

Clustering Systems of Phylogenetic Networks

Marc Hellmuth

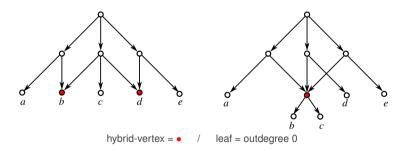
Department of Mathematics Faculty of Science Stockholm University

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Basics

All networks N = (V, E) considered here are

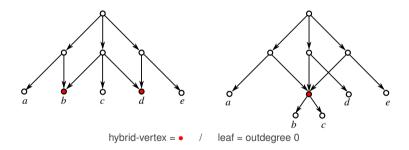
- 1. DAGs with a single root
- 2. phylogenetic (= no indegree 1 and outdegree 1 vertices)



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(In)comparable vertices

 $u \leq_N v$ if v is an ancestor of u, i.e., there is a directed path from v to u. u and v are \leq_N -comparable if $u \leq_N v$ or $v \leq_N u$ Otherwise, u and v are \leq_N -incomparable

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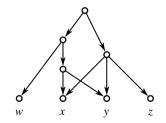
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 $\mathscr{C}_N := \{C(v) \mid v \in V\}$

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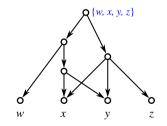
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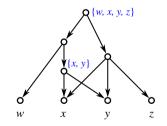
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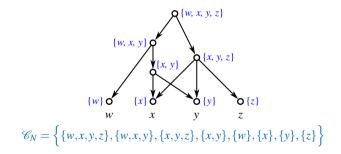
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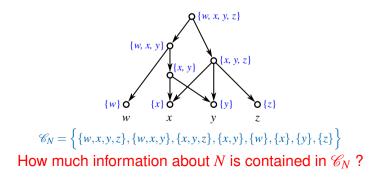
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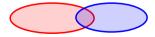
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where C(v) denotes the set of leaves "below" v.

Folklore (Trees):

• $C, C' \in \mathscr{C}$ overlap, if $C \cap C' \notin \{C, C', \emptyset\}$.

A clustering system is a hierarchy if it does not contain pairwise overlapping sets.



OVERLAP

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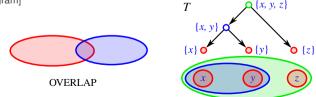
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• There is a 1-to-1 correspondence between rooted phylogenetic trees and hierarchies. [keyword: Hasse-diagram]



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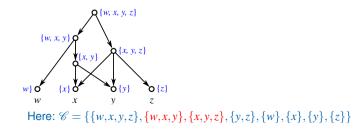
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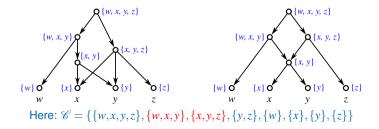
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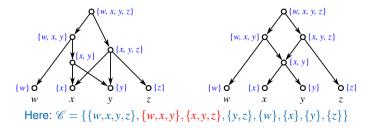
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- \mathscr{C}_N is usually *not* a hierarchy for general networks
- No 1-to-1 correspondence between networks and clustering systems.



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Central Questions: Given %.

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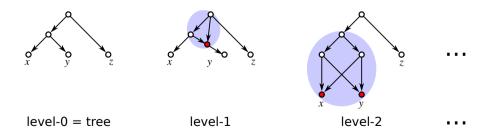
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Let us try to answer some of the questions for level-1 networks.

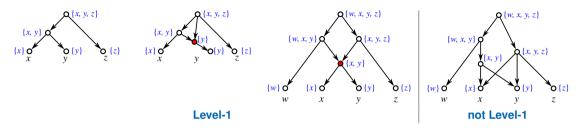
Level-k networks

A **block** *B* in a network is a maximal biconnected subgraph.

A network *N* is **level-***k* if each block *B* in *N* contains $\leq k$ hybrid-vertices (distinct from root ρ_B of *B*).

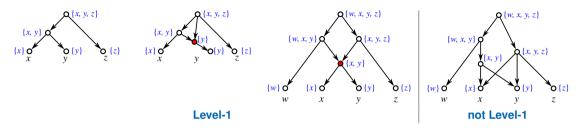






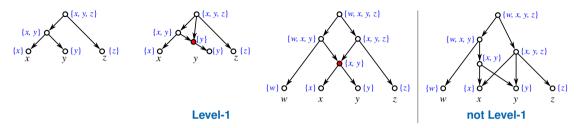
In trees *T*: $u \preceq_T v \iff C(u) \subseteq C(v)$





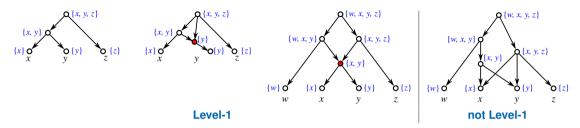
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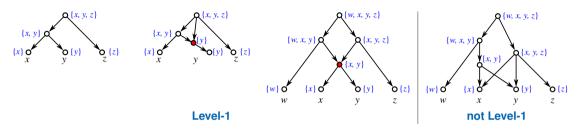


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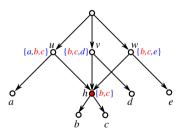
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Lemma (H., Schaller, Stadler, 2023)

In level-1 networks N: u and v are \leq_N -incomparable in $N \iff C(u) \cap C(v) \in \{\emptyset, C(h_B)\}$ where $h_B \neq u, v$ is the unique hybrid in block B that contains u and v.

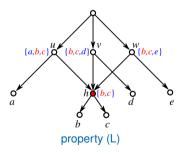
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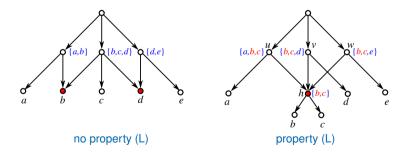
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A clustering system \mathscr{C} satisfies **property (L)** if $C \cap C_1 = C \cap C_2$ for all $C, C_1, C_2 \in \mathscr{C}$ where *C* overlaps both C_1 and C_2 .



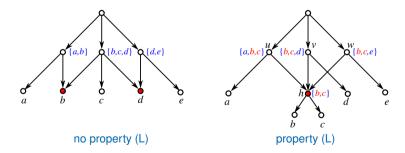
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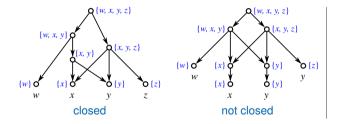
For all level-1 networks N, the set \mathcal{C}_N satisfies property (L).

Closed clustering systems

A clustering system \mathscr{C} is **closed** if $A \cap B \in \mathscr{C}$ for all $A, B \in \mathscr{C}$ with $A \cap B \neq \emptyset$.

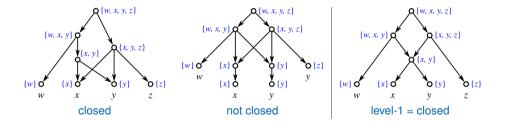
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Lemma (H., Schaller, Stadler, 2023)

For level-1 networks N, the set \mathscr{C}_N is always closed.

Characterization for level-1 Networks

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Theorem (H., Schaller, Stadler, 2023)

For \mathscr{C} , there is a level-1 network N with $\mathscr{C}_N = \mathscr{C} \iff \mathscr{C}$ is closed and satisfies (L).

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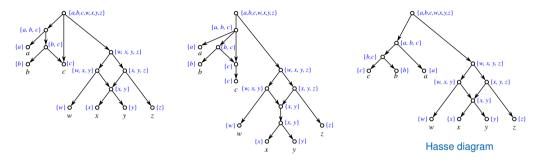
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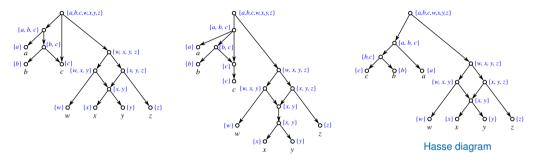
Recognition of such \mathscr{C} and reconstruction of such a level-1 network can be done in polynomial time.

Uniqueness Results



In general, several different level-1 networks may represent the same ${\mathscr C}$

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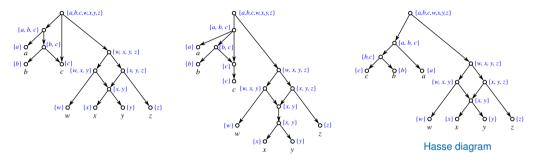


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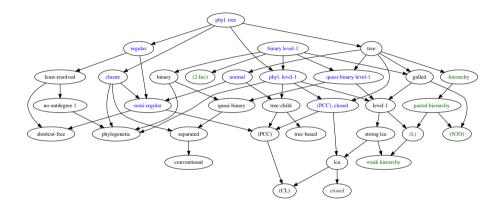
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Every level-1 network N is a refinement (=adding shortcuts + expand vertices) of the Hasse-diagram of \mathscr{C}_N .

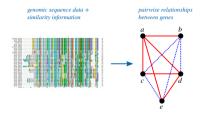
Other types of networks ...



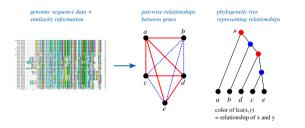
The latter results and plenty of other characterizations for dozens of other network types can be found in *Hellmuth, Stadler, Schaller, Clustering Systems of Phylogenetic Networks, Theory in Biosciences (142), 301-358, 2023*

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- we wanted to understand in more detail inference of horizontal gene transfer and orthology

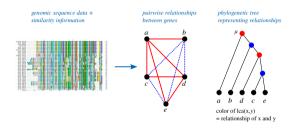
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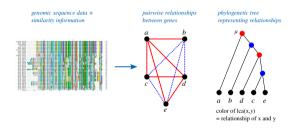
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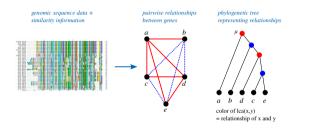
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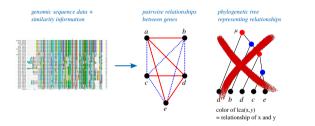
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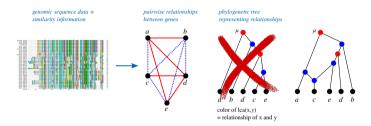
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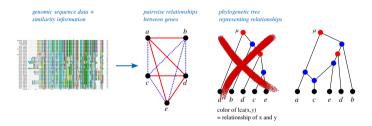
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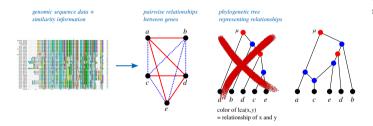
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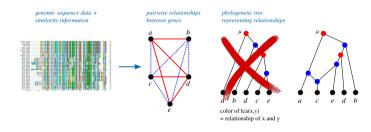
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A first starting point is provided by

Theorem (Shanavas, Changat, H., Stadler, 2024)

For \mathscr{C} , there is a network N with $\mathscr{C} = \mathscr{C}_N$ and pairwise lca-property $\iff \mathscr{C}_N$ is pre-binary, i.e., there is a unique incl.-min. cluster $C \in \mathscr{C}$ such that $\{x, y\} \subseteq C$ for all $x, y \in L(N)$

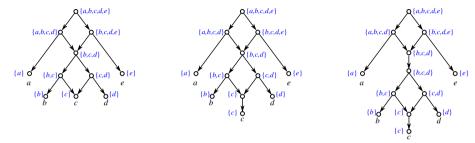
Shanavas, Changat, Hellmuth, Stadler Unique Least Common Ancestors and Clusters in Directed Acyclic Graphs, LNCS vol 14508, 2024

- David Schaller (Biontech, GER)
- Peter F. Stadler (Uni Leipzig, GER)
- Ameera Vaheeda Shanavas (Uni Kerala, IND)
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Thanks!

Uniqueness Results



We obtain uniqueness for mild restrictions!

Theorem

Let \mathscr{C} be a closed clustering system that satisfies (L). Then, there is a unique shortcut-free level-1 network N with $\mathscr{C}_N = \mathscr{C}$ that satisfies precisely one condition:

- *N* contains no vertex *v* with outdegree 1 (=Hasse diagram)
- every leaf in N has indegree 1 and all vertices v with outdegree 1 are adjacent to leaves.
- every hybrid in *N* has outdegree 1 (thus all leaves have indegree 1).

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Theorem

There is a galled tree N with $\mathscr{C}_N = \mathscr{C} \iff \mathscr{C}$ is **closed** and satisfies **Property (L)** and does not contain three distinct pairwise overlapping clusters.

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Binary network = network where all non-hybrids v is either a leaf or has outdeg(v) = 2, and every hybrid v satisfies indeg(v) = 2 and outdeg(v) = 1.

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Theorem

There is a binary level-1 network N with $\mathscr{C}_N = \mathscr{C} \iff \mathscr{C}$ is **closed** and satisfies **Property (L)** and for all clusters $C \in \mathscr{C}$, there are at most two inclusion-maximal clusters $A, B \in \mathscr{C}$ with $A, B \subsetneq C$ and at most two inclusion-minimal clusters $A, B \in \mathscr{C}$ with $C \subsetneq A, B$.

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Theorem

 \mathscr{C} is compatible w.r.t. a level-1 network $N \iff \mathscr{C}$ is satisfies **Property (L)**.

In this case, compute $A \cap B$ for all overlapping $A, B \in \mathscr{C}$ and add it to \mathscr{C} if their intersection is not present. (can be done in polynomial time)



Fig. 2. Consider the DAG G with leaf set X = L(G)where $\mathscr{C}_G = \{\{x\}, \{y\}, \{z\}, \{w\}, \{x, y, z\}, X\}$. Here, \mathscr{C}_G satisfies (**KS**) and (**KC**) for k = 2. By definition, \mathscr{C}_G is thus pre-binary. However, G is not a pairwise lea-network since lca(x, y), lea(x, z), and lea(y, z) are not defined. Moreover, \mathscr{C}_G also satisfies (**KC**) for k = 3 but G is not a 3-lea-network since lea(x, y, z) is not defined.

pre-binary clustering system but not pairwise lca-property