Fast, Flexible, and Exact Minimum Flow Decompositions via ILP

Fernando H.C. Dias¹, Lucia Williams², Brendan Mumey², Alexandru I. Tomescu¹

 1 Department of Computer Science, University of Helsinki, Finland 2 School of Computing, Montana State University, Bozeman, MT, USA

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Background and Motivation



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Flow decomposition (FD), the problem of decomposing a **network flow** into a set of source-to-sink **paths** and associated **weights** that perfectly explain the flow values on the edges, is a classical and well-studied concept in Computer Science.

Motivation



- The main bioinformatics motivation for this paper is *multiassembly* [1], reconstruct multiple genomic sequences from mixed samples using short substrings (called *reads*) generated cheaply and accurately from next-generation sequencing technology;
- One example is the reconstruction of RNA transcripts from sequencing reads, which is essential to characterize gene regulation and function, development and diseases such as cancer;
- Another is the **reconstruction of viral quasispecies** which can identify different strains of a virus in a samples.

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Preliminares



Definition (Flow network)

A tuple G=(V,E,f) is said to be a *flow network* if (V,E) is a DAG with unique source s and unique sink t, where for every edge $(u,v)\in E$ we have an associated positive integer *flow value* f_{uv} , satisfying *conservation of flow* for every $v\in V\setminus\{s,t\}$, namely:

$$\sum_{u,v)\in E} f_{uv} = \sum_{(v,w)\in E} f_{vw}.$$
 (1)

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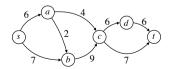
Preliminares

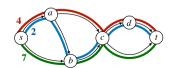


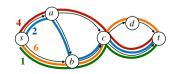
Definition (k-Flow Decomposition)

A k-flow decomposition (\mathcal{P},w) for a flow network G=(V,E,f) is a set of k s-t flow paths $\mathcal{P}=(P_1,\ldots,P_k)$ and associated weights $w=(w_1,\ldots,w_k)$, with each $w_i\in\mathbb{Z}^+$, such that for each edge $(u,v)\in E$ it holds that:

$$\sum_{\substack{i \in \{1,\dots,k\} \text{ s.t.} \\ (u,v) \in P_i}} w_i = f_{uv}. \tag{2}$$







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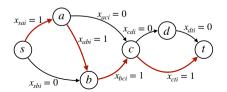
Flow Conservation



$$\sum_{(s,v)\in E} x_{svi} = 1, \qquad \forall i \in \{1,\dots,k\}, \tag{3a}$$

$$\sum_{(u,t)\in E} x_{uti} = 1, \qquad \forall i \in \{1,\dots,k\},$$
(3b)

$$\sum_{(u,v)\in E} x_{uvi} - \sum_{(v,w)\in E} x_{vwi} = 0, \quad \forall i \in \{1,\dots,k\}, \forall v \in V \setminus \{s,t\}.$$
 (3c)



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Flow Superposition



$$\sum_{i \in \{1,\dots,k\}} x_{uvi} w_i = f_{uv}, \qquad \forall (u,v) \in E.$$
 (4)

Linearized as:

$$f_{uv} = \sum_{i \in \{1, \dots, k\}} \pi_{uvi}, \qquad \forall (u, v) \in E,$$
(5a)

$$\pi_{uvi} \le \overline{w}x_{uvi}, \qquad \forall (u,v) \in E, \forall i \in \{1,\dots,k\},$$
 (5b)

$$\pi_{uvi} \le w_i, \qquad \forall (u, v) \in E, \forall i \in \{1, \dots, k\},$$
 (5c)

$$\pi_{uvi} \ge w_i - (1 - x_{uvi})\overline{w}, \qquad \forall (u, v) \in E, \forall i \in \{1, \dots, k\}.$$
 (5d)

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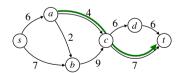
Subpath Constraints



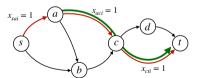
Definition (Flow decomposition with subpath constraints)

Let G=(V,E,f) be a flow network. Subpath constraints are defined to be a set of simple paths $\mathcal{R}=\{R_1,\ldots,R_\ell\}$ in G (not necessarily s-t paths). A flow decomposition (\mathcal{P},w) satisfies the subpath constraints if and only if

$$\forall R_j \in \mathcal{R}, \exists P_i \in \mathcal{P} \text{ such that } R_j \text{ is a subpath of } P_i.$$
 (6)



(a) A flow network with a single subpath constraint $R_1 = (a, c, t)$.



(b) Constraint R_1 is satisfied because for the $i^{\rm th}$ path we can set $r_{i1}=1$ so that $x_{aci}+x_{cti}\geq 2r_{i1}$ holds .

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Subpath Constraints - Formulation



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$$\forall R_j \in \mathcal{R}, \exists P_i \in \mathcal{P} \text{ such that } R_j \text{ is a subpath of } P_i.$$
 (7)

can be written:

$$\sum_{(u,v)\in R_i} x_{uvi} \ge |R_j| r_{ij}, \qquad \forall i \in \{1,\dots,k\}, \forall R_j \in \mathcal{R},$$
 (8a)

$$\sum_{i \in \{1, \dots, k\}} r_{ij} \ge 1, \qquad \forall R_j \in \mathcal{R}.$$
 (8b)

Inexact Flow



Definition (Inexact flow network)

A tuple $G=(V,E,\underline{f},\overline{f})$ is said to be an *inexact flow network* if (V,E) is a DAG with unique source s and unique sink t, where for every edge $(u,v)\in E$ we have associated two positive integer values $\underline{f_{uv}}$ and $\overline{f_{uv}}$, satisfying $\underline{f_{uv}}\in \overline{f_{uv}}$.

Given an inexact flow network $G=(V,E,\underline{f},\overline{f})$ the minimum inexact flow decomposition problem is to determine if there exists, and if so, find a minimum-size set of s-t paths $\mathcal{P}=(P_1,\ldots,P_k)$ and associated weights $w=(w_1,\ldots,w_k)$ with $w_i\in\mathbb{Z}^+$ such that for each edge $(u,v)\in E$ it holds that:

$$\underline{f_{uv}} \le \sum_{\substack{i \in \{1,\dots,k\} \text{ s.t.} \\ (u,v) \in P_i}} w_i \le \overline{f_{uv}}.$$
(9)

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 $\underline{f_{uv}} \le \sum_{uvi} \overline{f_{uv}},$ $i \in \{1,...,k\}$

 $\forall (u, v) \in E. \tag{10a}$



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Experiments Design



- Time limit of 60 seconds:
- 2 Comparison of STANDARD with Toboggan, the implementation by [2] for their exact FPT algorithm for MFD;
- **3** Comparison of *SUBPATH* with Coaster, the implementation by [3] for MFDSC, which is an exact FPT algorithm extending Toboggan.
- 4 Comparison of INEXACT with IFDSolver, which is an implementation of a heuristic algorithm for MIFD by [4];
- **6** Three different datasets composed of RNA transcripts with range of nodes between 4 and 50;

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- For the *STANDARD* formulation, we could solve all instances within **20** seconds. While the *TOBOGGAN* required at least **1** minute for instances up to 10 flow-paths. For instances with more than 10 flow-paths, it did **not** solve within the runtime limit;
- For the *SUBPATH* version, the runtime for our formulation is below **30 seconds**, while *COASTER* cannot solve most instances;
- For the *INEXACT*, the runtime of the heuristic is **faster**, but the solution is not optimal (overestimate the number of paths by 2, in average).

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Conclusions



- Fast: all instances were solved in under 20 seconds while state-of-the-start method requires a few minutes;
- Flexible: capable to be easily adjusted to incorporate new behaviour (as done
 with subpath constraints and inexact constraint) without compromising
 performance;
- Future Work: Cycles, improvement in inexact flow with Robust Optimization;
- https://github.com/algbio/MFD-ILP

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Acknowlegement



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