# Graphs of separability at most two: structural characterizations and their consequences

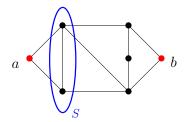
## Ferdinando Cicalese<sup>1</sup> Martin Milanič<sup>2</sup>

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Raziskovalni matematični seminar, FAMNIT, 18. oktober 2010

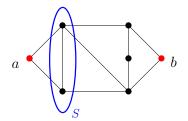
An (a, b)-separator is a set  $S \subseteq V(G)$  such that a and b are in different connected components of G - S.



Separability of  $\{a, b\}$ : the smallest size of an (a, b)-separator.

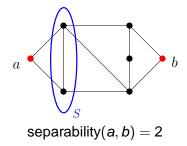
Cicalese–Milanič Graphs of separability at most two

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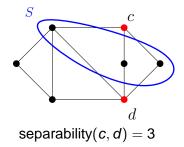


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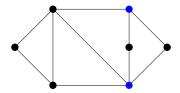
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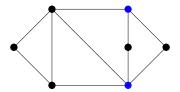
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a graph of separability 3

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## By Menger's Theorem,

separability(a, b) = min size of an (a, b)-separator

= max # internally vertex-disjoint (a, b)-paths.

## Therefore, for a non-complete graph G,

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## $\mathcal{G}_k = \{ \mathbf{G} : separability(\mathbf{G}) \leq k \}.$

Graphs in  $\mathcal{G}_k$ :

- generalize graphs of maximum degree k,
- generalize pairwise k-separable graphs,

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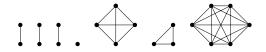
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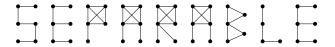
## Can we characterize graphs of separability at most k, at least for small values of k?

## Structure of graphs in $\mathcal{G}_0$ and $\mathcal{G}_1$

Graphs of separability 0 = disjoint unions of complete graphs



Graphs of separability at most 1 = block graphs: graphs every block of which is complete.



## Outline

## $\mathcal{G}_2$ , graphs of separability at most 2:

- generalize complete graphs, trees, cycles, block-cactus graphs
- characterizations
- algorithmic and complexity results

#### Graphs in $\mathcal{G}_k$ :

connection to the parsimony haplotyping problem

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## **Graphs in** $\mathcal{G}_k$ :

connection to the parsimony haplotyping problem

# Characterizations

#### Theorem

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### Corollary

Every graph in  $G_2$  contains either a simplicial vertex or two adjacent vertices of degree 2.

 $v \in V(G)$  is simplicial if its neighborhood is a clique.

#### Corollary

Graphs in  $\mathcal{G}_2$  are  $\chi$ -bounded: There exists a function f such that for every  $\mathbf{G} \in \mathcal{G}_2$ ,

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## Is tree-width of graphs in $\mathcal{G}_2$ bounded by a constant?

No  $\therefore$  complete graphs are in  $\mathcal{G}_2$ .

Corollary

For every  $G \in \mathcal{G}_2$ ,

 $tw(G) \leq \max\{2, \omega(G) - 1\}.$ 

(This is best possible: no similar result holds for  $G_3$ .)

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## Characterization by forbidden induced subgraphs

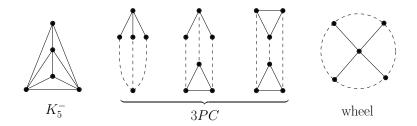
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#### Theorem

 $\mathcal{G}_2 = \{K_5^-, 3PC, wheels\}$ -induced-subgraph-free graphs.



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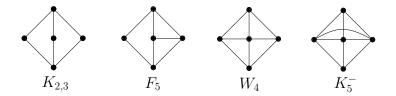
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 $\mathcal{G}_2 = \{\textit{K}_{2,3},\textit{F}_5,\textit{W}_4,\textit{K}_5^-\}\text{-induced-minor-free graphs}.$ 



#### Theorem

 $\mathcal{G}_k$  is closed under induced minors if and only if  $k \leq 2$ .



a graph from  $\mathcal{G}_3$  contracted to a graph of separability 6

## Algorithms and complexity

Some problems are solvable in polynomial time for graphs in  $\mathcal{G}_k$ , for every *k*:

- recognition
  - $O(|V(G)|^2)$  max flow computations
- finding a maximum weight clique
  - polynomially many maximal cliques

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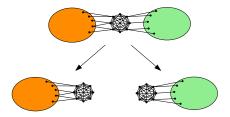
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## Good news

For graphs in  $\mathcal{G}_2$ , the structure theorem leads to polytime algorithms for:

- maximum weight independent set (NP-hard for  $\mathcal{G}_3$ )
- coloring (NP-hard for  $\mathcal{G}_3$ )

The algorithms are based on the decomposition by clique separators.



Whitesides 1981, Tarjan 1985

## Not so good news

#### Is clique-width of graphs in $\mathcal{G}_2$ bounded by a constant?

#### Proposition

Graphs in  $\mathcal{G}_2$  are of unbounded clique-width.

#### Proposition

- The dominating set problem for graphs in G<sub>2</sub>.
- The simple max cut problem for graphs in  $\mathcal{G}_2$ .
- The 3-colorability problem for graphs in *G*<sub>3</sub> (planar, of maximum degree 6).

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# Connection to the parsimony haplotyping problem

#### PARSIMONY HAPLOTYPING:

Given: a set of *n* vectors in  $\{0, 1, 2\}^m$  (genotypes).

Task: find the minimum size of a set of vectors in  $\{0, 1\}^m$  (*haplotypes*) such that every genotype can be expressed as the sum of two haplotypes from the set.

Addition rules: 0 + 0 = 0, 1 + 1 = 1, 0 + 1 = 1 + 0 = 2

## A problem from computational biology

Compatibility graph *G*: the graph with  $V(G) = \{\text{genotypes}\}$  and  $E(G) = \{gg' : \nexists r \text{ such that } \{g_r, g'_r\} = \{0, 1\}\}.$ 

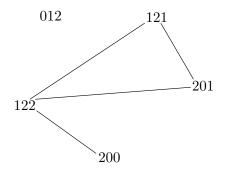
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A parsimony haplotyping instance is *k*-bounded if in every coordinate, at most *k* genotypes contain a 2.

#### Theorem

 $G_k = \{$  compatibility graphs of k-bounded PH instances $\}$ .

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#### Theorem

 $\mathcal{G}_k = \{ \text{compatibility graphs of } k \text{-bounded PH instances} \}.$ 

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	$\mathcal{G}_1$	$\mathcal{G}_2$	$\mathcal{G}_{3}$
PARSIMONY HAPLOTYPING	polynomial	?	NP-complete

van Iersel–Keijsper–Kelk–Stougie 2008 Sharan–Halldórsson–Istrail 2006 For  $k \ge 3$ , characterize graphs in  $\mathcal{G}_k$  in terms of:

- forbidden induced subgraphs,
- decomposition properties.

For  $k \ge 3$ , determine whether graphs in  $\mathcal{G}_k$  are  $\chi$ -bounded.

Determine the complexity of:

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## Conclusion

Separators of size at most 2 sometimes help...

- decomposition along separating cliques of size at most two into cycles and complete graphs,
- $tw(G) \leq f(\omega(G))$ ,
- $\chi(\mathbf{G}) \leq f(\omega(\mathbf{G})).$



...but not always:

- dominating set is NP-complete,
- simple max cut is NP-complete,
- clique-width is unbounded.



## HVALA ZA POZORNOST