The maximum infection time in the geodesic and monophonic convexities $\stackrel{\bigstar}{\Rightarrow}$

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Abstract

Recent papers investigated the maximum infection time $t_{P3}(G)$ of the P_3 convexity (also called maximum time of 2-neighbour bootstrap percolation) and the maximum infection time $t_{mo}(G)$ of the monophonic convexity. In 2014, it was proved that, for bipartite graphs, deciding whether $t_{P3}(G) \ge k$ is polynomial time solvable for $k \le 4$, but is NP-complete for $k \ge 5$ [23]. In 2015, it was proved that deciding whether $t_{mo}(G) \ge 2$ is NP-Complete even for bipartite graphs [12]. In this paper, we investigate the maximum infection time $t_{gd}(G)$ of the geodesic convexity. We prove that deciding whether $t_{gd}(G) \ge k$ is polynomial time solvable for k = 1, but is NP-complete for $k \ge 2$ even for bipartite graphs. We also present an $O(n^3m)$ -time algorithm to determine $t_{gd}(G)$ and $t_{mo}(G)$ in distance-hereditary graphs. For this, we characterize all minimal hull sets of a general graph in the monophonic convexity. Moreover, we improve the complexity of the fastest known algorithm for finding a minimum hull set of a general graph in the monophonic convexity.

Keywords: Geodesic convexity, maximum infection time, monophonic hull number, monophonic convexity

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1. Introduction

A family \mathcal{C} of subsets of a finite set V is a convexity on V if \emptyset and V belong to \mathcal{C} and \mathcal{C} is closed under intersections. A member of \mathcal{C} is said to be a convex set of \mathcal{C} or simply a \mathcal{C} -convex set. When there is no risk of confusion, we omit \mathcal{C} and write only convex set. When V is the vertex set of a graph G, a standard way to define a convexity \mathcal{C} on V is by fixing a family \mathcal{P} of paths of G and letting a subset S of V be \mathcal{C} -convex if and only if for every path P in \mathcal{P} whose endpoints belong to S we have that all vertices of P also belong to S. In that case we have a graph convexity. The most studied graph convexities are the geodesic convexity [17, 19], the monophonic convexity [14, 16, 18] and the P_3 convexity [5], where \mathcal{P} is, respectively, the family of all shortest paths, of all induced paths, and of all paths of order three of the graph. A rich source on abstract convexities is a book by van de Vel [26].

The *C*-convex hull of *S* is the smallest *C*-convex set, denoted $H_{\mathcal{C}}(S)$, containing *S*. We say that *S* is a *C*-hull set if $H_{\mathcal{C}}(S) = V(G)$. The *C*-hull number is the cardinality of the minimum *C*-hull set and is denoted by $h_{\mathcal{C}}(G)$. The *C*-interval of *S*, denoted $I_{\mathcal{C}}(S)$, consists of *S* and all vertices lying in some path of \mathcal{P} that has endpoints in *S*. Observe that, since *G* is finite, the *C*convex hull of *S* can be obtained by starting with *S* and repeatedly applying the *C*-interval function until we obtain a *C*-convex set.

We may also see the above process, used to obtain the \mathcal{C} -convex hull of S, as an infection that starts at the set S and spreads to other vertices through the paths (of \mathcal{P}) that connect two infected vertices. Here, we are interested in the maximum amount of time needed to infect all vertices starting with a \mathcal{C} -hull set, where one unit of time corresponds to applying the interval function once. More precisely, let $I^0_{\mathcal{C}}(S) = S$ and $I^k_{\mathcal{C}}(S) = I_{\mathcal{C}}(I^{k-1}_{\mathcal{C}}(S))$ for $k \geq 1$. We say that a hull set S takes time k to infect G if $I^k_{\mathcal{C}}(S) = V(G)$ but $I^{k-1}_{\mathcal{C}}(S) \neq V(G)$ (when S = V(G) we say it takes time 0 to infect G). The *infection time of G relative to* \mathcal{C} , called $t_{\mathcal{C}}(G)$, is the maximum k such that there is a \mathcal{C} -hull set S which takes time k to infect G. We considered the decision version of this problem.

MAX INFECTION TIME ON CONVEXITY CInput: A graph G and an integer k. Question: Is $t_{\mathcal{C}}(G) \geq k$? There is vast literature about "infection problems", also studied under the names "dissemination", "diffusion" or "conversion" [2, 3, 21]. However, the way such infections spread varies considerably. For example, another common model is r-neighbour bootstrap percolation in which a vertex becomes infected if it has at least r infected neighbors, a model introduced by Chalupa, Leath and Reich [10] that has found many applications in physics and computer science [15]. There are also applications in clustering phenomena, sandpiles [20], and many other areas of statistical physics, as well as in neural networks [1].

The question about the maximum infection time was originally posed by Bollobás for the 2-neighbour bootstrap percolation in the square grid, and solved by Benevides and Przykucki [8]. We remark that in general, counterintuitively, a hull set S that infects G in time $t_{\mathcal{C}}(G)$ may need to have more than $h_{\mathcal{C}}(G)$ elements. That is, a set that infect G at the maximum possible time does not have to be of minimum order (although it has to be minimal). This follows from the main results in [7] and [8], even when G is a grid and \mathcal{C} is the P_3 -convexity.

Note that the 2-neighbour bootstrap percolation model coincides with the infection problem for the P_3 convexity. The algorithm complexity of MAX INFECTION TIME ON THE P_3 CONVEXITY was considered in [6] where it was shown that it is NP-complete for general graphs and any fixed $k \ge 4$, and it was given polynomial algorithms for planar graphs, trees, chordal graphs, and for $k \le 2$. In 2014, Marcilon et al. obtained polynomial time algorithms for $k \le 3$ in general graphs and $k \le 4$ in bipartite graphs, and proved that it is NP-complete for $k \ge 5$ in bipartite graphs [23]. In 2015, Marcilon and Sampaio proved that it is NP-complete even in grid graphs with maximum degree 3 [24]. The question about solid grid graphs with maximum degree 4 is still open.

Regarding the monophonic convexity, it was proved in 2015 that the maximum infection time problem is NP-complete for $k \geq 2$ in bipartite graphs [12]. In 2010, Dourado et al. obtained an $O(n^3m)$ -time algorithm to compute a minimum hull set for the monophonic convexity [14].

In this work, we consider the MAX INFECTION TIME on the geodesic and on the monophonic convexities. This problem is, therefore, at the intersection of two large branches: "convexity problems" and "infection problems". In Section 3, we prove that the MAX INFECTION TIME ON GEODESIC CON-VEXITY is NP-complete even if the input graph is bipartite and $k \geq 2$ is fixed. In Section 4, we obtain an important characterization of all minimal hull sets in the monophonic convexity (see Theorem 4.9). We also consider the problem of determining the hull number of G in the monophonic convexity. As a matter of fact, we observe that the clique decomposition presented in [22] can be used to improve the time complexity of the fastest known algorithm [14] for this problem from $O(n^3m)$ -time to O(nm)-time. These results are useful for a polynomial time algorithm, which is given in Section 5, for computing the maximum infection time on distance-hereditary graphs, a class of graphs where the geodesic and monophonic convexities coincide. In the next section, we present some useful definitions and examples.

2. Preliminaries and notation

We consider only finite, simple, and undirected graphs. For a vertex u of a graph G, denote by N(u) the set of neighbors of u, and for $S \subseteq V(G)$, denote by N(S) the set $\{v : v \in N(u) \setminus S \text{ and } u \in S\}$. The subgraph of G induced by S is denoted by G[S]. For two vertices $u, v \in V(G)$, the distance of u to v, denoted d(u, v), is the number of edges in a shortest path between u and v. If every two vertices of S are adjacent, then S is called a *clique of* G; and if every two vertices are not adjacent, then S is called an *independent* set of G. A graph is *bipartite* if its vertex set can be partitioned into two independent sets. A vertex v is called simplicial if N(v) is a clique. For any positive integer n, we define [n] to be the set $\{t : t \text{ is integer and } 1 \leq t \leq n\}$.

When there is no ambiguity about which graph convexity we are using, we will drop the symbol of the convexity. For example, we will use H(S) to denote the convex hull of S in the considered underling convexity. On the other hand, sometimes only the type (e.g., geodetic, monophonic or P_3) of convexity is clear but not the convexity itself. For example, it may be clear that we are working on the geodesic convexity, but the family C of convex sets still depends on the underling graph G: a set S may be convex in a graph G but not in a subgraph G' of G, even if $S \subseteq V(G')$, and vice-versa. Therefore, in some places, it will be useful to have a subscript to identify which graph was used to define the convexity.

As an example of the infection process, we present a family of (distancehereditary) graphs H_k , for $k \ge 1$, where $t(H_k) = k$ in either the geodesic or monophonic convexity. This family will be useful in the proof that the MAX INFECTION TIME ON GEODESIC CONVEXITY problem is NP-complete for bipartite graphs (Theorem 3.1). The graph H_k is defined recursively as



Figure 1: The Graph H_9 , which is distance-hereditary. A value inside the vertex is the time which the vertex becomes infected starting from the hull set v_1, v_3 .

follows (see also Figure 1): H_1 is a cycle of length four, where $V(H_1) = \{v_1, v_2, v_3, v_4\}$ and $E(H_1) = \{v_1v_2, v_2v_3, v_3v_4, v_4v_1\}$, and for $k \ge 2$ we set

$$V(H_k) = V(H_{k-1}) \cup \{v_{k+3}\} \text{ and}$$
(1)

$$E(H_k) = E(H_{k-1}) \cup \{v_{k+3}v_k, v_{k+3}v_{k+2}\}.$$
(2)

For our NP-completeness result, we will make a polynomial reduction from the following well known NP-complete problem [11].

3Sat

Input: A formula \mathcal{F} in the CNF such that each clause has size 3. Question: Is there a truth assignment satisfying all clauses of \mathcal{F} ?

3. Geodetic infection time on Bipartite graphs

Throughout this section, we only consider the geodesic convexity. Our goal is to prove the following theorem.

Theorem 3.1. MAX INFECTION TIME ON GEODESIC CONVEXITY is NPcomplete even if the input graph is known to be bipartite and $k \ge 2$ is fixed. *Proof.* Let G be a graph, S be any subset of its vertices, and $k \ge 1$ be an integer. Since the set I(S), result of the interval function applied to S, can be obtained in polynomial time on the size of G [13], for a given k one can check in polynomial time if $I^k(S) = V(G)$. Therefore this problem belongs to NP.

For the proof of the hardness, we first treat the case where k = 2. We shall do a reduction from 3SAT. Consider a boolean formula in the conjunctive normal form with *m* clauses, say $\mathcal{F} = \{C_1, \ldots, C_m\}$, on the set of variables $\{x_1, \ldots, x_n\}$.

For each clause C_i of \mathcal{F} we build a clause gadget, the bipartite graph depicted in Figure 2, whose vertex set is $\{u_i, t_i\} \cup \{v_{i,l}, w_{i,l}, x_{i,l}, y_{i,l} : l \in [3]\}$, and edge set is $\{u_i t_i\} \cup \{u_i v_{i,l}, t_i w_{i,l}, v_{i,l} w_{i,l}, w_{i,l} x_{i,l}, x_{i,l} y_{i,l} : l \in [3]\}$.

We construct a graph G as follows. Add each clause gadget to G and further add vertices q, r and s along with the edges $rx_{i,l}, st_i$ for every $i \in [m]$ and $l \in [3]$. Finally, for every pair of literals $\ell_{i,a} \in C_i$ and $\ell_{j,b} \in C_j$, where $i \neq j$ and $a, b \in [3]$, such that $\ell_{i,a}$ and $\ell_{j,b}$ are not the negation of each other, add vertices $o_{i,a,j,b}, p_{i,a,j,b}$ along with the edges $o_{i,a,j,b}p_{i,a,j,b}, o_{i,a,j,b}q, o_{i,a,j,b}w_{i,a},$ $o_{i,a,j,b}w_{j,b}$, to complete the construction of G. See Figure 3 for a partial construction of the graph G, when $\mathcal{F} = \{C_1, C_2, C_3\}$ where $C_1 = \{x_1, x_2, x_4\}$, $C_2 = \{\overline{x_2}, x_3, \overline{x_4}\}, C_3 = \{\overline{x_1}, x_2, \overline{x_4}\}$. Note that the gray and white vertices define a bipartition of G.

We let $W = \{w_{i,l} : \text{ for } i \in [m], l \in [3]\}, T = \{t_i : \text{ for } i \in [m]\}$. Similarly *O* is the set of vertices $o_{i,a,j,b}$ for all values of i, a, j, b for which $o_{i,a,j,b}$ is defined. Define *X*, *Y*, *V*, *U* and *P* in a analogous way. Finally for every $i \in [m]$, let $D_i = \{u_i, v_{i,1}, v_{i,2}, v_{i,3}\}$.

Now we show that there exists a truth assignment \mathcal{A} , to the variables $\{x_1, \ldots, x_n\}$, satisfying all clauses of \mathcal{F} if and only if the infection time of G is at least 2. This shall be a consequence of a few claims.

Claim 1. Every hull set of G contains at least one vertex of D_i , for every $i \in [m]$.

Proof. Fix $i \in [m]$. It suffices to show that $V(G) \setminus D_i$ is a convex set of G. Define $E_i = N(D_i) = \{t_i, w_{i,1}, w_{i,2}, w_{i,3}\}$. Observe that $V(G) \setminus D_i$ is a convex set if and only if $I(E_i) \cap D_i = \emptyset$. We distinguish two possibilities for a pair of vertices $z, z' \in E_i$. The first one is $z = t_i$ and $z' = w_{i,a}$, for some $a \in \{1, 2, 3\}$. In this case $zz' \in E(G)$ and then $I(z, z') = \{z, z'\}$. The second one is $z = w_{i,a}$ and $z' = w_{i,b}$, for some $a, b \in \{1, 2, 3\}$ with $a \neq b$. Since



Figure 2: Clause gadget.

 $t_i \in N(z) \cap N(z')$, it holds d(z, z') = 2. In fact, t_i is the unique common neighbour of z and z'. Then, $I(z, z') = \{z, z', t_i\}$.

Claim 2. Let $z \in D_i$, for some $i \in [m]$. Then $D_i \cup \{t_i\} \subset I(\{z, y_{i,1}, y_{i,2}, y_{i,3}\})$.

Proof. If $z = u_i$, it is clear that $zv_{i,a}w_{i,a}x_{i,a}y_{i,a}$ is a shortest path between $z = u_i$ and $y_{i,a}$, for any $a \in \{1, 2, 3\}$,; and that $zt_iw_{i,1}x_{i,1}y_{i,1}$ is a shortest path between $z = u_i$ and $y_{i,1}$ (in G).

Now, if $z = v_{i,a}$, for some $a \in \{1, 2, 3\}$, it is also clear that $zu_i v_{i,b} w_{i,b} x_{i,b} y_{i,b}$, and $zu_i t_i w_{i,b} x_{i,b} y_{i,b}$ are shortest paths between $z = v_{i,a}$ and $y_{i,b}$, for any $b \neq a$, with $b \in \{1, 2, 3\}$.

Claim 3. $W \cup X \cup O \cup \{q, r\} \subset I(Y \cup P)$ and $s \notin I(Y \cup P)$.

Proof. For every vertex $z \in W \cup X \cup O \cup \{q, r\}$ we show that there exists a pair of vertices $z', z'' \in Y \cup P$ such that $z \in I(\{z', z''\})$. Recall that that G is bipartite, and color its vertices with "black" or "white" as in Figure 3. Note that $Y \cup P$ is a subset of the "black" vertices. Also, for two "black" vertices z', z'', we have that d(z', z'') < 4 if and only if z' and z'' have a common neighbor.

Therefore, for any two distinct vertices $z', z'' \in Y$, we have d(z', z'') = 4. Since there is a path of length 4 between z' and z'' that pass through r, we



Figure 3: Partial construction of G for $\mathcal{F} = \{C_1, C_2, C_3\}$ where $C_1 = \{x_1, x_2, x_4\}$, $C_2 = \{\overline{x_2}, x_3, \overline{x_4}\}$, $C_3 = \{\overline{x_1}, x_2, \overline{x_4}\}$. The figure does not contain all vertices of O and P. In fact, it shows only vertices of O and P related with the literal x_4 of C_1 .

have that $r \in I(\{z', z''\})$. Similarly, for any two distinct vertices $z', z'' \in P$, we have that $q \in I(\{z', z''\})$.

Let $i \in [m]$ and $a \in \{1, 2, 3\}$. Since there exists at least one pair (j, b) for which $p_{i,a,j,b}$ is a vertex, there exists the shortest path $y_{i,a}x_{i,a}w_{i,a}o_{i,a,j,b}p_{i,a,j,b}$. Then one can easily check that $W \cup X \cup O \subset I(Y \cup P)$.

It remains to show that $s \notin I(Y \cup P)$. Note that for any $p \in P$ and $y \in Y$, we have that d(q, p) = 2, d(r, y) = 2 and d(q, y) = 4. This implies that the distance between any two vertices of $Y \cup P$ is at most 6. Since d(z', s) = 4for any vertex $z' \in Y \cup P$, we have that any path between two vertices of $Y \cup P$ that pass through s has length at least 8. Therefore $s \notin I(Y \cup P)$ and the claim holds.

Claim 4. $V(G) \setminus \{s\} \subseteq I(S)$, for every hull set S of G.

Proof. Since all vertices of Y and P are pendant vertices we have $Y \cup P \subset S$ for any hull set S. Then, by Claim 3, it remains to show only that $V \cup U \cup T \subset I(S)$. By Claim 1, there exists $v \in S \cap D_i$, for every $i \in [m]$. Applying Claim 2 for each such v, we complete the proof of Claim 4.

Claim 5. Let $v_{i,a}$ and $v_{j,b}$ with $i \neq j$. Then $s \in I(\{v_{i,a}, v_{j,b}\})$ if and only if $\ell_{i,a}$ is the negation of $\ell_{j,b}$.

Proof. Note that $d(v_{i,a}, s) = d(v_{j,b}, s) = 3$. Then, $s \notin I(\{v_{i,a}, v_{j,b}\})$ if and only if $d(v_{i,a}, v_{j,b}) < 6$, i.e., $d(v_{i,a}, v_{j,b}) = 2$ or $d(v_{i,a}, v_{j,b}) = 4$. The first case is not possible because there is no possibility for $v_{i,a}$ and $v_{j,b}$ share a neighbour. The other possibility occurs if and only if two vertices, one neighbour of each one, share a neighbour. Since $N(v_{i,a}) = \{w_{i,a}, u_i\}$ and $N(v_{j,b}) = \{w_{j,b}, u_j\}$, the path of length 4 must be $v_{i,a}w_{i,a}zw_{j,b}v_{j,b}$. By the construction, $z = o_{i,a,j,b}$ and $\ell_{i,a}$ is the negation of $\ell_{j,b}$.

Now, let us finish the proof of Theorem 3.1. Let S be a hull set of G such that I(S) is a proper subset of V(G), that is, t(S) > 1. By Claim 4, $I(S) = V(G) \setminus \{s\}$. By Claim 1, for every $i \in [m]$, the set S contains some vertex of D_i . Suppose, by contradiction, that for some $j \in [m]$, it holds $u_j \in S \cap D_j$. Observe that s is in a shortest path between u_j and u, where u is some vertex of $S \cap D_k$, for any $k \neq j$, which is a contradiction. Then, for every $i \in [m]$, S contains v_{i,a_i} , where $a_i \in \{1, 2, 3\}$. By Claim 5, every pair of literals $\ell_{i,a} \in C_i$ and $\ell_{j,b} \in C_j$ associated to vertices $v_{i,a}$ and $v_{j,b}$ contained in S are not the negation of each other. Then, the truth assignment where $\ell_{i,a}$ is true if and only if $v_{i,a}$ belongs to S satisfies all clauses of \mathcal{F} .

Conversely, let \mathcal{A} be a truth assignment for $\{x_1, \ldots, x_n\}$ satisfying all clauses of \mathcal{F} . Consider the set S formed by the pendant vertices of G plus one vertex $v_{i,a}$ for every clause C_i of \mathcal{F} , where $v_{i,a}$ is chosen if $\ell_{i,a}$ assumes value true by \mathcal{A} . By Claim 3 and Claim 2, $V(G) \setminus \{s\} \subseteq I(S)$. And, since s is not a simplicial vertex of G, it follows that S is a hull set of G. By Claim 5, it remains to show that $s \notin I(u, v)$ for $u \in Y \cup P$ and $v \in V$. Note that d(u, s) = 4 and d(v, s) = 3, but $d(u, v) \leq 5$, completing the proof for k = 2.

Now suppose that k > 2. To finish the proof, it suffices to add to the construction of G a vertex s' twin to s and a copy of the graph H_{k-2} (as defined by equations (1) and (2) at the end of Section 2), such that s' and s are identified to the vertices v_1 and v_3 of the copy of H_{k-2} , respectively. It is not hard to check that this new graph G has infection time k if and only if \mathcal{F} is satisfiable.

4. Minimal and minimum hull sets in the monophonic convexity

Here, we call any hull set in the monophonic convexity by m-hull set. In this section we show how to use results of Leimer [22], about the clique separator decomposition, to improve the complexity of the fastest known (polynomial) algorithm [14] for finding a minimum m-hull set of a general graph. We also present a characterization of all minimum and of all minimal m-hull sets of general graphs. The later result is essential to our polynomial time algorithm for finding the maximum infection time in a distance-hereditary graph, that will be presented in Section 5.

A survey on clique separator decompositions can be found in [9]. Let us start with some preliminary notations. Given a graph G and a set $C \subseteq V(G)$, we say that C is a separator of G if there are non-empty sets $A, B \subset V(G) \setminus C$ such that every path in G, between some $a \in A$ and $b \in B$, contains a vertex in C. Separators that are cliques are called *clique separators*. If C is a clique separator of G and H is any connected component of G - C, the subgraph $G[V(H) \cup C]$ is called an C-component of G. The family of C-components of G is a decomposition of G. We say that G is reducible if it contains a clique separator, otherwise it is called a prime. We note that, in the literature, those are also called atoms. A maximal prime subgraph of G, or simply mp-subgraph of G, is a maximal induced subgraph of G that is prime.

A separator C is a minimal separator of G if no proper subset of C is a separator of G. And C is a relative minimal separator for G if there exist vertices $v, w \in V(G)$ such that v and w are in different components of G-C, but for every proper subset C' of C we have that u and w are in the same component of G-C'. Note that every minimal separator is a relative minimal separator for G, but the reverse is not true.

Assume that a reducible graph G is decomposed by a clique C and each C-component is decomposed further until all derived subgraphs are prime. A family of prime subgraphs of G that can be derived in that way will be called a *derived family*.

Lemma 4.1. Let C be a clique of G and G_1 be a C-component. Then any separator of G_1 is also a separator of G. Furthermore, if S is a relative minimal separator of G_1 then it is a relative minimal separator of G.

Proof. Let G, C, and G_1 be defined as in the statement. Let S be a separator of G_1 . So there exist vertices $x, y \in G_1$ such that any path from x to y in G_1 pass through S. Suppose that there is a path in G from x to y that avoids S. Since C is a clique, we can shorten this path to a path in G_1 (that shall also avoids S). Therefore, S also separates x from y in G. For the last part, it is trivial that if S is minimal for x, y in G_1 then it is also minimal for x, y in G.

By the previous proposition, we also notice that if G_1 is a *C*-component of *G*, where *C* is a clique, and $x, y \in V(G_1)$ then any induced path from *x* to *y* in *G* must have all its vertices in G_1 . That is the intuition on why the clique separator decomposition is usefull to solve problems in the monophonic convexity. The following result of [14] indicates why this decomposition is fruitful for finding *m*-hull sets.

Theorem 4.2. [14] If G is a prime graph that is not a complete graph, then every pair of non-adjacent vertices is an m-hull set of G.

Consider a graph G and a clique separator of C. Clearly, any mp-subgraph of a graph is "unbreakable" under a C-decomposition (that is, it must be contained in a C-component). Therefore, every mp-subgraph of G is an mpsubgraph of some C-component of G. The converse is not always true. But we have the following result.

Lemma 4.3. A derived family is exactly the family of mp-subgraphs of G if and only if it is obtained by a decomposition that uses only cliques that are relative minimal separators of G. *Proof.* This follows directly from (2.2) of [22] together with itens (i) and (iii) of Theorem 4.1 of [22]).

In our proofs, we will also need the following concepts. Given any graph G, consider an (arbitrary) ordering of the mp-subgraphs F_1, \ldots, F_t of G and then define $R_i = V(F_i) \cap (V(F_1) \cup \ldots \cup V(F_{i-1}))$, for $i \in [t]$. This ordering of the mp-subgraphs is a D-ordering if, for all $i \in \{2, \ldots, t\}$, there is j such that j < i and $R_i \subseteq V(F_j)$. According to Theorem 2.5 of [22], there is a D-ordering for the mp-subgraphs of any graph. Further, Proposition 2.4 of [22] says that every permutation of the mp-subgraphs of G that is a D-ordering has the same family of R_i sets. Then, one can define $\mathcal{R}(G) = \{R_2, \ldots, R_t\}$ and $R(G) = R_2 \cup \ldots \cup R_t$, taking as starting point any D-ordering of the mp-subgraphs of G. We will also use the following result of Leimer (again see Theorem 4.1 of [22], itens (i) and (iv)).

Theorem 4.4. [22] Let C be a set of vertices of a graph G. We have that C is a clique and a relative minimal separator for G if and only if $C \in \mathcal{R}(G)$.

In our algorithm, we shall adapt results of [14]. There, the authors have used a decomposition of the following type. Take a minimal clique C of G, look at the C-components of G, say G_1, \ldots, G_t . For each component take a minimal clique separator of that component and continue the process recursively to obtain a derived family. Note that the minimal separator of the components does not have to be a minimal separator of G. After that, they classify each set in the derived family into certain types and used this classification to find an m-hull set of G. We observe that any minimal clique separator in G_1 is a relative minimal clique for G_1 . By Lemma 4.1, it is also a relative minimal separator of G. By the same argument, for any component obtained at further steps in that decomposition, any minimal clique separator of that component is a relative minimal clique separator of G. By Lemma 4.3, the derived family obtained in [14] is precisely the family of mp-subgraphs.

As a consequence of the above, we can rewrite the characterization of minimum *m*-hull sets presented in [14]. Let F be any *mp*-subgraph. We say that F is:

type 0 if $V(F) \cap R(G)$ is not a clique;

type 1 if $V(F) \cap R(G)$ is a clique and there is a vertex $u \in V(F)$ not adjacent to some vertex of $V(F) \cap R(G)$; **type** 2 if $V(F) \cap R(G)$ is a clique, V(F) is not a clique, and every vertex of $V(F) \cap R(G)$ is universal in F;

type 3 if V(F) is a clique.

As observed in [14], one can check that F is of exactly one of the above types. This result implies that we can also rewrite the characterization of minimum *m*-hull sets from [14] using *mp*-subgraphs.

Theorem 4.5. A set S is a minimum m-hull set of a graph G if and only if for every mp-subgraph F of G, the set S satisfies the following conditions:

- if F is of type 0, then $S \cap V(F) = \emptyset$;
- if F is of type 1, then $S \cap (V(F) \setminus R(G)) = \{u\}$ for a vertex u not adjacent to some vertex of $V(F) \cap R(G)$;
- if F is of type 2, then $S \cap (V(F) \setminus R(G)) = \{u, v\}$ for some pair u, v of non-adjacent vertices;
- if F is of type 3, then $V(F) \setminus R(G) \subseteq S$.

The algorithm used in [14] to find the monophonic hull number of a graph on *n* vertices and *m* edges, had running time $O(n^3m)$ and consisted in finding minimal clique separators (using an algorithm for finding a clique separator of a graph [27]) and recursively obtaining the derived family. Now, using Theorem 4.5, the complexity analysis is the following. The *mp*-subgraphs and $\mathcal{R}(G)$ can be found in O(nm) time using the algorithm of Leimer [22]. The number of *mp*-subgraphs of *G* is O(n) [22, 25]. The type of each *mp*subgraph can be found in O(m) time. To compute the hull number we only have to add 1 for each *mp*-subgraph of type 1, two for each of type 2 and V(F), and $|V(F) \setminus R(G)|$ for each of type 3. We note that since the sets $V(F) \setminus R(G)$ are disjoint, there is no double counting in the previous summation. Therefore, one can find a minimum hull set of any graph in the monophonic convexity in O(nm) steps.

In the sequel, we present a characterization of minimal m-hull sets. First, we state two simple lemmas from [14] rewriting them using the equivalence of the atoms in their decomposition with the mp-subgraphs.

Lemma 4.6. [14] Let F be an mp-subgraph of G. If F is of type $t, t \in \{1, 2\}$, then every m-hull set of G contains at least t vertices of $V(F) \setminus R(G)$.

Lemma 4.7. [14] A vertex v is simplicial in G if and only if it belongs to a mp-subgraph of G of type 3 and $v \notin R(G)$.

Next, it is useful to show that R(G) is contained in the interval of any *m*-hull set of *G*.

Theorem 4.8. Let G be a graph and S a minimal m-hull set of G. Then $R(G) \cap S = \emptyset$ and $R(G) \subset I(S)$.

Proof. For every $R_i \in \mathcal{R}(G)$, Theorem 4.4 says that R_i is a minimal clique separator of G relative to some pair of vertices $v_{i,1}, v_{i,2}$. Denote by $H_{i,1}$ and $H_{i,2}$ the R_i -components of G containing $v_{i,1}$ and $v_{i,2}$, respectively. Note that the complement, in G, of $V(H_{i,j}) \setminus R_i$ is a m-convex set for $j \in 1, 2$. This implies that $S \cap (V(H_{i,j}) \setminus R_i)$ is non-empty. And, by the minimality of R_i , every vertex of R_i has a neighbor in $H_{i,1}$ and in $H_{i,2}$. This implies that there are vertices $v'_{i,j} \in S \cap (V(H_{i,j}) \setminus R_i)$, for $j \in \{1, 2\}$ such that, for every $u \in R_i$ there is an induced path P_1 from $v'_{i,1}$ to u in $H_{i,1}$ and an induced path P_2 from $v'_{i,2}$ to u in $H_{i,2}$. Since $V(P_j) \cap R_i = \{u\}$, for $j \in \{1, 2\}$, it is clear that the concatenation of P_1 and P_2 is an induced path P_u of G, concluding the proof that $R(G) \subset I(S)$. Since S is minimal, we also have that $R(G) \cap S = \emptyset$. \Box

Now, we can present the characterization of minimal m-hull sets of general graphs.

Theorem 4.9. A set S is a minimal m-hull set of a graph G if and only if for every mp-subgraph F of G, S satisfies the following conditions:

- if F is of type 0, then $S \cap V(F) = \emptyset$;
- if F is of type 1, then either $S \cap (V(F) \setminus R(G)) = \{u\}$ for a vertex u not adjacent to some vertex of $V(F) \setminus R(G)$ or $S \cap (V(F) \setminus R(G))) = \{u, v\}$ for some pair u, v of non-adjacent vertices;
- if F is of type 2, then $S \cap (V(F) \setminus R(G)) = \{u, v\}$ for some pair u, v of non-adjacent vertices;
- if F is of type 3, then $V(F) \setminus R(G) \subseteq S$.

Proof. Let k be the number of mp-subgraphs of G, where $k \ge 1$. Let S be a minimal m-hull set of G and F an mp-subgraph F of G of type t. By

Theorem 4.8, we know that $R(G) \cap S = \emptyset$, then the vertices of F contained in S belong to $V(F) \setminus R(G)$.

If t = 0, there are members R_x and R_y of $\mathcal{R}(G)$ contained in F such that $R_x \cup R_y$ is not a clique. Defining the vertices $v_{x,1}, v_{x,2}, v_{y,1}, v_{y,2}$ as in the proof of Theorem 4.8, we can conclude that at least one of $v_{x,1}, v_{x,2}$ and one of $v_{y,1}, v_{y,2}$ does not belong to V(F). Therefore, for every vertex u of $R_x \cup R_y$ there is an induced path P_u with endpoints in S that pass through u, similar to the one constructed in the proof of Theorem 4.8. Then, since S is minimal, S does not contain any vertex of F.

If t = 1, by Lemma 4.6, the set S contains at least one and at most two vertices of F. If $S \cap F$ contains one vertex, it is clear that it must be not adjacent to some vertex of $V(F) \cap R(G)$. Now, suppose that $S \cap F$ contains two vertices, say x and y. If these two vertices were adjacent, then at least one of them, say x, must be not adjacent to some vertex x' of some member $R_x \in \mathcal{R}(G)$ with $R_x \subseteq V(F) \cap R(G)$. Now, considering vertices $v'_{x,1}$ and $v'_{x,2}$ analogously defined as in the previous case, we can say that at least one of them, say $v_{x,1}$, is not contained in V(F). Then, we can conclude that there is an induced path from x to $v_{x,1}$ containing x'. This would imply that $y \in H(S \setminus \{y\})$. Therefore $xy \notin E(G)$.

If t = 2, by Lemma 4.6, the set S contains two non-adjacent vertices of $V(F) \setminus R(G)$. Since these two vertices form an *m*-hull set of G and S is minimal, we have $|S \cap (F \setminus R(G))| = 2$.

The result for t = 3 follows directly from Lemma 4.7 and the fact that Theorem 4.8 implies that $S \cap R(G) = \emptyset$.

Conversely, let S be a set of vertices of G satisfying, for every mpsubgraph F of G, the property associated with the type of F. If, for some mp-subgraph F of G of type 1, the set S contains two non-adjacent vertices, then $V(F) \subseteq H(S)$, by Theorem 4.2. This implies that H(S) contains a set satisfying all four conditions of Theorem 4.5, which implies that S is an m-hull set of G.

5. Infection time on distance-hereditary graphs

A connected graph G is called *distance-hereditary* if for every induced subgraph G' of G and every pair of vertices $u, v \in V(G')$ it holds that $d_G(u, v) = d_{G'}(u, v)$. This graph class admits a characterization by forbidden induced subgraphs and recognition in polynomial time [4]. Observe that geodesic and monophonic convexities coincide in distance-hereditary graphs, since for a graph in this class every induced path is also a shortest path. Although even the computation of the interval function is a NP-hard problem in the monophonic convexity for general graphs [14], the previous fact allows us to use the known algorithms for computing the interval function in the geodesic convexity together with the characterization of minimal m-hull sets for solving the maximum infection problem in distance-hereditary graphs. Next, we present a polynomial algorithm for that.

Since every hull set of a graph G contains a minimal hull set, t(G) can be determined by finding the maximum t(S) among all minimal hull sets of G. Theorem 4.9 characterizes the minimal *m*-hull sets of G according to their intersections with *mp*-subgraphs. The overall idea of our algorithm consists in decomposing G into *mp*-subgraphs, and reducing the problem of computing the maximum infection time of G to the problem of computing the infection time of a polynomial number of sets of these prime graphs.

We need of some definitions. Let G be a graph, S a hull set of G, and F an mp-subgraph of G. Let

- F^* be the graph obtained from F by adding, for every $C_j \in \mathcal{R}(G)$ such that $C_j \subset V(F)$, a vertex x_j and the edges $\{vx_j : v \in C_j\}$; and
- $S_F = (S \cap V(F)) \cup \{x_j : C_j \subseteq R(G) \cap V(F)\};$

The following result shows that, for every mp-subgraph F of a graph G and any hull set S of G, if we start an infection with S, we can determine which vertices of F become infected at any given time by looking only at infected vertices in S_F .

Lemma 5.1. Let F be an mp-subgraph and S an m-hull set of a graph G. Then $I_G^k(S) \cap V(F) = I_{F^*}^k(S_F) \cap V(F)$, for any $k \ge 0$.

Proof. For k = 0, the definitions imply that $S \cap V(F) = S_F \cap V(F)$. For k > 0, observe that it suffices to show $I_G(S) \cap V(F) = I_{F^*}(S_F) \cap V(F)$. And this is equivalent to show that a vertex $v \in V(F) \setminus S$ is in an induced path of G between two vertices of S if and only if v is in an induced path of F^* between two vertices of S_F .

Let $w \in V(F) \setminus S$, and P_{uv} be an induced (u, v)-path of G passing through w such that $u, v \in S$. Denote by P_{uw} the subpath of P_{uv} beginning at u and ending at v. If $u \in V(F)$, define $P'_{uw} = P_{uw}$. Otherwise P_{uw} contains a vertex u' of some clique $C_i \in \mathcal{R}(G)$ such that $C_i \subset V(F)$. In this case, define P'_{uw} as x_i concatenated with the subpath of P_{uw} beginning at u' and ending at

w. Define P_{vw} and P'_{vw} analogously. It is clear that P'_{uw} concatenated with P'_{vw} is a induced path of F^* passing through w such that the extremities are two vertices of S_F .

Conversely, let $w \in V(F^*) \setminus S_F$, and P_{uv} be an induced (u, v)-path of F^* passing through w such that $u, v \in S_F$. Denote by P_{uw} the subpath of P_{uv} beginning at u and ending at v. If $u \in V(F)$, define $P'_{uw} = P_{uw}$. Otherwise $u = x_j$ for some $R_i \in \mathcal{R}(G)$ such that $R_i \subset V(F)$. Theorem 4.4 says that R_i is a minimal separator of G relative to some pair of vertices $v_{i,1}, v_{i,2}$. Without loss of generality we can say that the R_i -component $H_{i,1}$ containing $v_{i,1}$ does not contain F. Observe that $H_{i,1}$ is a proper m-convex sets of G and that every vertex of R_i has a neighbor in $H_{i,1}$. This implies that there is a vertex $v'_{i,1} \in S \cap (V(H_{i,1}) \setminus R_i)$. Then, consider an induced path $P_{i,1}$ of $H_{i,1} - R_i$ from $v'_{i,1}$ to some vertex u'' that is neighbor to some vertex $u' \in R_i \cap V(P_{uw})$. Now, define P'_{uw} as the concatenation of the three paths $P_{i,1}$, the edge u'u'', and the subpath of P_{uw} from u' to w.

Next, define P_{vw} and P'_{vw} analogously. It is clear that P'_{uw} concatenated with P'_{vw} is a induced path of G passing through w such that the extremities are two vertices of S.

This result has the following important consequence for the algorithm.

Corollary 5.2. Let G be a graph. Then $t(G) = \max\{t_{F^*}(S_F) : S \text{ is a minimal m-hull set and F is an mp-subgraph of G}.$

We can now describe the algorithm for computing t(G) of a distancehereditary graph G with n vertices and m edges in the geodesic or monophonic convexity. The mp-subgraphs and $\mathcal{R}(G)$ can be found in O(nm)time [22]. The number of mp-subgraphs of G is O(n) [22, 25]. The type of each mp-subgraph can be determined in O(m) time. By Corollary 5.2, for every mp-subgraph F of G, we need to find the set maximizing $t_{F^*}(S_F)$ among all minimal hull sets S of G. However, by Lemma 5.1, it suffices to consider all distinct intersections of V(F) with all minimal hull sets of G. By Theorem 4.9, the number of such intersections is $O(n^2)$ for mp-subgraphs of types 1 and 2, and is O(1) for mp-subgraphs of types 0 and 3. Finding ksuch that $I^k(S) = H(S)$ can be done in O(nm) steps for general graphs in the geodesic convexity [13] for any set S. Then, a direct analysis leads to an algorithm with time complexity $O(n^4m)$. However, we observe that the sum of the distinct intersections of all mp-subgraphs of G with all minimal hull sets of G is $O(n^2)$. Then, doing an amortized analysis, we conclude that the total complexity of this algorithm is $O(n^3m)$.

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