Neighborhood-based Hypergraph Core Decomposition

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ABSTRACT

We propose neighborhood-based core decomposition: a novel way of decomposing hypergraphs into hierarchical neighborhood-cohesive subhypergraphs. Alternative approaches of decomposing hypergraphs such as reduction to clique graphs or bipartite graphs inflate the problem-size and are not meaningful in certain applications, while existing degree-based hypergraph decomposition does not distinguish nodes with different neighborhood sizes. Applications and our case-studies show that the proposed decomposition is more effective than degree-based decomposition in diffusion problems such as disease intervention and in extracting provably approximate and application-wise meaningful, non-trivial densest subhypergraphs. As technical contributions, we propose three algorithms: **Peel**, its efficient variant E-Peel, and a novel local algorithm: Local-core with parallel implementation. Our most efficient sequential algorithm Local-core(OPT) decomposes hypergraph with 27M nodes and 17M hyperedges in-memory within 16 minutes by adopting various optimizations that are unique to hypergraphs. Our parallel implementation Local-core(P) further expedites computation achieving at least 2-5x speedup over Local-core(OPT).

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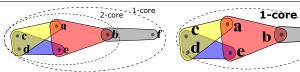
1 INTRODUCTION

Decomposition of a graph into hierarchically cohesive subgraphs is an important tool for solving many graph data management problems, e.g., community detection [49], densest subgraph discovery [15], identifying influential nodes [48], and network visualization [1, 8]. Depending on different notions of cohesiveness, there are

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decomposition of hypergraph H

Figure 1: Neighborhood-based core Figure 2: Degree-based hypergraph core decomposition of H

several decomposition approaches: core-decomposition [9], trussdecomposition [62], nucleus-decomposition [53], etc. In this work, we are interested in decomposing hypergraphs, a generalization of graphs where an edge may connect more than two entities.

Many real-world relations consist of poly-adic entities, e.g., relations between individuals in co-authorships [33], legislators in parliamentary voting [11], items in e-shopping carts [66], proteins in protein complexes, and metabolites in a metabolic process [24, 27]. Often such relations are reduced to a clique graph or a bipartite graph for convenience (§2.2). However, these reductions may not be desirable due to two reasons. First, such reductions might not be meaningful, e.g., three authors collaborating on an article does not necessarily mean that they must have collaborated pairwise and vice versa. A pair of proteins in a certain protein-complex may not necessarily interact pairwise to create a new functional protein-complex. We show in §2.2 that the core decomposition of the clique graph or bipartite graph representation of a hypergraph may not yield the same result as the core decomposition of the hypergraph. Second, reducing a hypergraph to a clique graph or a bipartite graph inflates the problem-size [36]: A hypergraph in [67] with 2M nodes and 15M hyperedges is converted to a bipartite graph with 17M nodes and 1B edges. Even worse, a k-uniform hypergraph with m hyperedges causes its clique graph to have $O(mk^2)$ edges.

To this end, we propose a novel neighborhood-cohesion based approach for hypergraph core decomposition. Neighborhood-based core decomposition is a decomposition of a hypergraph into nested, strongly-induced maximal subhypergraphs such that all the nodes in every subhypergraph have at least a certain number of neighbors in that subhypergraph. Being strongly-induced means that a hyperedge is only present in a subhypergraph if and only if all its constituent nodes are present in that subhypergraph.

EXAMPLE 1 (NEIGHBORHOOD-BASED CORE DECOMPOSITION). In the hypergraph H in Figure 1, the node a has 4 neighbors: $\{b, c, d, e\}$. Similarly, nodes b, c, d, e, and f have 3, 3, 3, 4, and I neighbors, respectively. As every node has ≥ 1 neighbor, the neighborhood-based 1-core, denoted by $H[V_1]$, is the hypergraph H

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itself. The neighborhood based 2-core is the subhypergraph $H[V_2] = \{\{a, c, d\}, \{c, d, e\}, \{a, b, e\}\}$ because nodes b, a, e, c, and d respectively have 2, 4, 4, 3, and 3 neighbors in $H[V_2]$.

Motivation. The only hypergraph decomposition existing in the

literature is that based on degree [51, 56]. The degree-based core decomposition decomposes a hypergraph into a sequence of nested maximal subhypergraphs (cores) such that every node in the k-th core has degree at least k in that core. Degree-based core decomposition does not take hyperedge sizes into consideration. As a result, nodes in the same core may have vastly different neighborhood sizes. For instance, nodes f and a have 1 and 4 neighbors, respectively, yet they belong to the same core in the degree-based decomposition, as illustrated in Figure 2. There are applications, e.g., propagation of contagions in epidemiology, diffusion of information in viralmarketing, where it is desirable to capture such differences, because nodes with the same number of neighbors in a subhypergraph are known to exhibit similar diffusion characteristics [40]. Indeed from an intuitive viewpoint, node f (as a seed) propagates information (disease) to only 1 node, whereas a can do the same to 4 nodes. Thus, targeting (intervening) node a should be more important than f for devising targeted campaigns (disease intervention strategies). Clearly, we need a new measure of node-importance different from degree, and hence a new decomposition which respects that measure. Applications. First, we demonstrate the usefulness of neighborhoodbased core decomposition in diffusion-related domains [10, 55]. Specifically, we show in § 6.1 that nodes in the innermost core of our decomposition are not only the most influential in spreading information, but also the earliest adopters of diffused information. Besides, deleting an innermost core node is more likely to disrupt the spread of an infectious disease (e.g., COVID19, Ebola) compared to an outer core node. We show such an intervention to be generally effective due to decrease in connectivity and increase in the average length of shortest paths after intervention. Furthermore, an intervention based on our neighborhood-based decomposition is much more effective compared to a degree-based decomposition strategy as discussed in the Motivation.

Second, the proposed core-decomposition gives rise to a new type of densest subhypergraph, which we refer to as the *volume-densest subhypergraph*. Nodes in the volume-densest subhypergraph have the largest number of average neighbors (in the subhypergraph) among all subhypergraphs. Our neighborhood-based decomposition induces a node-ordering which we exploit to obtain the volume-densest subhypergraph approximately with a theoretical guarantee (§6.2). In §6.2.1, we show that the proposed volume-densest subhypergraphs capture neighborhood-cohesive regions more effectively than the existing degree-densest subhypergraphs [35]. Case-study on human protein-complexes (§6.2.2) shows that the volume-densest subhypergraph extracts complexes that participate in RNA metabolism and localization. Case-study on organizational emails (§6.2.3) shows that the volume-densest subhypergraph extracts emails about internal announcements, meetings, and employee gatherings.

Challenges. Hypergraph core decomposition is challenging because a hyperedge can relate to more than two nodes. Furthermore, a pair of nodes may be related by multiple, yet distinct hyperedges. Thus, trivial adaptation of core-decomposition algorithms for graphs to hypergraphs is difficult. For instance, in a neighborhood-based

hypergraph core decomposition, deleting a node may reduce the neighborhood size of its neighboring node by more than 1. Hence, to recompute the number of neighbors of a deleted node's neighbor, one must construct the residual hypergraph after deletion, which is expensive. In the following, we discuss the challenges associated with adopting the local approach [46], one of the most efficient methods for graph core decomposition, to hypergraphs.

Challenges in adopting a local approach [46, 50]. In this approach, a core-number estimate is updated iteratively [46] or in a distributed manner [50] for every node in a graph. The initial value of a node's core-number estimate is a known upper bound of its core-number. In subsequent rounds, this estimate is iteratively decreased based on estimates of neighboring nodes. [46] uses Hirsch's index (h-index) [34] for such an update. They have shown that the following invariant must hold: every node with core-number k has h-index at least k, and the subgraph induced by nodes with h-index at least k has at least k neighbors per node in that subgraph. The former holds but the later may not hold in a hypergraph, because the subhypergraph induced by nodes with h-index at least k may not include hyperedges that partially contain other nodes. As we illustrate in §3.3, due to those 'missing' hyperedges, the number of neighbors of some nodes in that subhypergraph may drop below k violating the coreness condition. Therefore, while the local approach is used for computing the k-core [46] or more general (k, h)-core [45] in graphs, the same approach results in incorrect neighborhood-based hypergraph cores, as demonstrated empirically in §5.1.

Our contributions and roadmap. Our contributions are summarized as follows:

Novel problem and characterization (§ 2). We are the first to define and investigate the novel problem of neighborhood-based core decomposition in hypergraphs. Our core decomposition is well-defined and logical: We prove that neighborhood-based k-cores are unique, and the k-core contains the (k+1)-core.

Exact algorithms (§ 3). We propose three exact algorithms to compute neighborhood-based cores in a hypergraph, with their formal correctness and time complexity analyses. Two of them, **Peel** and its enhancement **E-Peel** adopt the classic peeling approach [9] incurring global changes to the hypergraph. For **E-Peel**, we derive *novel lower-bound* on core-number that eliminates many redundant neighborhood recomputations. Our third algorithm, called **Local-core** is the most efficient one, it only makes node-level local computations. Even though the existing local method [46, 50] fails to correctly find neighborhood-based core-numbers in a hypergraph, our algorithm **Local-core** applies a *novel* **Core-correction** procedure after local updates, ensuring correct core-number computations.

Optimization and parallelization strategies (§ 4). We propose four optimization strategies to improve the efficiency of **Local-core**. Compressed representations for hypergraph (optimization-I) and the family of optimizations for efficient **Core-correction** (optimization-II) are novel to core-decomposition literature. The other optimizations, though inspired from graph literature, have not been adopted in earlier hypergraph-related works. We also propose parallelization of **Local-core** for the shared-memory programming paradigm.

Empirical evaluation (§ 5). Empirical evaluation on real and synthetic hypergraphs shows that the proposed algorithms are effective, efficient, and practical. **Local-core** with optimizations decomposes

hypergraph with 27M nodes and 17M hyperedges within 16 minutes. Furthermore, our OpenMP parallel implementation **Local-core(P)** achieves 2-5x speedup compared to its sequential counterpart.

Applications (§ 6). In diffusion-related applications, we show our decomposition to be more effective in disrupting diffusion than other decompositions. Such diffusion-interruption is significant for intervening epidemics or selecting target-groups for marketing.

Our greedy algorithm proposed for the volume-densest subhypergraph recovery achieves a $(d_{pair}(d_{card}-2)+2)$ -approximation guarantee, where hyperedge-cardinality and node-pair co-occurrence are at most d_{card} and d_{pair} , respectively. If the hypergraph is a graph $(d_{card}=2)$, our result generalizes Charikar's 2-approximation guarantee for the densest subgraph discovery [15]. The proposed volume-densest subhypergraphs capture neighborhood-cohesive regions more effectively than existing degree-densest subhypergraphs [35]. Our case-studies show that the proposed subhypergraphs captures functionally significant complexes in human cells and important emails in organizations.

2 OUR PROBLEM AND CHARACTERIZATION

Hypergraph. A hypergraph H = (V, E) consists of a set of nodes V and a set of hyperedges $E \subseteq P(V) \setminus \phi$, where P(V) is the power set of V. A hyperedge is modeled as an unordered set of nodes.

Neighbors. Neighbors N(v) of a node v in a hypergraph H = (V, E) is the set of nodes $u \in V$ that co-occur with v in some hyperedge $e \in E$. That is, $N(v) = \{u \in V \mid u \neq v \land \exists e \in E \text{ s.t. } u, v \in e\}$.

Strongly induced subhypergraph [6, 17, 31]. A strongly induced subhypergraph H[S] of a hypergraph H = (V, E), induced by a node set $S \subseteq V$, is a hypergraph with the node set S and the hyperedge set $E[S] \subseteq E$, consisting of all the hyperedges that are subsets of S.

$$H[S] = (S, E[S]), \text{ where } E[S] = \{e \mid e \in E \land e \subseteq S\}$$
 (1)

In other words, every hyperedge in a strongly induced subhypergraph must exist in its parent hypergraph.

2.1 Problem Formulation

Nbr-*k***-core.** The *nbr-k*-*core* $H[V_k] = (V_k, E[V_k])$ of a hypergraph H = (V, E) is the maximal (strongly) induced subhypergraph such that every node $u \in V_k$ has at least k neighbours in $H[V_k]$. For simplicity of notation, we denote nbr-*k*-core, $H[V_k]$ as H_k .

The maximum core of H is the largest k for which H_k is nonempty. The core-number c(v) of a node $v \in V$ is the largest k such that $v \in V_k$ and $v \notin V_{k+1}$. The core decomposition of a hypergraph assigns to each node its core-number. Given a hypergraph, the problem studied in this paper is to correctly and efficiently compute its neighborhood-based core decomposition.

2.2 Differences with Other Core Decompositions

There are broadly two kinds of approaches that one can adapt from the literature towards decomposing hypergraphs.

Approach-1. One may transform the hypergraph into other objects (e.g., a graph), apply existing decomposition approaches [9, 14] on that object, and then project the decomposition back to the hypergraph. For instance, a hypergraph can be transformed into a clique graph by replacing the hyperedges with cliques and then classical graph decomposition is applied (Figure 3). A hypergraph can also be

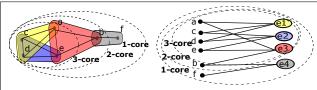


Figure 3: Core decomposition of the clique-graph of *H*

Figure 4: Distance-2 core decomposition of the bipartite graph of H

transformed into a bipartite graph by representing the hyperedges as nodes in the second partition and creating an edge between two crosspartition nodes if the hyperedge in the second partition contains a node in the first partition. Finally, distance-2 core decomposition is applied (Figure 4). In distance-2 core-decomposition, nodes in k-core has at least k 2-hop neighbors in the subgraph. Both clique graph and bipartite graph representations inflate the problem size as discussed in the Introduction. Besides, the decomposition they yield may be different from that yielded by ours. For instance, core(d) = 3 in both clique graph decomposition and dist-2 bipartite graph decomposition, whereas d has core-number 2 in our decomposition.

Approach-2. Sun et al. [56] define the k-core (i.e., deg-k-core) of a hypergraph from the notion of node degree in a hypergraph. The $degree\ d(v)$ of a node v in hypergraph H is the number of hyperedges incident on v [12], i.e., $d(v) = |\{e \in E \mid v \in e\}|$. The deg-k-core H_k^{deg} of a hypergraph H is the maximal (strongly) induced subhypergraph of H such that every node u in H_k^{deg} has degree at least k in H_k^{deg} . This approach does not take hyperedge sizes into consideration as stated in §1. Therefore, it does not necessarily yield the same decomposition as our approach, which can be seen from the difference between Figures 2 and 1. This difference originates from the fact that, unlike in a graph, the number of neighbors (|N(v)|) of a node v in a hypergraph is not equal to its degree.

2.3 Nbr-k-Core: Properties

THEOREM 1. The nbr-k-core H_k is unique for any k > 0.

PROOF. Let, if possible, there be two distinct nbr-k-cores: $H_{k_1} = (V_{k_1}, E[k_1])$ and $H_{k_2} = (V_{k_2}, E[k_2])$ of a hypergraph H = (V, E). By definition, both H_{k_1} and H_{k_2} are maximal strongly induced subhypergraphs of H. Construct the union hypergraph $H_k = (V_{k_1} \cup V_{k_2}, E[V_{k_1} \cup V_{k_2}])$. For any $u \in V_{k_1} \cup V_{k_2}$, u must be in either V_{k_1} or V_{k_2} . In both cases, u must have at least k neighbours in the respective subhypergraph H_{k_1} or H_{k_2} . Since $E[V_{k_1}] \cup E[V_{k_2}] \subseteq E[V_{k_1} \cup V_{k_2}]$, u must also have at least k neighbours in H_k . Since H_k is a supergraph of both H_{k_1} and H_{k_2} , it follows that, H_{k_1} and H_{k_2} are not maximal, and thus are not nbr-k-cores, leading to a contradiction.

THEOREM 2. The (k+1)-core is contained in the k-core, $\forall k > 0$.

PROOF. Let, if possible, for some node $u \in V_{k+1}$, $u \notin V_k$. Construct $S = V_k \cup V_{k+1}$. Since $u \notin V_k$, but $u \in V_{k+1} \subset S$, the set S is larger than V_k , to be precise, $|S| \ge |V_k| + 1$. It is easy to verify that every node $v \in S$ has at least k neighbours in H[S] and $|S| > |V_k|$. Then, V_k is not maximal and thus not nbr-k-core, which is a contradiction. The theorem follows.

Algorithm 1 Peeling algorithm: Peel

```
\begin{array}{lll} \textbf{Input:} & \text{Hypergraph } H = (V, E) \\ \textbf{Output:} & \text{Core-number } c(u) \text{ for each node } u \in V \\ \textbf{1:} & \textbf{ for all } u \in V \textbf{ do} \\ \textbf{2:} & \text{Compute } N_V(u) \\ \textbf{3:} & B[|N_V(u)|] \leftarrow B[|N_V(u)|] \cup \{u\} \\ \textbf{4:} & \textbf{ for all } k = 1, 2, \dots, |V| \textbf{ do} \\ \textbf{5:} & \textbf{ while } B[k] \neq \phi \textbf{ do} \\ \textbf{6:} & \text{Remove a node } v \text{ from } B[k] \\ \textbf{7:} & c(v) \leftarrow k \\ \textbf{8:} & \textbf{ for all } u \in N_V(v) \textbf{ do} \\ \textbf{9:} & \text{Move } u \text{ to } B[\max\left(|N_{V\setminus \{v\}}(u)|, k\right)] \\ \textbf{10:} & V \leftarrow V \setminus \{v\} \\ \textbf{11:} & \text{ return } c \\ \end{array}
```

3 ALGORITHMS

We propose three algorithms to exactly compute neighborhood-based hypergraph cores. The algorithms **Peel** and its efficient variant **E-Peel** are inspired by a family of peeling-based algorithms similar to graph core computations [9, 14]. The algorithm **Local-core** is inspired by a family of local approaches to graph core computation [46, 50]. The algorithms **E-Peel** and **Local-core**, despite being inspired by the existing family of graph algorithms, are by no means trivial adaptations. For **E-Peel**, we devise a new local lower-bound for core-numbers because hypergraphs generalize graphs and the lower-bound for graph core is naturally insufficient for our purpose. For **Local-core**, we illustrate how a direct adaptation of local algorithm [46] may lead to incorrect core computations. Hence, we devise the notions of *hypergraph h-index* and *local coreness constraint* and employ them to compute hypergraph cores correctly.

3.1 Peeling Algorithm

Following Theorem 2, the (k+1)-core can be computed from the k-core by "peeling" all nodes whose neighborhood sizes are less than k+1. Algorithm 1 describes our peeling algorithm: **Peel**, which processes the nodes in increasing order of their neighborhood sizes (Lines 4-10). B is a vector of lists: Each cell B[i] is a list storing all nodes whose neighborhood sizes are i (Line 3). When a node v is processed at iteration k, its core-number is assigned to c(v) = k (Line 7), it is deleted from the set of "remaining" nodes V (Line 10). The neighborhood sizes of the nodes in v's neighborhood are recomputed (each neighborhood size can decrease by more than 1, since when v is deleted, all hyperedges involving v are also deleted), and these nodes are moved to the appropriate cells in v0. The algorithm completes when all nodes in the hypergraph are processed and have their respective core-numbers computed.

Proof of correctness. Initially B[i] contains all nodes whose neighborhood sizes are i. When we delete some neighbor of a node u, the neighborhood size of u is recomputed, and u is reassigned to a new cell corresponding to its reduced neighborhood size until we find that the removal of a neighbor v of u reduces u's neighborhood size even below the current iteration number k (Line 9). When this happens, we correctly assign u's core-number c(u) = k. (1) Consider the remaining subhypergraph formed by the remaining nodes and hyperedges at the end of the $(k-1)^{th}$ iteration. Clearly, u is in the k-core since u has at least k neighbors in this remaining subhypergraph, where all nodes in the remaining subhypergraph also have neighborhood sizes $\geq k$. (2) The removal of v decreases u's neighborhood size smaller than the current iteration number k, thus

Algorithm 2 Efficient peeling algorithm with bounding: E-Peel

```
Input: Hypergraph H = (V, E)
Output: Core-number c[u] for each node u \in V
  1: for all u \in V do
           Compute LB(u)
           B[LB(u)] \leftarrow B[LB(u)] \cup \{u\}
setLB(u) \leftarrow True
  3.
  4:
  5:
      for all k = 1, 2, ..., |V| do while B[k] \neq \phi do
  6:
7:
8:
                 if setLB(v) then
                     B[|N_V(v)|] \leftarrow B[\max(|N_V(v)|,k)] \cup \{v\}
10:
11:
                     setLB(v) \leftarrow False
                     c(v) \leftarrow k
 12:
 13:
                     for all u \in N_V(v) do
 14:
                         if \neg setLB(u) then
                              Move u to B\left[\max\left(|N_{V\setminus\{v\}}(u)|,k\right)\right]
 15:
16:
                     V \leftarrow V \setminus \{v\}
17: return c
```

when the current iteration number increases to k + 1, u will not have enough neighbors to remain in the (k + 1)-core.

Time complexity. Each node v is processed exactly once from B in Algorithm 1; when it is processed and thereby deleted from V, neighborhood sizes of the nodes in v's neighborhood are recomputed. Assume that the maximum number of neighbors and hyperedges of a node be d_{nbr} and d_{hpe} , respectively. Thus, Algorithm 1 has time complexity $O(|V| \cdot d_{nbr} \cdot (d_{nbr} + d_{hpe}))$.

3.2 Efficient Peeling with Bounding

An inefficiency in Algorithm 1 is that it updates the cell index of every node u that is a neighbor of a deleted node v. To do so, it has to compute the number of neighbors of u in the newly constructed subhypergraph. Can we delay this recomputation for some neighbors u of v? We derive a local lower-bound for c(u) via Lemma 1 and use it to eliminate many redundant neighborhood recomputations and cell updates (Algorithm 2). The intuition is that a node u will not be deleted at some iteration u0 the lower-bound on u0, thus we do not require computing u0 s neighborhood size until the value of u0 reaches the lower-bound on u0. Our lower-bound is local since it is specific to each node in the hypergraph.

LEMMA 1 (LOCAL LOWER-BOUND). Let $e_m(v) = \arg \max\{|e| : e \in E \land v \in e\}$ be the highest-cardinality hyperedge incident on $v \in V$. For all $v \in V$,

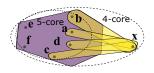
$$c(v) \ge \max\left(|e_m(v)| - 1, \min_{u \in V} |N(u)|\right) = \mathsf{LB}(v) \tag{2}$$

PROOF. Notice that $c(v) \geq \min_{u \in V} |N(u)|$, since all nodes in the input hypergraph must be in the $(\min_{u \in V} |N(u)|)$ -core. Next, we show that $c(v) \geq |e_m(v)|-1$, by contradiction. Let, if possible, $|e_m|-1 > c(v)$. This implies that v is not in the $(|e_m|-1)$ -core, denoted by $H[V_{|e_m|-1}]$. Consider $V' = V_{|e_m|-1} \cup \{u: u \in e_m\}$. Clearly, $|V'| \geq |V_{|e_m|-1}| + 1$, since $v \notin V_{|e_m|-1}$, but $v \in e_m$, so $v \in V'$. We next show that $H[V_{|e_m|-1}]$ is not the maximal subhypergraph where every node has at least $|e_m|-1$ neighbors, which is a contradiction.

To prove non-maximality of $H[V_{|e_m|-1}]$, it suffices to show that for any $u \in V'$, $N_{V'}(u) \ge |e_m| - 1$. If $u \in V_{|e_m|-1} \subset V'$, $|N_{V'}(u)| \ge |N_{V_{|e_m|-1}}(u)| \ge |e_m| - 1$. If $u \in e_m$, $N_{V'}(u) \ge N_{e_m}(u) = |e_m| - 1$.

Since our premise $|e_m| - 1 > c(v)$ contradicts the fact that $H[V_{|e_m|-1}]$ is the $(|e_m|-1)$ -core, $|e_m|-1 \le c(v)$.

4



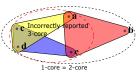


Figure 5: (a) During x's core-number computation, Algorithm 2 does not perform neighborhood recomputations and cell updates for x's neighbors $\{a,b,c,d\}$; thus saving four redundant neighborhood recomputations and cell updates. (b) For any n>1, the h-index (Definition 2) of node a never reduces from $h_a^{(1)}=\mathcal{H}(2,3,3,4)=3$ to its core-number 2: $\lim_{n\to\infty}h_a^{(n)}=3\neq$ core-number of a. Because a will always have at least 3 neighbors (c,d, and e) whose h-indices are at least 3. As a result, the naïve approach reports an incorrect 3-core.

Algorithm. Our efficient peeling approach is given in Algorithm 2: **E-Peel**. In Line 14, we do not recompute neighborhoods and update cells for those neighboring nodes u for which setLB is True, thereby improving the efficiency. setLB is True for nodes for which LB() is known, but $N_V()$ at the current iteration is unknown.

Proof of correctness. The proof of correctness follows that of Algorithm 1. When a node v is extracted from B[k] at iteration k, we check setLB(v). (1) Lemma 1 ensures that, if we extract a node v from B[k] and setLB(v) is True, then $c(v) \geq k$. In that case, we compute the current value of $N_V()$, where V denotes the set of remaining nodes, and insert v into the cell: $B[\max{(|N_V(v)|, k)}]$. We also set setLB(v) = False, implying that $N_V(v)$ at the current iteration is known. (2) In contrast, if we extract a node v from B[k] and setLB(v) is False, this indicates that c(v) = k, following the same arguments as in Algorithm 1. In this case, we correctly assign v's core-number to k, and v is removed from V. Moreover, for those neighbors v of v for which v is True, implying that v is v we appropriately delay recomputing their neighborhood sizes.

Time complexity. Following similar analysis as in Algorithm 1, **E-Peel** has time complexity $O(\alpha \cdot |V| \cdot d_{nbr} \cdot (d_{nbr} + d_{hpe}))$, where $\alpha \le 1$ is the ratio of the number of neighborhood recomputations in Algorithm 2 over that in Algorithm 1. Based on our experimental results in § 5, **E-Peel** can be up to 14x faster than **Peel**.

3.3 Local Algorithm

Although **Peel** and its more efficient variant **E-Peel** correctly computes core-numbers, they must modify the remaining hypergraph at

every iteration by peeling (deleting) nodes and hyperedges. Peeling operation may impact the hypergraph data structure globally and must be performed in sequence. Thus, there is little scope for making **Peel** and **E-Peel** more efficient via parallelization. Furthermore, they are not suitable in a time-constrained setting where a high-quality partial solution is sufficient. We propose a novel local algorithm that is able to provide partial solutions, amenable to a number of optimizations, as well as parallelizable.

Naïve adoption of local algorithm in hypergraphs: a negative result. Eugene et al. [46] adopt Hirsch's index [34], popularly known as the h-index (Definition 1), to propose a local algorithm for core computation in graphs. This algorithm relies on a recurrence relation that defines higher-order h-index (Definition 2). The local algorithm for graph core computation starts by computing h-indices of order 0 for every node in the graph. At each iteration n > 0, it computes order-n h-indices using order-(n-1) h-indices computed in the previous iteration. It is well-known that higher-order h-indices monotonically converge to the core-number of that node in the graph.

DEFINITION 1 (\mathcal{H} -OPERATOR [34, 46]). Given a finite set of positive integers $\{x_1, x_2, ..., x_t\}$, $\mathcal{H}(x_1, x_2, ..., x_t) = y > 0$, where y is the maximum integer such that there exist at least y elements in $\{x_1, x_2, ..., x_t\}$, each of which is at least y.

EXAMPLE 3 (*H*-OPERATOR). $\mathcal{H}(1,1,1,1)=1$, $\mathcal{H}(1,1,1,2)=1$, $\mathcal{H}(1,1,2,2)=2$, $\mathcal{H}(1,2,2,2)=2$, $\mathcal{H}(1,2,3,3)=2$, $\mathcal{H}(1,3,3,3)=3$

DEFINITION 2 (h-INDEX OF ORDER n [46]). Let $\{u_1, u_2, \ldots, u_t\}$ be the set of neighbors of node $v \in V$ in graph G = (V, E). The n-order h-index of node $v \in V$, denoted as $h_v^{(n)}$, is defined for any $n \in \mathbb{N}$ by the recurrence relation

$$h_v^{(n)} = \begin{cases} |N(v)| & n = 0\\ \mathcal{H}\left(h_{u_1}^{(n-1)}, h_{u_2}^{(n-1)}, \dots, h_{u_t}^{(n-1)}\right) & n \in \mathbb{N} \setminus \{0\} \end{cases}$$
(3)

For neighborhood-based hypergraph core decomposition via local algorithm, we define $h_n^{(0)}$ as the number of neighbors of node v in hypergraph H = (V, E) (instead of graph G). The definition of $h_n^{(n)}$ for n > 0 remains the same. However, this direct adoption of local algorithm to compute hypergraph cores does not work. Although one can prove that the sequence $(h_v^{(n)})$ adopted for hypergraph has a limit, that value in-the-limit is not necessarily the core-number c(v)for every $v \in V$. For some node, the value in-the-limit of its h-indices is strictly greater than the core-number of that node. The reason is as follows. \mathcal{H} -operator acts as both necessary and sufficient condition for computing graph cores. It has been shown that the subgraph induced by G[S], where S contains all neighbors u of a node vsuch that $h_u^{(\infty)} \ge c(v)$, satisfies $h_v^{(\infty)} = c(v)$ [46, p.5 Theorem 1]. However, Definition 2 is not sufficient to show $h_n^{(\infty)} = c(v)$ for a hypergraph. Because it is not guaranteed that v will have at least h_v^{∞} neighbors in the subhypergraph $H[\{u: h_u^{\infty} \geq h_u^{\infty}\}]$ that is reported as the c(v)-core. So, thr reported c(v)-core can be incorrect.

EXAMPLE 4. For the hypergraph in Figure 5(b), The values inthe-limit of h-indices are $h_a^{(\infty)} = h_c^{(\infty)} = h_d^{(\infty)} = h_e^{(\infty)} = 3$ and $h_b^{(\infty)} = 2$. No matter how large n is chosen, Equation (3) does not help $h_a^{(n)}$ to reach the correct core-number (=2) for a. Three neighbors of node a, namely c, d, and e have their h^{∞} -values at least

5

Algorithm 3 Local algorithm with local coreness constraint: Local-core

```
\begin{array}{lll} \text{Input:} & \text{Hypergraph } H = (V, E) \\ \text{Output:} & \text{Core-number } c [v] \text{ for each node } v \in V \\ 1: & \text{ for all } v \in V \text{ do} \\ 2: & \hat{h}_v^{(0)} = h_v^{(0)} \leftarrow |N(v)|. \\ 3: & \text{ for all } n = 1, 2, \ldots, \infty \text{ do} \\ 4: & \text{ for all } v \in V \text{ do} \\ 5: & h_v^{(n)} \leftarrow \min \left( \mathcal{H}(\{\hat{h}_u^{(n-1)} : u \in N(v)\}), \hat{h}_v^{(n-1)} \right) \\ 6: & \text{ for all } v \notin V \text{ do} \\ 7: & c[v] \leftarrow \hat{h}_v^{(n)} \leftarrow \text{Core-correction } (v, h_v^{(n)}, H) \\ 8: & \text{ if } \forall v, \hat{h}_v^{(n)} = h_v^{(n)} \text{ then} \\ 9: & \text{Terminate Loop} \\ 10: & \text{return } c \end{array}
```

 $3 = h_v^{\infty}$ in Figure 5(b). But, a does not have at least 3 neighbors in the subhypergraph $H[\{a,c,d,e\}] = H[\{u:h_u^{\infty} \geq h_v^{\infty}\}]$. Thus, the 3-core $H[\{a,c,d,e\}]$ reported by the naïve h-index based local approach is incorrect. The reason is that a and e are no longer neighbors to each other in $H[\{a,c,d,e\}]$ due to the absence of b.

Local algorithm with local coreness constraint. Motivated by the observation mentioned above, we define a constraint as a sufficient condition, upon satisfying which we can guarantee that for every node v, 1) the sequence of its h-indices converges and 2) the value in-the-limit $h_v^{(\infty)}$ is such that v has at least $h_v^{(\infty)}$ neighbors in the subhypergraph induced by $H[u:h_u^{(\infty)} \geq h_v^{(\infty)}]$. The first condition is critical for algorithm termination. The second condition is critical for correct computation of core-numbers as discussed in Example 4.

DEFINITION 3 (LOCAL CORENESS CONSTRAINT (LCC)). Given a positive integer k, for any node $v \in V$, let $H^+(v) = (N^+(v), E^+(v))$ be the subhypergraph of H such that for any n > 0

$$E^{+}(v) = \{ e \in Incident(v) : h_u^{(n)} \ge k, \forall u \in e \}$$

$$N^{+}(v) = \{ u : u \in e, \forall e \in E^{+}(v) \} \setminus \{v\}$$

$$(4)$$

Local coreness constraint (for node v) is satisfied at k, denoted as LCCSAT(k), iff $\exists H^+(v)$ contains at least k nodes, i.e., $|N^+(v)| \ge k$. Here, Incident(v) is the set of hyperedges incident on v.

We define *Hypergraph h-index* based on the notion of *LCCSAT* and a re-defined recurrence relation for $h_n^{(n)}$.

DEFINITION 4 (HYPERGRAPH h-INDEX OF ORDER n). The Hypergraph h-index of order n for node v, denoted as $\hat{h}_v^{(n)}$, is defined for any natural number $n \in \mathbb{N}$ by the following recurrence relation:

$$\hat{h}_{v}^{(n)} = \begin{cases} |N(v)| & n = 0\\ h_{v}^{(n)} & n > 0 \land LCCSAT(h_{v}^{(n)})\\ \max\{k \mid k < h_{v}^{(n)} \land LCCSAT(k)\} & n > 0 \land \neg LCCSAT(h_{v}^{(n)}) \end{cases}$$
(5)

 $h_v^{(n)}$ is a newly defined recurrence relation on hypergraphs:

$$h_{v}^{(n)} = \begin{cases} |N(v)| & n = 0\\ \min\left(\mathcal{H}\left(\hat{h}_{u_{1}}^{(n-1)}, \hat{h}_{u_{2}}^{(n-1)}, \dots, \hat{h}_{u_{t}}^{(n-1)}\right), \hat{h}_{v}^{(n-1)}\right) & n \in \mathbb{N} \setminus \{0\} \end{cases}$$

The recurrence relations in Equations (5) and (6) are coupled: $\hat{h}_v^{(n)}$ depends on the evaluation of $h_v^{(n)}$, which in turn depends on the evaluation of $\hat{h}_v^{(n-1)}$. Such inter-dependency causes both sequences to converge, as proven in our correctness analysis.

Local-core (Algorithm 3) initializes $h_v^{(0)}$ and $\hat{h}_v^{(0)}$ to |N(v)| for every node $v \in V$ (Lines 1-2) following Equation (6) and Equation (5),

Algorithm 4 Core-correction procedure

```
 \begin{array}{lll} \textbf{Input:} & \text{node } v,v \text{'s hypergraph } h \text{ index } h_v^{(n)}, \text{ hypergraph } H \\ 1: & \textbf{while } h_v^{(n)} > 0 \textbf{ do} \\ 2: & & \text{Compute } E^+(v) \leftarrow \{e \in Incident(v) : h_u^{(n)} \geq h_v^{(n)}, \forall u \in e\} \\ 3: & & \text{Compute } N^+(v) = \{u : u \in e, \forall e \in E^+(v)\} \setminus \{v\} \\ 4: & \textbf{if } |N^+(v)| \geq h_v^{(n)} \textbf{ then} \\ 5: & \text{return } h_v^{(n)} \\ 6: & \textbf{else} \\ 7: & & h_v^{(n)} \leftarrow h_v^{(n)} - 1 \end{array}
```

respectively. At every iteration n>0, Algorithm 3 first computes $h_v^{(n)}$ for every node $v\in V$ (Lines 4-5) following Equation (6). In order to decide whether the algorithm should terminate at iteration n (Lines 8-9), the algorithm computes $\hat{h}_v^{(n)}$ using Algorithm 4. Algorithm 4 checks for every node $v\in V$, whether $LCCSAT(h_v^{(n)})$ is True or False. Following Equation (5), if $LCCSAT(h_v^{(n)})$ is True it returns $h_v^{(n)}$; if $LCCSAT(h_v^{(n)})$ is False, a suitable value lower than $h_v^{(n)}$ is returned. The returned value $\hat{h}_v^{(n)}$ is considered as the estimate of core-number c(v) at that iteration (Line 7).

To compute $LCCSAT(h_v^{(n)})$, Algorithm 4 checks in Line 4 if the subhypergraph $H^+(v) = (N^+(v), E^+(v))$ constructed in Lines 2-3 contains at least $h_v^{(n)}$ neighbors of v. If v has at least $h_v^{(n)}$ neighbors in the subhypergraph $H^+(v)$, due to Equation (5) no correction to $h_v^{(n)}$ is required. In this case, Algorithm 4 returns $h_v^{(n)}$ in Line 5. If v does not have at least $h_v^{(n)}$ neighbors in the subhypergraph $H^+(v)$ (Line 4), $LCCSAT(h_v^{(n)})$ is False by Definition 3. Following Equation (5), a correction to $h_v^{(n)}$ is required. In search for a suitable corrected value lower than $h_v^{(n)}$ and a suitable subhypergraph $H^+(v)$, Line 7 keeps reducing $h_v^{(n)}$ by 1. Reduction to $h_v^{(n)}$ causes $|N^+(v)|$ to increase, while $h_v^{(n)}$ decreases, until the condition in Line 4 is satisfied. At some point a suitable subhypergraph must be found.

Theorem 4 proves that the numbers returned by Algorithm 3 at that point indeed coincide with the true core-numbers. The termination condition $\hat{h}_v^{(n)} = h_v^{(n)}$ must be satisfied at some point because Theorem 3 proves that $\lim_{n \to \infty} \hat{h}_v^{(n)} = \lim_{n \to \infty} h_v^{(n)} \ \forall v \in V$.

Example 5. Consider iteration n=1 of Algorithm 3, when the input to the algorithm is the hypergraph in Figure 5(b). The algorithm corrects the core-estimate $h_a^{(1)}=3$ to $\hat{h}_a^{(1)}=2$ in Line 7. Because in Line 4 of Core-correction, the algorithm finds that for $h_a^{(1)}=3$, a only has $|N^+(v)|=2$ neighbors in $H^+(v)=H[\{a,c,d,e\}]$ thus violating the condition that $|N^+(v)|>h_a^{(1)}$. Hence $h_a^{(1)}$ needs to be corrected to satisfy LCCSAT($h_a^{(1)}$). In Line 7 of Core-correction, it reduces $h_a^{(1)}$ by 1 and subsequently for $h_a^{(1)}=2$ the subhypergraph $H^+(a)=H$ indeed satisfies LCCSAT($h_a^{(1)}$). This is how Algorithm 3 corrects the case of incorrect core-numbers discussed in Example 4.

3.4 Theoretical Analysis of Local-core

Proof of Correctness. Algorithm 3 terminates after a finite number of iterations because for any $v \in V$, both sequences $(h_v^{(n)})$ and $(\hat{h}_v^{(n)})$ are finite and have the same limit by Theorem 3. At the limit, $\forall v \in V$, $\lim_{n \to \infty} h_v^{(n)} = \lim_{n \to \infty} \hat{h}_v^{(n)}$ holds and it follows from Theorem 4 that $\forall v \in V$, $\lim_{n \to \infty} h_v^{(n)} = \lim_{n \to \infty} \hat{h}_v^{(n)} = c(v)$. Due to limitation

of space, we only provide proof sketches, while the formal proofs are given in our extended version [3].

THEOREM 3. For any node $v \in V$ of a hypergraph H = (V, E), the two sequences $(h_v^{(n)})$ defined by Equation (6) and $(\hat{h}_v^{(n)})$ defined by Equation (5) have the same limit: $\lim_{n\to\infty} h_v^{(n)} = \lim_{n\to\infty} \hat{h}_v^{(n)}$.

Proof sketch. Construct a new sequence $(h\hat{h}_v)$ by interleaving components from $(h_v^{(n)})$ and $(\hat{h}_v^{(n)})$ as the following:

$$(h\hat{h}_{v}) = (h_{v}^{(0)}, \hat{h}_{v}^{(0)}, h_{v}^{(1)}, \hat{h}_{v}^{(1)}, h_{v}^{(2)}, \hat{h}_{v}^{(2)}, \hat{h}_{v}^{(2)} \dots)$$

It can be verified that this *interleave sequence* [59, Defn 680], denoted as $(h\hat{h}_v)$, is monontonically non-increasing and is lower-bounded by 0. By Monotone convergence theorem [7], $(h\hat{h}_v)$ has a limit. An interleave sequence has a limit if and only if its constituent sequence pairs are convergent and have the same limit [59]. Since $(h\hat{h}_v)$ has a limit, the sequences $(h_v^{(n)})$ and $(h_v^{(n)})$ converges to the same limit: $\lim_{n\to\infty} h_v^{(n)} = \lim_{n\to\infty} \hat{h}_v^{(n)}$.

THEOREM 4. If the local coreness-constraint is satisfied for all nodes $v \in V$ at the terminal iteration, the corrected h-index at the terminal iteration $\hat{h}_v^{(\infty)}$ satisfies: $\hat{h}_v^{(\infty)} = c(v)$.

Proof sketch. Given any $v \in V$, we first show that $c(v) \ge \hat{h}_v^\infty$, followed by showing $\hat{h}_v^\infty \ge c(v)$. The former holds since it can be proved that $v \in \hat{h}_v^\infty$ -core. However, by definition of core-number, c(v) is the largest integer for which $v \in c(v)$ -core. Thus, $c(v) \ge \hat{h}_v^\infty$. Next, $\hat{h}_v^\infty \ge c(v)$ can be proved by induction on the number of iteration. Hence, the theorem.

Time complexity. Assume that Algorithm 3 terminates at iteration τ of the for-loop at Line 3. Each iteration has time-complexity $O(\sum_u |N(u)| (d(u) + |N(u)|) + \sum_u |N(u)|)$, the first term is due to Lines 6-7 and the second term is due to Lines 4-5. Computation of \mathcal{H} -operator requires hypergraph h-indices of u's neighbors and can be done in linear-time: O(|N(u)|). Core-correcting u requires at most |N(u)| iterations of while-loop (Line 1, Algorithm 4), at each iteration the construction of $E^+(u)$ costs O(d(u) + |N(u)|). Thus, **Local-core** has time complexity $O(\tau * (\sum_u d(u)|N(u)| + |N(u)|^2))$.

4 OPTIMIZATION AND PARALLELIZATION OF THE LOCAL-CORE ALGORITHM

We propose four optimizations to further improve the efficiency of **Local-core** (§3.3). Algorithm 5 presents the pseudocode for the optimized algorithm, **Local-core(OPT)** where all four optimizations are indicated. Optimization-I adopts sparse representations to efficiently evaluate neighborhood queries in hypergraphs, while Optimization-II consists of three implementation-specific improvements to efficiently perform **Core-correction**. Optimizations-I and II have not been used in earlier core-decomposition works for both graphs and hypergraphs. Our Optimizations-III and IV are motivated from [45], where similar optimizations are proposed for graph (k, h)-core decomposition to improve convergence and eliminate redundant computations, respectively, though such optimizations have not been adopted in earlier hypergraph-related works.

Algorithm 5 Optimized Local algorithm: Local-core(OPT)

```
Input: Hypergraph H = (V, E)
Output: Core-number c[v] for each node v \in V
  1: Construct CSR representations /* Opt-I */
            Compute LB(v) /* local lower-bounds for Opt-IV */
            c[v] \leftarrow \hat{h}_v^{(0)} \leftarrow h_v^{(0)} \leftarrow |N(v)| /* \text{ core-estimate c[v] initialized for Opt-III*} /
       for all n = 1, 2, ..., \infty do
            for all v \in V do

if h_v^{(n)} > LB(v) then /* Opt-IV*/
  7:
                      c[v] \leftarrow h_v^{(n)} \leftarrow \min\left(\mathcal{H}(\{c[u] : u \in N(v)\}), c[v]\right) /* \text{Opt-III*}/
  8:
  9:
 10:
                 if h_n^{(n)} > LB(v) then /* Opt-IV */
                      c[v] \leftarrow \hat{h}_{v}^{(n)} \leftarrow \text{Core-correction } (v, h_{v}^{(n)}, \mathcal{H}) \text{ /* Opt-II & Opt-III*/}
11:
            if \forall v, \hat{h}_v^{(n)} == h_v^{(n)} then
12:
13:
                 Terminate Loop
14: return c
```

Optimization-I (Compressed hypergraph representation): Local-core makes two primitive neighborhood queries on hypergraph structures: neighbors enumeration (for h-index computation) and incident-hyperedges enumeration (during Core-correction). A naive implementation keeps an $N \times N$ matrix for neighbors counting queries and an $N \times E$ matrix for incident-hyperedge queries. However, storing such matrices in main memory is not only expensive, but also unnecessary for large hypergraphs, since these matrices are sparse in practice. Hence, it is imperative to adopt a compressed sparse representation for these matrices. *Compressed sparse row* (CSR) is one such widely-used representation in scientific computations.

CSR representation for storing neighbors: We use two arrays F and N. N[v] stores the starting index in F containing neighbors of node v. Neighbors of node v are stored in contiguous locations $(F[N[v]], F[N[v]+1], \ldots, F[N[v+1]-1])$. N[v+1]-N[v] gives us the number of neighbors |N(v)| of node v.

CSR representation for storing incident-hyperedges works analogously. Alternatively, one can use hash tables for storing a node's neighbors and incident hyperedges. In §5.2, we empirically show that **Local-core + Optimization-I** is more efficient than **Local-core** which uses hash tables on large datasets.

Optimization-II (Efficient Core-correction and LCCSAT): We design three optimization methods for more efficient Core-correction. Hyperedge-index for efficient E⁺ computation: In Line 2 of Core**correction** (Algorithm 4), we check if $h_u^{(n)} \ge h_v^{(n)}$ for every node $u \in e$ such that $e \in Incident(v)$. This computation incurs O(d(v))|N(v)|) at every while-loop (Line 1) of **Core-correction**. We reduce this cost to O(d(v)) by maintaining an index E_e for hyperedges. $E_e^{(n)}$ records for every hyperedge $e \in E$ the minimum of h-indices of its constituent nodes (min_{$u \in e$} $h_u^{(n)}$). We compute $E^+(v)$ by traversing only the incident hyperedges whose $E_e^{(n)} \ge h_v^{(n)}$. Storing E_e for all hyperedges costs O(|E|) space and constructing the indices costs $O(\sum_{e \in E} |e|)$ time. However, hyperedge-indices are constructed only once before every iteration in Local-core (Line 3, Algorithm 3). Moreover, hyperedge-index helps efficiently compute N^+ as follows. Dynamic programming for efficient N^+ computation: The while loop in the **Core-correction** procedure computes $k = h_v^{(n)}, h_v^{(n)}$ – 1,..., $\hat{h}_v^{(n)}$; and for every $h_v^{(n)} \leq k \leq \hat{h}_v^{(n)}$, it recomputes $E^+(v)$ and $N^+(v)$ until returning $\hat{h}_v^{(n)}$ as output. The cost of computing $N^+(v)$ for every k, without any optimization, is $O(\sum_i |e_v^j|)$, where

 e_v^j is the j^{th} hyperedge incident on v. Thus, the total cost of the **Core-correction** procedure, without any optimization, is $O((h_v^{(n)} - \hat{h}_v^{(n)}) \sum_j |e_v^j|)$. We reduce this cost to $O(\sum_j |e_v^j|)$ by constructing an index B such that $B[h_v^{(n)}]$ records the set of incident hyperedges whose hyperedge-index $E_e^{(n)} \ge h_v^{(n)}$, and for every $k < h_v^{(n)}$, B[k] records the incident hyperedges whose hyperedge-index $E_e^{(n)} = k$.

Let us denote $E^+(v)$ and $N^+(v)$ at k as $E^+(v,k)$ and $N^+(v,k)$, respectively. Exploiting the index structure B, we have the following dynamic programming paradigm for efficiently computing $N^+(v,k)$.

$$N^{+}(v,k) = \begin{cases} \bigcup_{e} B[k] & k = h_{v}^{(n)} \\ N^{+}(v,k+1) \cup (\bigcup_{e} B[k]) & \hat{h}_{v}^{(n)} \le k < h_{v}^{(n)} \end{cases}$$
(7)

Instead of traversing all incident hyperedges at every $\hat{h}_v^{(n)} \leq$ $k \le h_v^{(n)}$, we only traverse hyperedges at B[k] to compute $N^+(v,k)$. Since the indices B[k] are mutually exclusive, each incident hyperedge is traversed at most once during the entire Core-correction procedure. This reduces the runtime complexity of Core-correction to $O(\sum_j |e_v^j|)$. The storage cost of B is $O(h_v^{(n)} + d(v))$, as there are at most $h_n^{(n)}$ keys in B and exactly d(v) hyperedges are stored in B. Efficient LCCSAT computation: We return True immediately upon finding the first hyperedge $e \in Incident(v)$ adding which to $E^+(v)$ causes $|N^+(v)| \ge h_v^{(n)}$. Adding subsequent incident hyperedges to $E^+(v)$ increases $|N^+(v)|$ more without affecting LCCSAT(k)=True. Optimization III (Faster convergence): In Line 8, Algorithm 5, we use the most-recent core-estimates $c[u_1], c[u_2], ... c[u_t]$ to update the node $v \in N(u_i)$'s core-number estimate c[v] (and vice versa): $c[v] \leftarrow \min(\mathcal{H}(c[u_1], c[u_2], ..., c[u_t]), c[v])$. Notice that due to the min() operator, c[v] and $c[u_i]$'s are non-increasing with more iterations. Assuming that $c[u_1]$ is updated before c[v], c[v] might decrease more if v uses the most-recent $c[u_1]$ instead of using $c[u_1]$ from the previous iteration. The reason for faster decrease is that lowering a few arguments might cause the output of $\mathcal{H}()$ to decrease as well, e.g. $\mathcal{H}(1, 1, 2, 2) = 2$, whereas $\mathcal{H}(1, 1, 1, 2) = 1$.

Optimization-IV (Reducing redundant \mathcal{H} computations and LCCSAT checks): We use local lower-bound LB(v) on core-numbers (Lemma 1) to reduce the number of h-index computations and core corrections. At some iteration n, if $\hat{h}_v^{(n)}$ is equal to LB(v), we can ensure that core-number c(v) = LB(v). In other words, h-index for v would no longer reduce in future iterations; otherwise, $c(v) = \hat{h}_v^{(\infty)} < \hat{h}_v^{(n)} = LB(v)$, which is a contradiction. Hence, it must be that $c(v) = \hat{h}^{(\infty)} = \hat{h}^{(n)}$. We need not compute the h-index and core corrections for node v at future iterations.

Note that Liu et al. [45] determine the redundancy of \mathcal{H} computation for node v based on the convergence of both $h_v^{(n)}$ and $h_u^{(n)}$ of neighbors $u \in N(v)$. This does not work for our problem, because as we have shown in Example 4 that even if a node and its neighbors' h-indices have converged, node v may still need core correction.

Parallelization of Local-core. We propose a parallel implementation of the local algorithm, **Local-core(P)** following the shared-memory, data parallel programming paradigm. The algorithm partitions the nodes into *T* partitions, where *T* is the number of threads. Each thread is responsible for computing core-numbers of nodes in

Table 1: Dataset statistics: |V| #nodes, |E| #hyperedges, d(v) (mean) degree of a node, |e| (mean) cardinality of a hyperedges, |N(v)| (mean) #neighbors per node

	hypergraph	V	E	d(v)	e	N(v)
Syn.	bin4U	500	12424	99.4±8.5	4±0	225.3±15.5
	bin3U	500	16590	99.5±8	3±0	164.1±11.6
	pref3U	125329	250000	5.9±915.9	3±0	4.5±412.4
Real	enron	4423	5734	6.8±32	5.2±5	25.3±44
	contact	242	12704	127±55.2	2.4±0.5	68.7±26.6
	congress	1718	83105	426.2±475.8	8.8±6.8	494.7±248.6
	dblp	1836596	2170260	4±11.6	3.4±1.8	9±21.4
	aminer	27850748	17120546	2.3±5	3.7±2.6	8.4±24.1

its own partition. To improve load-balancing, we adopt the longest-processing-time-first scheduling approach [30] such that the aggregated number of neighbors of nodes in different threads are roughly the same. **Local-core(P)** has three core differences compared to its sequential counterpart **Local-core(OPT)** (Algorithm 5).

First, at iteration 0 (Line 3), every thread initializes hypergraph hindices for its allocated nodes asynchronously in parallel. Note that concurrent computation of |N(v)| and |N(u)| for nodes allocated to different threads requires concurrent reads to the CSR representation. Since the CSR representation does not change across queries, both queries will produce the same result as their sequential counterparts. **Second**, at subsequent iterations (n > 0), every thread computes $h_v^{(n)}$ and corrected value $\hat{h}^{(v)}$ for its allocated nodes asynchronously in parallel (Lines 6-11). At some iteration n, the computation of c[v]requires c[u] of its neighbor u. The same goes for Core-correction procedure. One may wonder if the parallel algorithm outputs wrong core-number, since due to asynchronous processing, the computation order of c[v] and c[u] is not fixed any more. Interestingly, our algorithm still terminates with the correct output because none of the theorems in § 3.4 rely on any particular computation-order of nodes. The computation-order only affects the number of iterations required for convergence, not the converged value. Finally, Localcore(P) does not use optimization-IV, as we empirically find that Optimization-IV does not improve the execution time significantly.

5 EMPIRICAL EVALUATION

We empirically evaluate the performance of our algorithms on four synthetic and five real-world datasets (Table 1). We implement our algorithms in GNU C++11 and OpenMP API version 3.1. All experiments are conducted on a server with 80 core Intel(R) Xeon(R) 2.8GHz CPU and 128GB RAM. Our code and datasets are at [4].

Datasets. Among synthetic hypergraphs, bin4U and bin3U are 4-uniform and 3-uniform hypergraphs, respectively, generated using state-of-the-art hypergraph configuration model [2]. pref3U is a 3-uniform hypergraph generated using the hypergraph preferential-attachment model [5] with parameter p=0.5, where p is the probability of a new node being preferentially attached to existing nodes in the hypergraph. The node degrees in pref3U approximately follows a power-law distribution with exponent $\beta=2.2$. Among real-world hypergraphs, enron is a hypergraph of emails, where each email correspondence is a hyperedge and users are nodes [11]. We derive the contact (in a school) dataset from a graph where each maximal clique is viewed as a hyperedge and individuals are nodes [11]. In the congress dataset, nodes are congress-persons and each hyperedge comprises of sponsors and co-sponsors (supporters) of a legislative

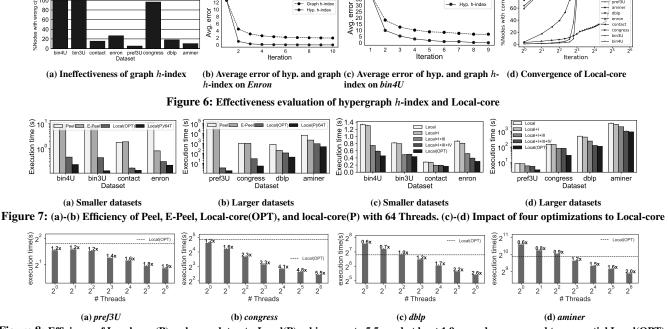


Figure 8: Efficiency of Local-core(P) on larger datasets. Local(P) achieves up to 5.5x and at least 1.9x speedup compared to sequential Local(OPT).

bill put forth in both the House of representatives and the senate [11]. In *dblp*, nodes are authors and a hyperedge consists of authors in a publication recorded on DBLP [11]. Similarly, *aminer* consists of authors and publications recorded on Aminer [58].

5.1 Effectiveness of Local-core Algorithm

Exp-1: Novelty & importance of hypergraph h-index. We demonstrate the novelty of the proposed hypergraph h-index (Definition 4) by showing that a direct adaptation of graph h-index (Definition 2) without any core correction, that is, running the local algorithm from [46, 50] may produce incorrect hypergraph core-numbers. Figure 6(a) depicts that a local algorithm that only considers graph h-index without adopting our novel Core-correction (§ 3.3) generates incorrect core-numbers for at least 90% nodes on bin4U, bin3U, and congress. On contact, enron, pref3U, dblp, and aminer, corenumbers for at least 15%, 26%, 5%, 17%, and 10% nodes are incorrect, respectively. As nodes in bin4U, bin3U, and congress have relatively higher mean(|N(v)|) (Table 1), there are more correlated neighbors in these datasets. Incorrect h-indices of correlated neighbors have a domino-effect: A few nodes with wrong h-values, unless corrected, may cause all their neighbors to have wrong h-values, which may in turn cause the neighbors' neighbors to have wrong h-values, and so on. Unless all such correlated nodes are corrected, almost all nodes eventually end up with wrong core-numbers.

We also compare the average error in the core-number estimates at each iteration by graph h-index and our hypergraph h-index. Here, avg. error at iteration $n = \sum_{u \in V} (h^{(n)}(u) - core(u))/|V|$. Figures 6(b)-(c) show avg. errors incurred at the end of each iteration on enron and bin4U. At initialization, both indices have the same error on a specific dataset. For both indices, avg. error at a given

iteration is less than or equal to that in the previous iteration. However at higher iterations, hypergraph h-index incurs less avg. error compared to graph h-index. At termination, although hypergraph h-index produces correct core-numbers, graph h-index has non-zero avg. error. These results suggest that hypergraph h-index estimates core-numbers more accurately than graph h-index at intermediate iterations. We notice similar trends in other datasets, however for the same iteration number, graph h-index produces higher avg. error on bin4U than that on enron. This is due to more number of correlated neighbors in bin4U than that in enron as stated earlier.

Exp-2: Convergence of Local-core. Figure 6(d) shows that as the number of iteration increases in our **Local-core** algorithm (§ 3.3), the percentage of nodes with correctly converged core-numbers increases. The number of iterations for convergence depend on the hypergraph structure and on the computation ordering of nodes. As pref3U, dblp, and enron has more nodes with fewer correlated neighbors, more nodes achieve correct core-numbers after the first iteration. On pref3U at least 98% nodes already converge by iteration 2. This observation also suggests that one can terminate the proposed **Local-core** algorithm early at the expense of a fraction of incorrect results for the pref3U hypergraph. Notice that even though contact has only 3% nodes with correct core-numbers at the first iteration, its convergence rate is steeper than that of bin3U and bin4U, since contact has a lower mean(|N(v)|), and thus lesser correlated neighbors compared to that for bin3U and bin4U.

5.2 Efficiency Evaluation

Figures 7(a)-(b) show the execution times of **Peel**, **E-Peel**, optimized sequential **Local-core** algorithm **Local(OPT)**, and its parallel variant **Local(P)**. We find that (1) **E-Peel** is more efficient than **Peel** on

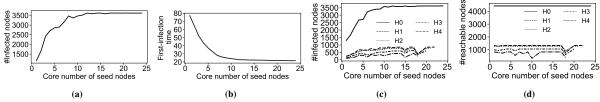


Figure 9: (a) The number of infected nodes increases as the core-number of the seed node increases. (b) The infection time of nodes decreases for nodes with higher core-numbers indicating innermost cores are infected earlier. (c) The impact of deleting the innermost-core for disrupting diffusion. H0 is the input hypergraph H, H1 is constructed by deleting nodes in the innermost-core of H0, H2 is constructed by deleting nodes in the innermost-core of H1, and so on. The number of infected nodes generally decreases after each deletion of the innermost-core. (d) The reason behind such intervention to be effective is that the average number of nodes reachable from the seed node decreases due to deletion of the innermost-core (Enron dataset.).

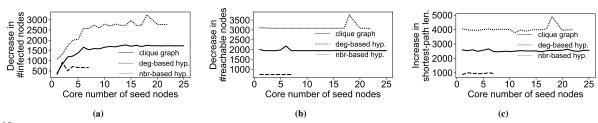


Figure 10: Comparison between different decomposition approaches on *enron*: Different decomposition methods assign different core-numbers to the same node. Thus, not all methods have the same core-number range and the curves representing different methods have different spans along X-axis. (a) Decrease in the number of infected population due to deletion of the innermost-core from different decomposition approaches. (b) Deleting an innermost neighborhood-based core causes the most disruption in reachability among nodes. (c) Nodes becomes either unreachable (∞ is replaced by |E|=5 734 to compute the avg. shortest path lengths from seed nodes) or the cost of reaching them from seed nodes after deleting the innermost-core becomes higher.

majority datasets. (2) Local(OPT) is more efficient than Peel and E-Peel on all datasets. (3) Local(P) is more efficient than Peel, E-Peel, and Local(OPT) on all datasets.

Exp-3: Efficiency of E-Peel. As stated in § 3.2, the speedup of **E-Peel** over **Peel** is related to α , which is the ratio of the #|N(u)|queries made by **E-Peel** to that of **Peel**. In Figures 7(a)-(b), **E-Peel** achieves the highest speedup on enron because $\alpha = 0.38$ is the smallest for this dataset. We also find that $\alpha \cong 1$ on *congress*, bin3U, and contact, thus E-Peel gains almost no speedup on these datasets. Exp-4: Efficiency and impact of optimizations to Local-core. We next analyze the efficiency of the proposed optimizations (§ 4) with respect to Local-core without any optimization. Local+I incorporates optimization-I to Local-core, Local+I+III incorporates optimization-III on top of Local+I, Local+I+III+IV incorporates optimization-III on top of Local+I+III. Finally, Local(OPT) incorporates all four optimizations. Figures 7(c)-(d) show the execution times of Local-core, Local+I, Local+I+III, Local+I+III+IV, and Local(OPT) on all datasets. We observe that adding each optimization generally reduces the execution time. The impact of adding the same optimization on different datasets is different in general. For instance, on enron the speedup of Local+I+III+IV w.r.t Local+I+III is 1.5x, whereas on congress the speedup is 1x (no speedup). This is because on enron, a large number of nodes (90%) have their corenumbers equal to their respective local lower-bounds (Lemma 1). However, on congress, only 0.05% nodes have core-numbers equal to their local lower-bounds. As a result, more redundant \mathcal{H} computations are saved on enron compared to that in congress.

Exp-5: Impact of parallelization. We test the parallelization performance of **Local(P)** by varying the number of threads from 1 to 64 (Figure 8). Adding more threads reduces the per-thread workload in a single iteration. As a result, individual iterations takes less time culminating in the reduction of overall execution time. For example,

in Figure 8(b), **Local(P)** is **5.5x** faster on *congress* when the number of threads increases from 1 to 64. However in Figure 8(a), **Local(P)** achieves only up to **1.9x** speedup. The limited speedup is due to two reasons: 1) As shown in Exp-2, 98% nodes converge by 2 iterations in *pref3U*. In the remaining iterations, only 2% nodes are processed in parallel. This limits the parallelism because many threads remain idle. 2) *pref3U* has a skewed neighborhood-size distribution as the standard deviation (412.4 in Table 1) is the largest among all datasets. Thus, there are a few non-converged nodes with significantly large numbers of neighbors. The thread handling such a node will take longer than the rest, causing the overall execution time to increase.

6 APPLICATIONS AND CASE STUDIES

6.1 Influence Spreading and Intervention

We empirically study the significance of nodes in the innermost-core derived from neighborhood-based core decomposition. We consider the *SIR* diffusion process [40]: Initially, all nodes except one—called a *seed*—are at the *susceptible* state. The seed node is initially at the *infectious* state. At each time step, each infected node infects its susceptible neighbors with probability β and then enters into *immunized* state. Once a node is immunized, it is never re-infected.

Innermost-core contains influential spreaders. We randomly select a seed node from each core-number in *enron* and run the SIR model (β =0.3). We compute the number of infected nodes after 100 time-steps. We repeat this process 1000 times and report the average number of infected nodes in Figure 9(a). We observe that inner-core nodes infect larger population compared to outer-core nodes.

Innermost-core contracts diffusion early. We select a seed node from the hypergraph uniformly at random, and run the *SIR* model for 100 time-steps. For each seed node, we record the time-step at which other nodes are infected. Repeating 1000 times with different seeds,

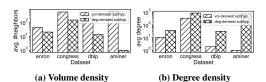


Figure 11: Comparison between different subhypergraphs based on average #neighbors (left) and average degree (right) per node

we report the average infection times. Figure 9(b) indicates that nodes in inner-cores are infected earlier than those in outer-cores.

Innermost-core deletion for maximum intervention in spreading. We next devise an intervention strategy to disrupt diffusion, which has significance in mitigating the spread of contagions (in epidemiology), limiting the spread of misinformation, or blocking competitive campaigns (in marketing). We delete nodes and hyperedges in the innermost-core, and measure the number of infected population before and after deletion. Figure 9(c) shows the effectiveness of our intervention strategy over *enron*, where H0 is the input hypergraph H, H1 is constructed by deleting nodes in the innermost-core of H0, H2 is constructed by deleting nodes in the innermost-core of H1, and so on. We select a seed node from the remaining hypergraph uniformly at random. We find that the number of infected population generally decreases after each deletion of the innermost-core, irrespective of the origin of seed nodes. Moreover, deleting the innermost-core also reduces the number of nodes reachable from seed nodes, making our intervention quite effective (Figure 9(d)).

Comparative analysis of intervention strategies. We compare the effectiveness of different core decomposition approaches on hypergraphs, while applying the same intervention strategy. For that, we take the difference in the number of infected population, average number of reachable nodes from the seed, and average length of shortest paths from the seed between H0 and H1, separately for all three decomposition approaches: neighborhood-based (this work), degree-based [51, 56], and clique-graph based (§ 2.2).

In Figure 10(a), we observe that the neighborhood-based intervention results in the largest decrease in infected population compared to other two core decomposition approaches, thereby showing superior effectiveness of our decomposition-based intervention. To explain why our intervention is more effective than others, Figures 10(b)-(c) show the highest decrease in the average number of reachable nodes and the highest increase in the average length of shortest paths in our decomposition-based intervention.

6.2 Densest SubHypergraph Finding

The densest subgraph can be defined as a subgraph with the maximum average node-degree among all subgraphs of a given graph [15, 22, 29], which may correspond to communities [19], filter bubbles and echo chambers [41] in social networks, and brain regions responding to stimuli [43] or diseases [65]. Following the same principal, we define a new notion of densest subhypergraph, called the *volume-densest subhypergraph*, based on the number of neighbors of nodes in a hypergraph. The **volume-density** $\rho^N[S]$ of a subset $S \subseteq V$ of nodes in a hypergraph H = (V, E) is defined as the ratio of the summation of neighborhood sizes of all nodes $u \in S$ in the induced subhypergraph H[S] to the number of nodes in H[S].

$$\rho^{N}[S] = \frac{\sum_{u \in S} |N_{S}(u)|}{|S|} \tag{8}$$

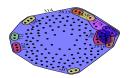


Figure 12: Volume-densest subhypergraph of human protein complex

The **volume-densest subhypergraph** is a subhypergraph which has the largest volume-density among all subhypergraphs.

Approximation algorithm. Inspired by Charikar [15], our approach to find the (approximate) volume-densest subhypergraph follows the peeling paradigm: In each round, we remove the node with the smallest number of neighbors in the current subhypergraph. In particular, we sort the nodes in ascending order of their neighborhood-based core-numbers, obtained from any neighborhood-based core-decomposition algorithm (e.g., Peel, E-Peel, Local(OPT), or Local(P)). We peel nodes in that order. Among nodes with the same core-number, the one with the smallest number of neighbors in the current subhypergraph is selected earlier for peeling. We finally return the subhypergraph that achieves the largest volume-density.

THEOREM 5. Our volume-densest subhypergraph finding algorithm returns $(d_{pair}(d_{card}-2)+2)$ -approximate densest subhypergraph if the maximum cardinality of any hyperedge is d_{card} and the maximum number of hyperedges between any pair of nodes is d_{pair} in the input hypergraph.

The proof is given in our extended version [3]. Notice that $d_{pair} = 2$ if the hypergraph is a graph and our result gives 2-approximation guarantee for the densest subgraph discovery [15].

6.2.1 Effectiveness of volume-densest subhypergraphs. We compute volume-densest subhypergraphs from our datasets using our approximation algorithm, and in those subhypergraphs we compute the avg. number of neighbors (neighborhood-cohesion measure) and the avg. degree (degree-cohesion measure) per node. As baseline for comparison, we compute those two measures on the degree-densest subhypergraphs of the same hypergraphs extracted using [35].

In Figure 11(a), we find that the nodes in the volume-densest subhypergraph have more neighbors (on average) than that in the degree-densest subhypergraph. In Figure 11(b), we observe that the nodes in the degree-densest subhypergraph have a higher degree (on average) than that in the volume-densest subhypergraph. These observations suggest that volume-densest subhypergraph is more effective than degree-densest subhypergraph in capturing neighborhood-cohesive regions. In contrast, degree-densest subhypergraph is more effective in capturing degree-cohesive regions. Neighborhood-cohesiveness is important in many applications as follows.

6.2.2 Case Study 1: Biology. We analyze a real-world hypergraph of manually-annotated human protein complexes collected from the CORUM database [28]. We consider each protein complex as a hyperedge consisting of proteins as nodes. There are 2611 hyperedges and 3622 nodes in the Human protein complex hypergraph. The volume-densest subhypergraph shown in Figure 12 has several interesting characteristics. First, the complexes in the subhypergraph are correlated as they participate in two fundamental biological processes: RNA metabolism and RNA localisation (e.g., some of them are listed in Table 2). Second, the largest hyperedge (314) is the Spliceosome complex, responsible for RNA splicing, which assists

Table 3: The top-5 highest degree nodes in volumedensest and degree-densest subhypergraphs. High-ranking executives of *enron* who make key decisions are captured in both subhypergraphs.

Table 2: Functions of some complexes in the volume-densest subhypergraph of human protein complex

Hyper- edge id	Complex name	Function
314	Spliceosome	RNA metabolic process
708	TREX	RNA Localization
712	THO	RNA Localization
713	CBC	RNA Localization
2835	TRA2B1-	RNA Metabolic
	SRSF9-	process
	SRSF6	
2836	SRSF9-	RNA Metabolic
	SRSF6	process
2916	hTREX84	RNA Metabolic
		process

	Top-5 highest degree nodes	Designation	Degree in sub hyp.
	Kenneth Lay	CEO	202
Degree	Greg Whalley	President	118
densest	Mark Koenig	Head (Investor	107
subhyp.		Relations)	
	Jeffrey McMahon	Chief Financial	88
		Officer	
	Mark A. Frevert	Chairman and	86
		CEO (EWS)	
	Kenneth Lay	CEO	122
Volume	Greg Whalley	President	28
densest	Jeffery Skilling	CEO	27
subhyp.	Mark A. Frevert	Chairman	22
		and CEO (EWS)	
	Mark Koenig	Head (Investor	20
		Relations)	

Table 5: Subject and intent of the top-3 emails with the highest number of participants in the volume-densest subhypergraph

Email subject	Intent of the email	
	Seeking applicants	
Haland Comme OF and a Tital	for Board of directors position at	
Updated Cougars@Enron email list	Cougars@Enron (U Housten	
	alumni group at Enron)	
A	Announcing about	
Associate/ analyst program	talent-seeking program of Enron.	
Analyst & associate program -	Total and a second and all and a second	
e-speak invitation from Billy Lemmons	Invitation to attend an online seminer	

in cell-evolution process and in the making of new and improved proteins in human body. Two subsets of Splicesome (2835 and 2836) are responsible for regulating mRNA splicing, which are known to be affected by a genetic disease called TAU-mutation causing frontotemporal dementia. **Finally**, the TREX complex (708) is responsible for transporting mRNA from the nucleus to the cytoplasm. One of its subsets, hTREX84 (2916) is found to be highly correlated with ovarian and breast cancers [32].

6.2.3 Case study II: Email communication. We extract all emails involving Kenneth Lay, who was the founder, chief executive officer, and chairman of Enron [64]. The ego-hypergraph of such a keyperson, having 4718 nodes (person) and 1190 hyperedges (emails), can provide insights and difference between the volume-densest and degree-densest subhypergraphs. The degree-densest subhypergraph has 166 nodes and 202 hyperedges, whereas the volumedensest subhypergraph has 1949 nodes and 122 hyperedges. We analyze the degree and neighborhood-sizes of the top-5 nodes in these two subhypergraphs in Tables 3 and 4. In Table 3, we find that both degree-densest and volume-densest subhypergraphs contain high-ranking key-personnel in Enron. Such personnel (nodes) participate in many emails (hyperedges) in the extracted subhypergraph. However, in Table 4, we notice that ordinary employees are communicated the most in emails (i.e., they are nodes with many neighbors), and such employees can only be extracted by analyzing the volume-densest subhypergraph. We further investigate the reason why ordinary employees have more neighbors in the volume-densest

Table 4: The top-5 highest neighborhood-size nodes in volumedensest and degree-densest subhypergraphs. In the degree-densest subhypergraph, the top-5 highest neighborhood-size nodes are quite similar to those in the top-5 highest-degree nodes (highranking executives). However, the top-5 highest neighborhoodsize nodes in the volume-densest subhypergraph are ordinary employees.

	Top-5 highest neighborhood-size nodes	Designation	#nbrs in subhyp.
	Kenneth Lay	CEO	165
Degree	Greg Whalley	President	158
densest	Mark Koenig	Head(Investor	153
subhyp.		Relations)	
	Jeffrey McMahon	Chief Financial	152
		Officer	
	John Sherriff	President and	151
		CEO (Enron Europe)	
	Gregory Martin	Analyst	1948
Volume	Dustin Collins	Associate (Enron	1948
densest		Global Commodities)	1948
subhyp.	Andrea Richards	Ordinary Employee	1948
	Sladana-anna Kulic	Ordinary Employee	1948
	Maureen Mcvicker	Assistant	1948

subhypergraph. We extract hyperedges (emails) where the top-5 highest neighborhood-size nodes were involved. We found that many such emails were about internal announcements, meetings/seminar invitation, and employee social gatherings (Table 5).

7 RELATED WORK

Recently, there has been a growing interest on hypergraphs data management [23, 38, 42, 54, 57, 63]. Since the focus of this paper is on core decomposition, we urge readers with a much broader interest to refer to recent surveys [10, 20, 60] for a general exposition.

Core decomposition in hypergraphs. Although core decomposition on graphs have been studied for decades [47], there are relatively few works on hypergraph core decomposition. Ramadan et al. [51] propose a peeling algorithm to find the maximal-degree-based *k*-core of a hypergraph. [37, 54] discuss parallel implementations of degree-based hypergraph core computation based on peeling approach. Sun et al. [56] propose a fully dynamic approximation algorithm that maintains approximate degree-based core-numbers of an unweighted hypergraph. Unlike ours, none of these works explore neighborhood-based hypergraph core decomposition, which is different from degree-based hypergraph core computation (§1), nor they consider algorithmic approaches other than peeling.

Core decomposition in graphs. The linear-time peeling algorithm for graph core decomposition was given by Batagelj and Zaveršnik [9]. Core decomposition has also been studied in disk-based [16, 39], distributed [50], parallel [18], and streaming [52] settings, and for varieties of graphs, e.g., weighted [26], directed [44], temporal [26], uncertain [13], and multi-layer [25] networks. Higher-order cores in a graph (e.g., (k, h)-core [14, 45], triangle k-core [69], k-clique-core, and pattern-core [22, 61]) and more complex cores in an attributed network (e.g., meta-path-based core [21] and (k, r)-core [68]) have been proposed. In §2.2, we reasoned that existing approaches for graph core decomposition cannot be easily adapted for neighborhood-based hypergraph core decomposition. We also depicted that the local approach [46, 50], an efficient method for

graph core decomposition, produces incorrect core-numbers for neighborhood-based hypergraph core decomposition (§3.3, §5).

8 CONCLUSIONS

In this paper, we introduced the notion of neighborhood-cohesive core decomposition of hypergraphs. We proved that our decomposition has desirable properties such as Uniqueness and Corecontainment. We then proposed three algorithms – Peel, E-Peel, and novel Local-core for hypergraph core decomposition. We also adopted four hypergraph-specific optimizations to Local-core and its shared-memory parallel implementation. Empirical evaluation on synthetic and real-world hypergraphs depicted that the novel Localcore with optimizations and parallel implementation is the most efficient among all proposed algorithms. In diffusion applications, the proposed decomposition is more effective than degree-based decomposition in intervening diffusion. In the densest subhypergraph finding application, our decomposition helps extract the volumedensest subhypergraph with an approximation guarantee. Our novel volume-densest subhypergraphs are more effective in maintaining neighborhood cohesiveness than existing degree-densest subhypergraphs. Two case studies illustrated that volume-densest subhypergraphs capture functionally significant protein-complexes in biology domains and important emails in organizational communications.

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