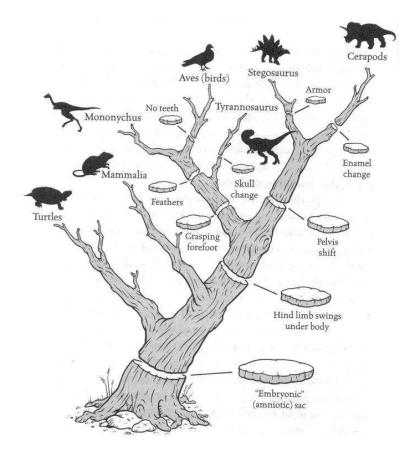
Small phylogeny problem: character evolution trees

Arvind Gupta Ján Maňuch Ladislav Stacho and Chenchen Zhu

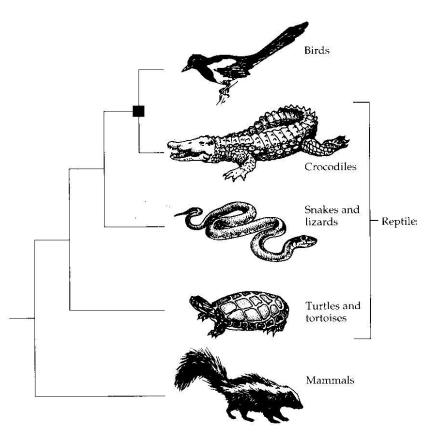
School of Computing Science and Department of Mathematics Simon Fraser University, Canada

 science determining ancestor/descendent relationships between species

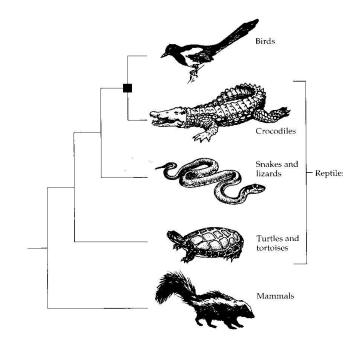
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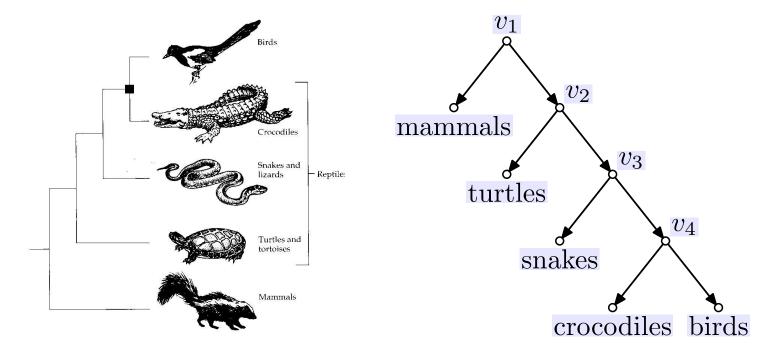


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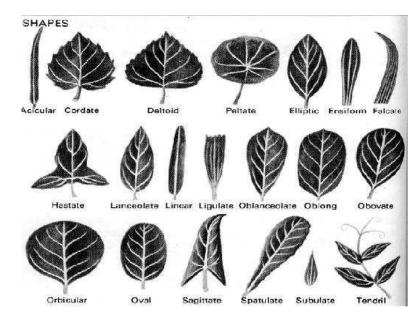
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- Each character (a morphological feature, a site in a DNA sequence, etc.) takes on one of a few possible states.
- Species can be modeled as vectors of states of a group of characters.

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- given:
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 - set of states for each character
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- This problem is NP-hard [Foulds, Graham (1982)].

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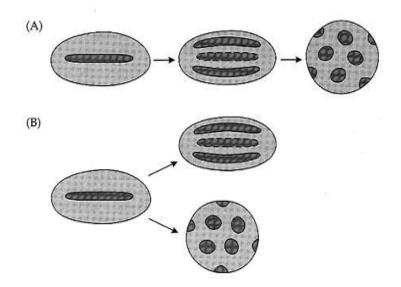
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- There are polynomial algorithms: [Fitch (1971)] (uniform costs), [Sankoff (1975)] (non-uniform costs).

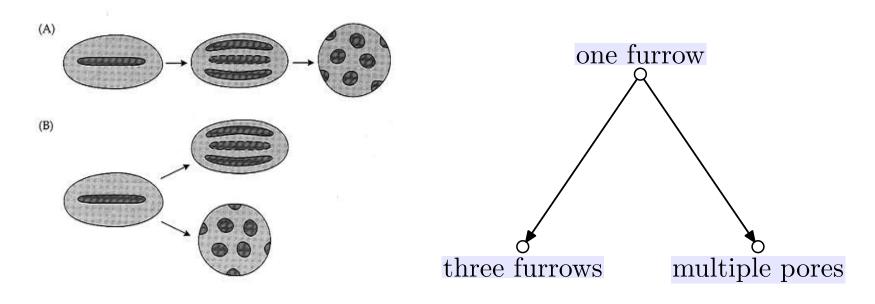
Character evolution tree

- So far we assumed that during one evolutionary step one state of a character can change to any other state. However, for many characters character state order and character state polarity can be observed.
 - **Example:** character evolution trees for pollen



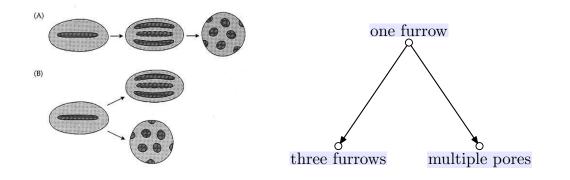
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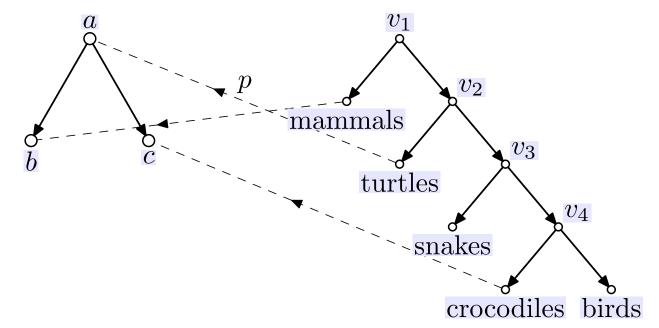
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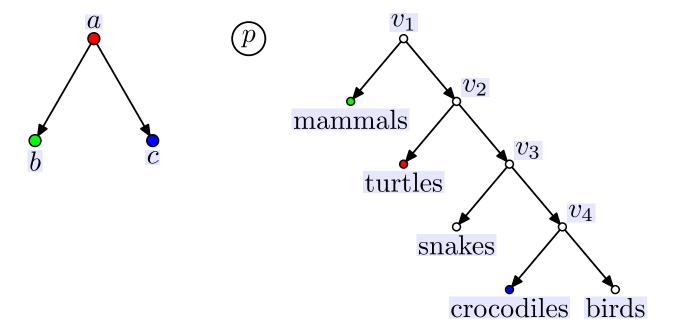


Our goal is to find a method of directly comparing a character evolution trees with a phylogenetic trees.

- **given:**
 - character evolution tree H_h with V(H) being states of the character
 - a phylogeny tree G_g with leaves L(G) being extant species
 - a *leaf labeling* $p : L(G) \rightarrow V(H)$ (a partial function)

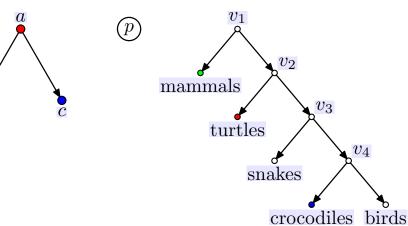


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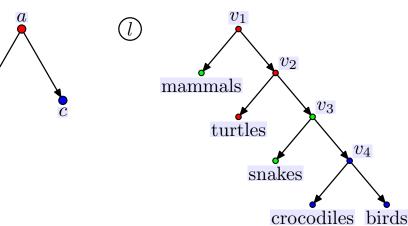


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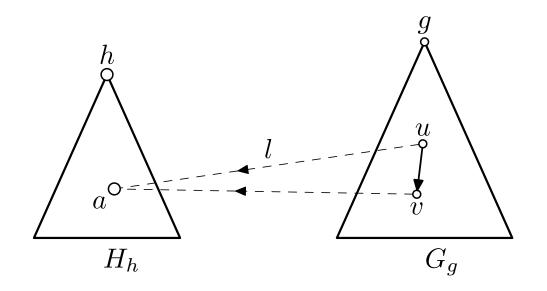


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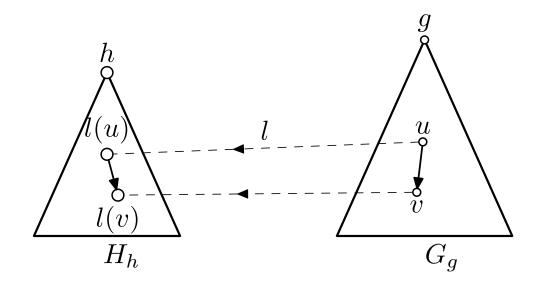


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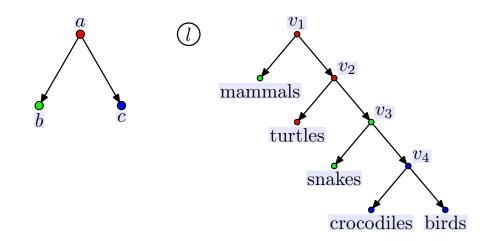


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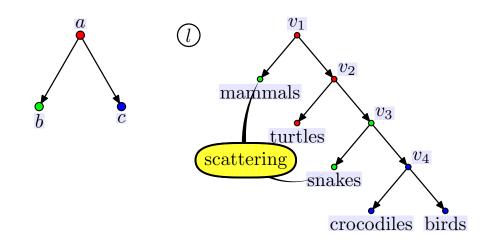


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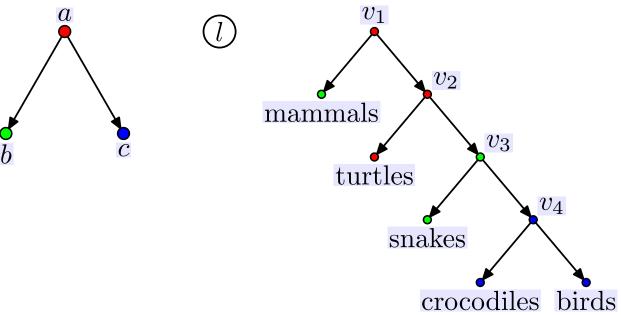
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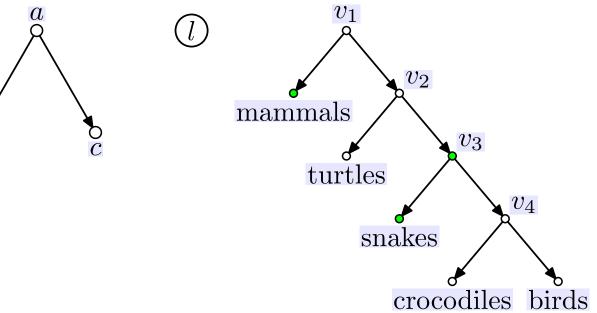
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- existence of such labeling *l* is very close to graph-theoretical notion of graph minors

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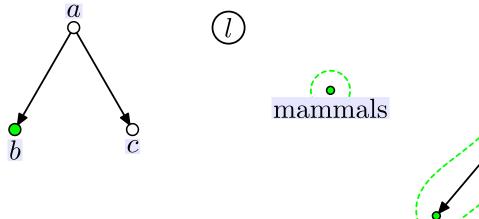


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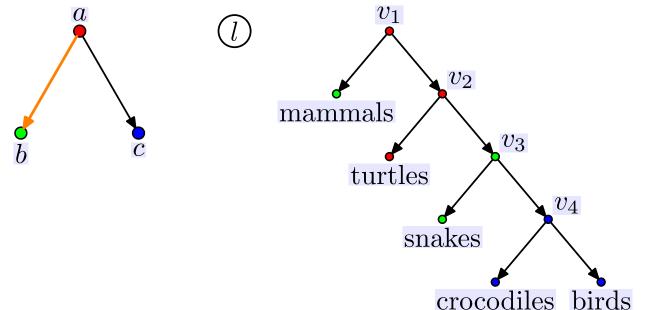
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snakes

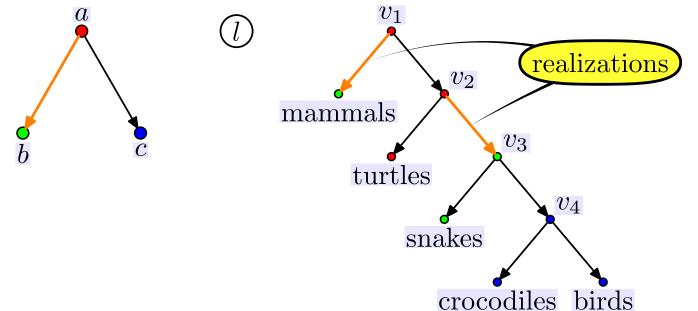


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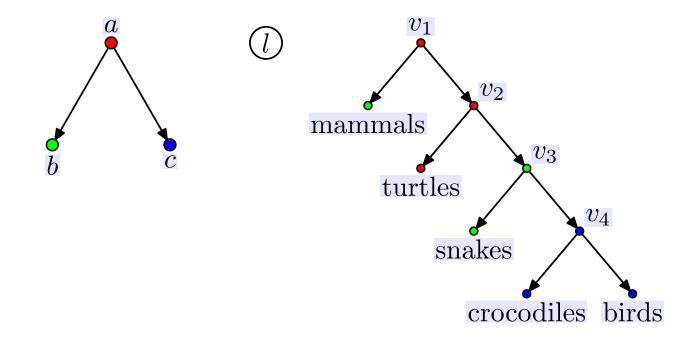


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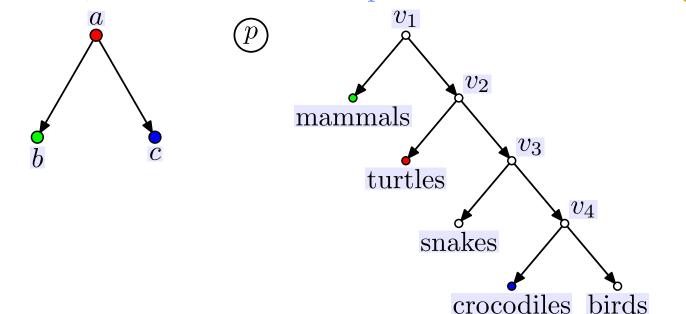
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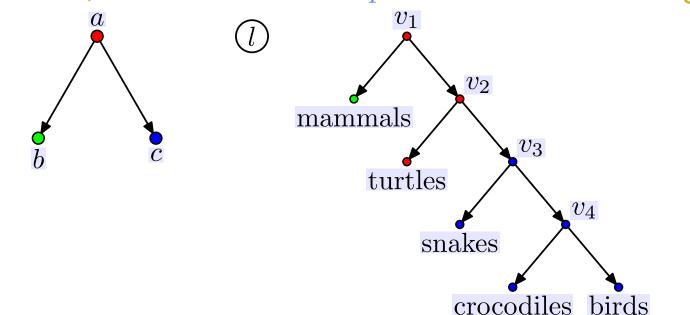
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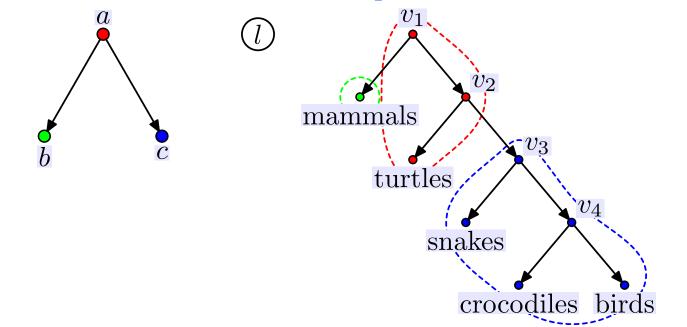
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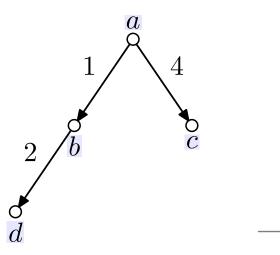
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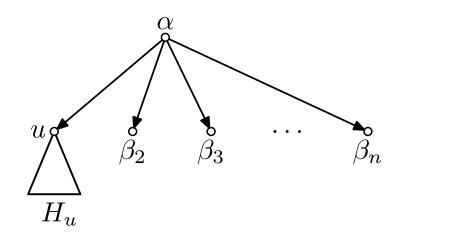
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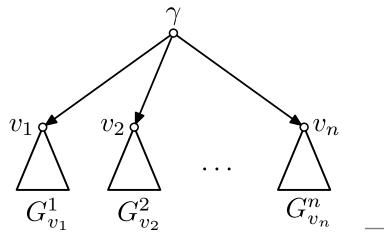


Consider an instance of Tree minor problem:

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Rooted-tree minor problem. Given two rooted trees H_h and G_g , and a leaf labeling $p : L(G_g) \to V(H)$. Decide whether H_h is a rooted-tree minor of G_g with respect to p.

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- Theorem 2. If the leaf labeling p is complete, then Rooted-tree minor problem can be decided in linear time.

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- Theorem 2. If the leaf labeling p is complete, then Rooted-tree minor problem can be decided in linear time.
 - Compute the LCA-tree.

(The label of each species is the least common ancestor of labels of its children.)

This can be done in linear time (requires preprocessing on the character evolution tree [Harel, Tarjan (1984)]).

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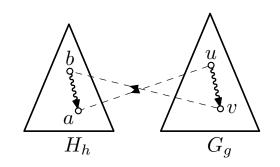
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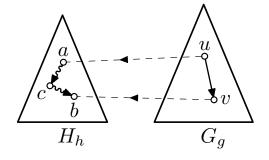
Crucial lemma: the ends of single branch paths are already fixed correctly for any labeling *l* satisfying the definition of *Rooted-tree minor*.

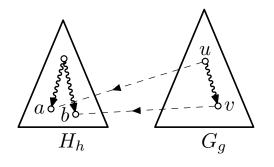
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 - Compute the LCA-tree.
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 - If each evolutionary transition has exactly one realization accept the input.

Incongruences



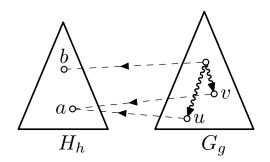




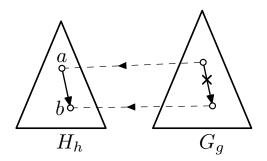
Inversion

Transitivity

Addition



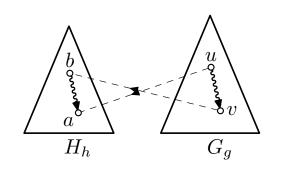
Separation

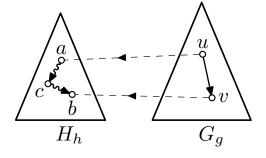


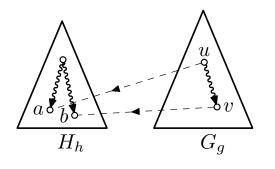
Negligence

Remark: solid lines represent arcs, wavy lines directed paths of length at least one.

Incongruences



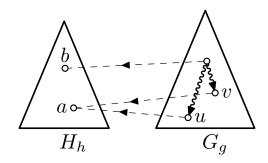




Inversion

Transitivity

Addition



Separation

Negligence

find labeling of nodes in phylogenetic tree minimizing the number of incongruences

Parsimony criteria

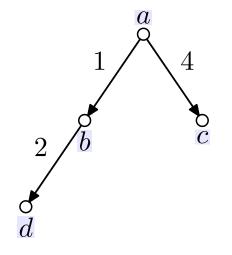
Given two rooted trees H_h and G_g with a labeling $l : G \to H$ and a weight function d on H_h , the *arc cost* and the *bag cost* of l are defined as follows:

$$\operatorname{arccost}(H_h, G_g, l) := \sum_{\langle u, v \rangle \in A(G_g)} d(l(u), l(v)),$$
$$\operatorname{bagcost}(H_h, G_g, l) := \sum_{v \in V(H)} \text{size of the bag-set of } v.$$

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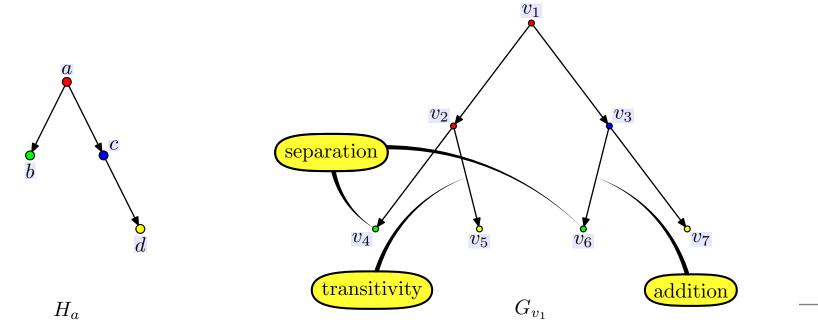
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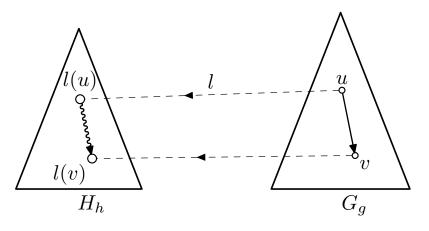
- **Relax-minor:** Given two rooted trees H_h and G_g with a leaf labeling p, we say that H_h is a *relax-minor* of G_g with respect to p if there exists a p-constrained labeling function $l: G \to H$ satisfying the following two conditions:
 - each evolutionary transition has a realization (no negligence); and
 - if u is an ancestor of v, then l(v) cannot be a proper ancestor of l(u) (no inversion).

Relax-minor:

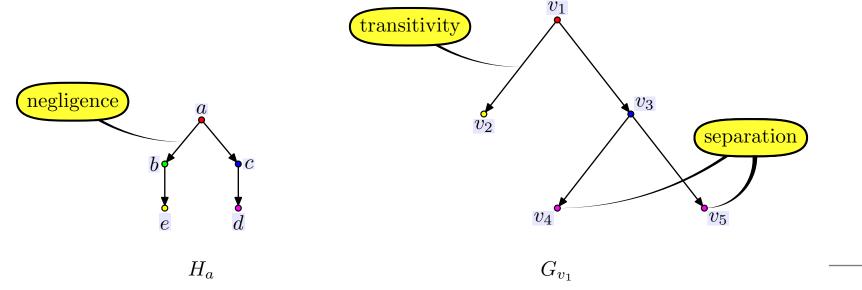
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- **Example** of other incongruences:



- Given two rooted trees H_h and G_g with a leaf labeling p, we say that H_h is a *pseudo-minor* of G_g with respect to p if there exists a p-constrained labeling function $l: G \to H$ such that
 - for every evolutionary step $\langle u, v \rangle$, l(u) is an ancestor of l(v) (no addition and inversion).



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- **Example** of other incongruences:



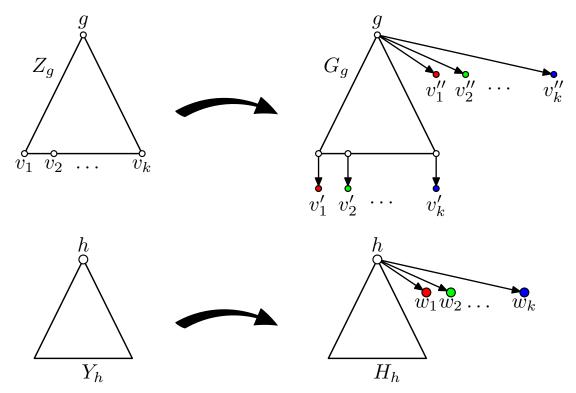
summary:

	rooted minor	relax-minor	pseudo-minor
inversion	Ν	Ν	Ν
transitivity	Ν	Y	Y
addition	Ν	Y	Ν
separation	Ν	Y	Y
negligence	Ν	Ν	Y

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 - Open problems.
 - Is it possible to solve Problem 2 in polynomial time when p is a complete leaf labeling?
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