therefore, each path is processed for $O(\delta)$ node, with each cost $O(\delta)$ time. Since there are $O(n \cdot 3^{\delta/3})$ nodes in the SCT. Therefore the time complexity for CCAS for one iteration is $O(n \cdot 3^{\delta/3})$.

Lemma 5.3 Given any $t \ge 1$, denote $\bar{\alpha}$ by the vector that has a minimum inner product with the gradient of $Q_G(\alpha^{(t)})$

$$\bar{\alpha} = \begin{bmatrix} \bar{\alpha}^{C_1}, & \bar{\alpha}^{C_2}, & \cdots &, \bar{\alpha}^{C_{|\Psi_k(G)|}} \end{bmatrix} = \arg \min_{\beta \in D(G, k)} \langle \beta, \nabla Q_G(\alpha^{(t)}) \rangle.$$

We have $\widehat{\alpha}$ is an approximate linear minimizer, i.e.,

$$\langle \widehat{\alpha}, \nabla Q_G(\alpha^{(t)}) \rangle \le \langle \overline{\alpha}, \nabla Q_G(\alpha^{(t)}) \rangle + \frac{1}{2} \gamma_t \theta \xi(Q_G),$$

where $\gamma_t = \frac{1}{t+1}$, $\theta = 2$,and $\xi(Q_G) = 2\Delta |\Psi_k(G)|$.

PROOF. We first use $\nabla_C Q_G(\alpha)$ to denote the projection of $\nabla Q_G(\alpha)$ onto \mathbb{R}^C . By a straightforward calculation from proof of Lemma 4.5 [23], the (C, u)-coordinate of $\nabla Q_G(\alpha)$ is

$$2 \cdot \frac{l(u)}{t} = 2 \cdot \sum_{\bar{C} \in \Psi_k(G): u \in \bar{C}} \alpha_u^{\bar{C}}$$
 (51)

It has been noted in the proof of Lemma 4.5 in [23] that one can consider each k-clique $C \in \Psi_k(G)$ independently and Therefore, for $t \ge 1$, we have:

$$<\widehat{\alpha}, \nabla Q_G(\alpha^{(t)})> -<\bar{\alpha}, \nabla Q_G(\alpha^{(t)})>$$
 (52)

$$= \sum_{i=1}^{|\Psi_k(G)|} \langle \widehat{\alpha}^i, \nabla_{C^i} Q_G(\alpha^{(t)}) \rangle - \langle \bar{\alpha}^i, \nabla_{C^i} Q_G(\alpha^{(t)}) \rangle$$
 (53)

$$= \sum_{i=1}^{|\Psi_k(G)|} < \widehat{\alpha}^i, \nabla_{C^i} Q_G(\alpha^{(t)}) > -\frac{2}{t} \min_{v \in C_i} l^{(t)}(v)$$
 (54)

Consider the value of $\langle \widehat{\alpha}^i, \nabla_{C^i} Q_G(\alpha^{(t)}) \rangle$, it should be equal to $\frac{2l^{(t)}(v)}{t}$, where v is the first vertex removed from C_i in the t-th iteration. Therefore, we have $l^{(t)}(v) \leq \min_{v' \in C_i} l^{(t)}(v') + |\Psi_k(v', G)| \leq$ $\min_{v' \in C_i} l^{(t)}(v') + \Delta.$

$$<\widehat{\alpha}, \nabla Q_G(\frac{\alpha^{(t)}}{t})> -<\bar{\alpha}, \nabla Q_G(\frac{\alpha^{(t)}}{t})>$$
 (55)

$$= \sum_{i=1}^{|\Psi_k(G)|} < \widehat{\alpha}^i, \nabla_{C^i} Q_G(\alpha^{(t)}) > -\frac{2}{t} \min_{v \in C_i} l^{(t)}(v)$$
 (56)

$$\leq \frac{1}{t} \sum_{i=1}^{|\Psi_k(G)|} \min_{v \in C_i} l^{(t)}(v) + \Delta - \min_{v \in C_i} l^{(t)}(v)$$
 (57)

$$= \frac{2}{t} |\Psi_k(G)| \Delta \le \frac{1}{2} \gamma_t \theta \xi(Q_G), \tag{58}$$

Theorem 5.4. CCAS improves the overall running time at least by $(\rho_k^*(G)\sqrt{k})$ over KCCA and by $(\frac{(\frac{\delta}{2})^{k-2}}{\mathcal{F}})$ over SuperGreedy++.

PROOF. To obtain a $(1-\epsilon)$ -approximation ratio solution, theoretically, KCCA, SuperGreedy++, and CCAS take $O\left(\frac{1}{\xi^2} \cdot \sqrt{k}\Delta |\Psi_k(G)| \cdot \xi \cdot \delta^2 \log \delta\right)$ $O\left(\frac{\Delta \log |\Psi_k(G)|}{\rho_k^*(G) \cdot \epsilon^2} \cdot km \cdot (\frac{\delta}{2})^{k-1}\right)$, and $O\left(\frac{\Delta \log |\Psi_k(G)|}{\rho_k^*(G) \cdot \epsilon^2} \cdot \xi \cdot \delta^3\right)$ time, respectively. Compared to KCCA, our method archives an improverespectively. Compared to KCCA, our method archives an improvement of $O(\frac{\sqrt{k}\Psi_k(G)\rho_k^*(G)\log\delta}{\log\Psi_k(G)\delta})$, which can be simplified to $O(\sqrt{k}\rho_k^*(G))$. This is because, $\frac{\Psi_k(G)\log\delta}{\log\Psi_k(G)\delta} > 1$, as shown by the function $f(x) = \frac{x}{\log x}$. Since $f'(x) = \frac{\log x - 1}{(\log x)^2}$ is positive for x > 2, f(x) is an increasing function, and thus $\frac{\Psi_k(G)}{\log\Psi_k(G)} > \frac{\delta}{\log\delta}$. Besides, compared to SuperGreedy++, our method achieves an improvement of $O(\frac{km\cdot(\frac{\delta}{2})^{k-1}}{\xi\delta^3})$, which simplifies to $O(\frac{k\cdot(\frac{\delta}{2})^{k-1}}{\xi\delta})$,

since $\delta^2 < m$. This equation can be further simplified to $O(\frac{(\frac{\delta}{2})^{k-2}}{\xi})$.

Lемма 6.1. Given a graph G, and an approximate k-clique CDS, $S_k(G)$, for any k' < k, we have $\rho_{k'}^*(G) \ge \frac{\binom{\gamma}{k'}}{|S_k(G)|}$, where γ is the maximum integer such that $\binom{\gamma}{k} \le \rho_k(S_k(G)) \cdot |S_k(G)|$.

PROOF. The number of k-cliques contained in approximate kclique CDS $S_k(G)$ should be $|\Psi_k(S_k(G))| = \rho_k(S_k(G)) \cdot |S_k(G)|$. Then, we compute the largest integer γ such that $|\Psi_k(S_k(G))| \ge$ $\binom{\gamma}{k}$. By Lemma 4.2 and mathematical induction, we can conclude

that
$$|\Psi_{k'}(\mathcal{S}_k(G))| \ge {\mu \choose k'}$$
 and $\rho_{k'}^*(G) \ge \rho_{k'}(\mathcal{S}_k(G)) \ge \frac{{\gamma \choose k'}}{|\mathcal{S}_k(G)|}$.

LEMMA 6.2 Given a graph G, an approximate k-clique CDS, $S_k(G)$, for any integers $3 \le x < k$, we use $\rho_x(G)$ and $\rho_{x+1}(G)$ to denote the lower bound of optimal densities obtained by the lemma 6.1, then we have $\mu_x \le \mu_{x+1}$, where μ_x and μ_{x+1} are derived by Lemma 4.3, using $\rho_x(G)$ and $\rho_{x+1}(G)$.

PROOF. Recall that μ_x is the maximum integer such that:

$$\binom{\mu_x}{x-1} \le \frac{\binom{\gamma}{x}}{|\mathcal{S}_{\omega}(G)|},$$
 (59)

where γ is the maximum integer s.t. $\binom{\gamma}{\omega} \leq \rho_{\omega}(\mathcal{S}_{\omega}(G)) \cdot |\mathcal{S}_{\omega}(G)|$. Based on the formulation (59) we have:

$$\frac{\mu_{x}!}{(x-1)!(\mu_{x}-x+1)!} \leq \frac{\gamma!}{x!(\gamma-x)!|S_{\omega}(G)|}$$

$$\frac{\mu_{x}!}{(\mu_{x}-x+1)!} \leq \frac{\gamma!}{x(\gamma-x)!|S_{\omega}(G)|}$$
(60)

$$\frac{\mu_{x}!}{(\mu_{x}-x+1)!} \le \frac{\gamma!}{x(\gamma-x)!|\mathcal{S}_{\omega}(G)|} \tag{61}$$

Similarly, μ_{x+1} is the maximum integer satisfy

$$\frac{\mu_{x+1}!}{(\mu_{x+1}-x)!} \le \frac{\gamma!}{(x+1)(\gamma-x-1)!|\mathcal{S}_{\omega}(G)|}$$
 (62)

Therefore, to show $\mu_x \leq \mu_{x+1}$, we only need to prove the following formulation:

$$\frac{\mu_{x}!}{(\mu_{x} - x)!} \le \frac{\gamma!}{(x+1)(\gamma - x - 1)! |S_{\omega}(G)|}$$
(63)

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Table 8: Additional two graphs.

Dataset	Category		E	δ
web-Google (WG)	Web	916,428	4,322,051	44
Wikipedia (WP)	Hyperlink	3,033,374	43,845,958	175

Here, we can transfer the formulation (45) to (47) by multiplying (μ_x-x+1) and $\frac{x(\gamma-x)}{x+1}$ for the left and right hand side of it, respectively. That is to say , if we can prove:

$$\mu_{x} - x + 1 \le \frac{x(\gamma - x)}{x + 1}$$

$$\mu_{x}(x + 1) + 1 - x^{2} \le x\gamma - x^{2}$$
(64)
(65)

$$\mu_X(x+1) + 1 - x^2 \le x\gamma - x^2 \tag{65}$$

$$\mu_X \le \frac{x\gamma - 1}{x + 1} \tag{66}$$

$$\mu_x \le \frac{x\gamma - 1}{x + 1}$$

$$\mu_x \le \gamma - \frac{\gamma + 1}{x + 1}$$

$$(66)$$

then the formulation (63) can also be proved

Since $\rho_{\omega}(S_{\omega}(G)) \cdot |S_{\omega}(G)| \ge 1$, we have $\gamma \ge \omega > x$. Therefore, we can rewrite y in terms of $a \cdot (x+1) + b$, where $a \ge 1$ and $0 \le b \le k$.

$$\mu_{x} \le a(x+1) + b - \frac{ax + a + b + 1}{x+1}$$

$$\mu_{x} \le ax + b - \frac{b+1}{x+1}$$
(68)

$$\mu_X \le ax + b - \frac{b+1}{x+1} \tag{69}$$

Since μ_x is an integer, we only need to prove:

$$\mu_X \le ax + b - 1 \tag{70}$$

We prove this by contradiction, where we aim to show that if $\mu_x > ax + b - 1$, it must violate formulation (61). Since the left-hand side of (61) is monotonically non-decreasing with μ_x , we only need to prove the case for $\mu_x = ax + b$, i.e.,:

$$\frac{(ax+b)!}{(ax+b-x+1)!} > \frac{(a\cdot (x+1)+b)!}{x(a\cdot (x+1)+b-x)!|S_{\omega}(G)|}$$
(71)

Recall that $\rho_{\omega}(S_{\omega}(G)) \cdot |S_{\omega}(G)| = |\Psi_{\omega}(S_{\omega}(G))| \geq {\gamma \choose \omega}$, so we have $|S_{\omega}(G)| \ge \gamma = ax + b$. Hence a stricter inequality can be built:

$$\frac{(ax+b)!}{(ax+b-x+1)!} > \frac{(a\cdot(x+1)+b-1)!}{x(a\cdot(x+1)+b-x)!}$$
(72)

$$x \cdot \frac{(ax+b)!}{(ax+b-x+1)!} > \frac{(a \cdot (x+1)+b-1)!}{(a \cdot (x+1)+b-x)!}$$
 (73)

$$x \cdot \frac{(a \cdot (x+1) + b - x)!}{(ax+b-x+1)!} > \frac{(a \cdot (x+1) + b - 1)!}{(ax+b)!}$$
 (74)

$$S_{\omega}(G)| \ge \gamma = ax + b. \text{ Hence a stricter inequality can be built:}$$

$$\frac{(ax+b)!}{(ax+b-x+1)!} > \frac{(a \cdot (x+1)+b-1)!}{x(a \cdot (x+1)+b-x)!}$$

$$x \cdot \frac{(ax+b)!}{(ax+b-x+1)!} > \frac{(a \cdot (x+1)+b-1)!}{(a \cdot (x+1)+b-x)!}$$

$$x \cdot \frac{(a \cdot (x+1)+b-x)!}{(ax+b-x+1)!} > \frac{(a \cdot (x+1)+b-1)!}{(ax+b)!}$$

$$x \cdot \prod_{i=ax+b-x+a}^{ax+b-x+a} i > \prod_{i=ax+b+1}^{ax+b+a-1} i$$

$$x \cdot \prod_{i=ax+b-x+a}^{ax+b-x+a} i > \prod_{i=ax+b-x+a}^{ax+b-x+a} (i+x-1)$$

$$(75)$$

$$x \cdot \prod_{i=ax+b-x+2}^{ax+b-x+a} i > \prod_{i=ax+b-x+2}^{ax+b-x+a} (i+x-1)$$
 (76)

$$x > \prod_{i=ax+b-x+a}^{i=ax+b-x+a} \frac{(i+x-1)}{i}$$
 (77)

If a = 1, the right-hand side of formulation (77) is 1, which is clearly smaller than x. For the case of $a \ge 2$, we complete our proof by bounding on the right-hand side of formulation (77).

$$\lim_{i=ax+b-x+2} \sin \theta = \lim_{i=ax+b-x+2} \frac{(i+x-1)}{i} \le \lim_{i=ax+b-x+2} \frac{ax+b+1}{ax+b-x+2}$$
 (78)
$$= \left(\frac{ax+b+1}{ax+b-x+2}\right)^{a-1}$$
 (80)
$$= \left(1 + \frac{1}{a-1}\right)^{a-1}$$
 (81)
$$\le \lim_{a \to \infty} \left(1 + \frac{1}{a-1}\right)^{a-1} = e$$
 (82)
$$\lim_{a \to \infty} \left(1 + \frac{1}{a-1}\right)^{a-1} = e$$
 (82)

$$= \left(\frac{ax+b+1}{ax+b-x+2}\right)^{a-1} \tag{79}$$

$$\leq \left(\frac{ax}{(a-1)\cdot x}\right)^{a-1} \tag{80}$$

$$= \left(1 + \frac{1}{a - 1}\right)^{a - 1} \tag{81}$$

$$\leq \lim_{a \to \infty} (1 + \frac{1}{a-1})^{a-1} = e$$
 (82)

Hence, we finish our proof since $x \ge 3 > e$.

ADDITIONAL EXPERIMENTS

Additional Datasets.

We present the statistics of the additional two graphs on Table 8 from the different domains. They are available on the Stanford Network Analysis Platform, and Laboratory of Web Algorithmics.

C.2 The k-clique densities

In Figure 12, we report the actual k-clique-densities of the approximation solutions returned by all the CDS algorithms on BG, EB, WT, and SD datasets. We observe that CCAS and SuperGreedy++ always perform better than the Frank-Wolfe-based algorithms, and typically, CCAS and SuperGreedy++ only require one iteration to achieve a solution with the same density that other algorithms obtain after ten iterations. In addition, both KClist++ and PSCTL yield comparable performance, and they slightly outperform KCCA, since they can enable a more balanced weight distribution among vertices, making them converge faster.

C.3 Effectiveness of the graph reduction

In this experiment, we evaluate the effectiveness of our graph reduction technique, by reporting the number of vertices in the (k-1)core, CDS, and CCAS for k = 7. We present results only for k = 7, as other *k* values yield similar results. In addition, more results for different *k* values can be found in our supplementary material. As shown in Table 9, we observe that: (1) the number of vertices in the CDS is remarkably smaller, usually no more than a few thousand; (2) across nearly all datasets, the number of vertices after HCGR closely matches the number of vertices in the CDS; (3) the number of vertices in the 6-core remains very high, being up to 100,000× larger than the number of vertices remaining after HCGR.

C.4 Case studies

Here, we would conduct the case studies on two real datasets, namely S-DBLP and Yeast. S-DBLP (|V|=478, |E|=1,086) is a subgraph of the DBLP dataset, which is about the co-authorship network of authors who published at least two DB/DM papers between 2013 and 2015. The triangle densest subgraph is depicted in Figure 13(a). In a triangle (3-clique), every pair of vertices is connected, so the CDS tends to be a near-clique [28, 33, 71]. The researchers involved in this CDS possess a close collaboration relationship: any two researchers have published papers together. In addition, Figure 13(b) shows the 4-clique densest subgraph in the yeast PPI network (|V|=1,116, |E|=2,148). This CDS shows a protein-subnetwork with the "subcellular localization", "cellular transport", and "protein synthesis" functional classes, which are all 4-cliques. Hence, tasks such as analyzing the conservation and evolution of cellular components [28] can then be performed on this subnetwork.

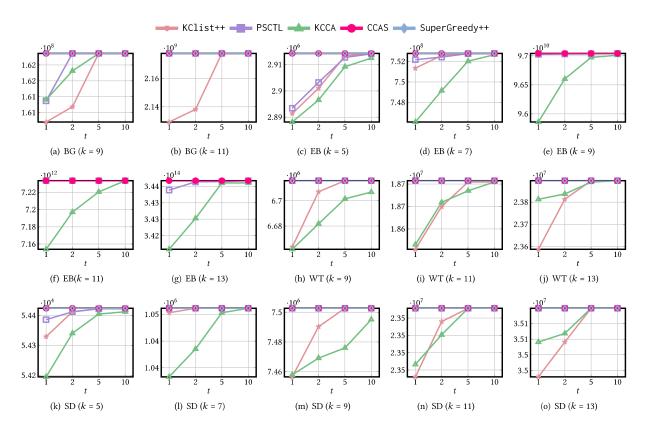


Figure 12: The k-clique densities of CDS obtained by SuperGreedy++, KClist++, KCCA, PSCTL, and CCAS.

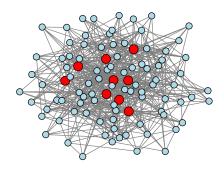


Figure 14: A case study to illustrate the HCGR.

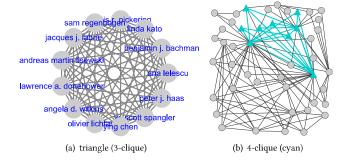


Figure 13: The densest subgraphs found in small DBLP and Yeast networks.

Besides, we also present a case study to illustrate our HCGR algorithm, as shown in Figure 14. The k-core-based method does not remove any vertices, whereas HCGR eliminates the blue vertices while retaining the red ones. This demonstrates that HCGR effectively removes nearly all vertices that are not part of the CDS.

C.5 Effect of ϵ

We evaluate the effect of ϵ using five datasets from different domains, where each domain has a dataset, and the values of ϵ are set to 1, 0.1, 0.05, and 0.01, respectively. The experimental results are reported in Table 10, which clearly shows that CCAS outperforms the other algorithms on all datasets. For around half of the datasets, CCAS is over three orders of magnitude faster than its competitors, those results are similar to the results for 0.1 and 0.01 cases, as shown in Table 6.

C.6 Efficiency of finding the CDS's for all k.

We compare the efficiency of our algorithm CCAS-A and others for finding the CDS's for all k values. Specifically, for each graph, we record the running time of the three algorithms PSCTL, KCCA, and CCAS-A as they process the k values from 3 to $25\% \cdot \omega$, $50\% \cdot \omega$, $75\% \cdot \omega$, and $100\% \cdot \omega$ across six datasets in Table 11. We present the results for ϵ from 1 to 0.01. The conclusions are similar to the Experiment 3 in the Section 7.2. Besides, to our best knowledge, CCAS is the first algorithm that can produce solutions of all k values

Table 9: # vertices in the (k-1)-core, CDS, and CCAS (k=7).

Datasets	# vertices in 6-core	$ V(\mathcal{D}_k(G)) $	CCAS
BG	1,306	116	268 (43.4 %)
EB	487	231	444 (52.0 %)
WT	2,619	127	551 (23.0 %)
SD	21,686	117	575 (20.3 %)
DP	68,070	114	216 (52.8 %)
HT	21,239	563	563 (100 %)
WG	367,361	116	256 (45.3 %)
WP	1,466,413	178	1,668 (10.7 %)
HW	1,069,126	2209	2209 (100 %)
ZB	3,391,200	268	325 (82.4 %)
UK	11,404,146	944	1,665 (56.7 %)
AC	16,063,729	3,250	3,250 (100 %)
IT	27,201,499	4,279	4,279 (100 %)
FS	36,821,624	141	141 (100 %)

with an approximation ratio of 0.99 for graphs with billions of edges in one hour.

D MORE DISCUSSIONS

The CDS solution has been used in many fundamental graph data management tasks, such as supporting graph visualization [82, 83],

community detection (or search) [4, 15, 28, 71, 72], identifying near-cliques [27, 46, 48, 53, 71], and path/reachability queries [17, 42]. In addition, the CDS problem also as a key task in graph data management has gained notable attention at top-tier data management conferences, including VLDB 2019 [28], 2020 [66], 2022 [27], SIG-MOD 2023 [36], and 2024 [84].

Finding CDS efficiently is very useful in many graph data mining applications. Specifically, it can help identify research communities in the DBLP network [23,59,60], detect subnetworks with a specific function in the biology network [28, 71, 72] and clusters in senators' networks on US bill voting [71], and discover compact dense subgraphs from e-commerce and social networks [28] when k is relatively small. In addition, identifying CDS with large k also has many applications. As shown in [27, 46, 48, 53, 71], a CDS becomes more akin to a large near-clique as k grows, useful for tasks like finding biologically relevant groups in protein interactions [19, 41, 71, 72], community detection [4, 15, 72], and anomaly detection [31, 67, 79], etc. In many of these applications, finding a "near-clique" is very important since a "near-clique" can be considered a clique in the forming stage or one with missing edges due to data corruption.

Table 10: Effect of ϵ and k. (Processing time (in seconds); Best performers are highlighted in bold.)

Dataset	Method		k = 11				k = 15				k = 19						
Dataset	Method	1	0.1	0.05	0.01	1	0.1	0.05	0.01	1	0.1	0.05	0.01	1	0.1	0.05	0.01
НТ	PSCTL KCCA CCAS	79,752 76,191 8.6	79,752 76,191 8.6	155,885 150,316 8.6	268,701 261,231 8.6	- 8.5	- 8.5	- 8.5	- 8.5	- 8.5	- 8.5	- 8.5	- 8.5	- - 8.4	- 8.4	- 8.4	- 8.4
WG	PSCTL KCCA CCAS	2.8 7.0 0.2	2.8 8.2 0.2	6.9 11.5 0.3	18.3 16.0 0.3	1.4 8.3 0.2	3.2 8.3 0.2	3.2 8.3 0.2	4.8 8.6 0.2	1.3 6.9 0.1	1.3 6.9 0.1	1.3 6.9 0.1	2.5 6.9 0.1	0.9 6.1 0.1	0.9 6.1 0.1	0.9 6.1 0.1	2.0 6.1 0.1
HW	PSCTL KCCA CCAS	52.3	- 52.3	_ _ 52.3	_ 52.3	- - 52.2	- 52.2	- 52.2	- 52.2	53.9	- - 53.9	- - 53.9	- - 53.9	- - 51.1	- - 51.1	- 51.1	- 51.1
ZB	PSCTL KCCA CCAS	89.4 185.5 11.5	123.5 261.5 11.5	182.1 261.5 11.5	507.6 339.2 12.1	100.4 150.2 9.3	143.9 176.9 9.3	143.9 176.9 9.3	634.7 405.3 10.0	79.4 105.7 7.0	112.6 127.4 7.0	112.6 127.4 7.2	466.0 323.2 8.4	65.1 122.6 5.4	94.9 147.9 5.4	94.9 147.9 5.5	404.9 370.6 6.7
UK	PSCTL KCCA CCAS	78.6	- - 78.6	- - 78.6	- - 78.6	_ _ 20.0	_ _ 20.0	_ _ 20.0	_ _ 20.0	- - 18.9	- - 18.9	- - 18.9	- - 18.9	- - 17.8	- - 17.8	- - 17.8	- - 17.8
AC	PSCTL KCCA CCAS	- - 5,853	- 5,853	- 5,853	- 5,853	- 6,703	- 6,703	- 6,703	- 6,703	- - 6,967	- 6,967	- 6,967	- 6,967	- 7,056	- 7,056	- 7,056	- 7,056

 $\label{thm:conds:cond$

Dataset	Method	$\epsilon=1$ $\epsilon=0.1$ $\epsilon=0.05$								$\epsilon = 0.01$							
Dataset	Method	25%	50%	75%	100%	25%	50%	75%	100%	25%	50%	75%	100%	25%	50%	75%	100%
DP	PSCTL KCCA CCAS-A	5.3 5.0 0.7	9.3 5.7 0.7	12.9 6.1 0. 7	16.0 6.3 0.7	6.2 6.4 0.7	10.2 7.1 0.7	13.8 7.5 0.7	16.9 7.7 0.7	6.4 7.3 0.7	10.4 8.0 0. 7	14.0 8.4 0. 7	17.1 9.6 0.7	9.1 8.1 0.7	13.2 8.8 0.7	16.7 9.3 0.7	19.8 9.5 0.7
НТ	PSCTL KCCA CCAS-A	- - 322.4	- 322.4	- 322.4	- 322.4	- - 322.4	- 322.4	- 322.4	- 322.4	- 322.4	- 322.4	- 322.4	- 322.4	- 322.4	- 322.4	_ _ 322.4	- 322.4
WP	PSCTL KCCA CCAS-A	3,219 4,557 28.5	3,855 6,071 30.8	4,127 6,556 32.7	4,211 6,702 33.9	4,798 5,713 39.0	5,434 7,338 44.2	5,707 7,869 46.8	5,797 8,056 48.0	7,080 6,597 48.6	7,719 8,254 55.1	7,995 8,840 60.7	8,085 9,050 62.2	21,103 13,970 243.1	21,789 15,767 255.7	22,080 16,622 268.0	22,172 16,853 274.5
ZB	PSCTL KCCA CCAS-A	2,438 3,960 66.0	2,983 4,206 103.0	3,441 4,334 136.9	3,707 4,396 161.7	3,509 4,781 66.0	4,053 5,033 103.0	4,511 5,163 136.7	4,777 5,230 161.7	3,788 4,996 72.8	4,390 5,258 114.1	4,884 5,393 151.1	5,150 5,468 176.7	12,181 9,949 106.0	12,968 10,273 166.6	13,569 10,453 217.3	13,835 10,581 247.5
UK	PSCTL KCCA CCAS-A	- - 456.6	- - 456.6	- 456.6	- 456.6	- - 456.6	- - 456.6	- - 456.6	- - 456.6	- - 456.6	- 456.6	- 456.6	- 456.6	- - 456.6	- - 456.6	- - 456.6	- - 456.6
FS	PSCTL KCCA CCAS-A	119,740 — 2,167	185,679 — 2,168	239,801 - 2,169	280,842 — 2,169	125,544 — 2,167	191,483 — 2,168	245,604 - 2,169	286,646 — 2,169	135,619 - 2,273	201,558 — 2,274	255,679 — 2,275	296,721 - 2,276	201,621 - 3,090	267,560 — 3,091	321,681 - 3,092	- 3,092