Level-k Phylogenetic Networks

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• Combine a set of small trees (triplets) into a single network that is as simple as possible



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Phylogenetic Networks





Phylogenetic Networks





How Simple is a Network?

Number of reticulations: total number of reticulation vertices (indegree two vertices)

Level: maximum number of reticulation vertices in a biconnected component

- Level-0 networks are trees
- Level-1 networks are galled trees



Example: level-2 network with 4 reticulations



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 A triplet set is *dense* if for each combination of three leaves it contains at least one of the three possible triplets



Triplet Consistency











These two triplets can be combined into a tree







three triplets







But this level-1 networks is.

A level-(n-1) network consistent with **any** triplet set on n leaves



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consistent with a maximum	NP-hard for all k
number of input triplets	



consistent with a maximum number of input triplets	NP-hard for all k
consistent with all input triplets	NP-hard for all k > 0

consistent with a maximum number of input triplets	NP-hard for all k
consistent with all input triplets	NP-hard for all k > 0
consistent with a maximum number of input triplets from a dense triplet set	NP-hard for all k



Approach for proving NP-hardness for all k

• For the consistency problem we generalise the reduction for level-1.

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- For the consistency problem we generalise the reduction for level-1.
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Approach for proving NP-hardness for all k

- For the consistency problem we generalise the reduction for level-1.
- For the maximisation problem we generalise the reduction for level-0, and add triplets to make the instance dense.
- To be able to generalise the different reductions we show that the following network is "unique"...

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Let T be the set of triplets consistent with this network.

Then this is the only network consistent with T.





This network is even "unique" for the maximisation problem



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 A triplet set is *dense* if for each combination of three leaves it contains at least one of the three possible triplets



consistent with all input triplets from a dense triplet set	polynomial-time for $k = 1$ and $k = 2$; open for $k > 2$



consistent with all input triplets from a dense triplet set	polynomial-time for $k = 1$ and $k = 2$; open for $k > 2$
and containing a minimum number of reticulations	polynomial-time for $k = 1$ and $k = 2$; open for $k > 2$



consistent with all input triplets from a dense triplet set	polynomial-time for $k = 1$ and $k = 2$; open for $k > 2$
and containing a minimum number of reticulations	polynomial-time for k = 1 and k = 2; open for k > 2
consistent with precisely the input triplets	polynomial-time for all k

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Given: general set of triplets **Construct:** level-k network consistent with a maximum number of input triplets

- Bang Ye Wu gave an O(3ⁿ) algorithm for level-0
- We give an O(4ⁿ) algorithm for level-1

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Given: general set of triplets **Construct:** level-k network consistent with a maximum number of input triplets

- Bang Ye Wu gave an O(3ⁿ) algorithm for level-0
- We give an O(4ⁿ) algorithm for level-1
- Are faster algorithms possible?
- Is anything better than a 3-approximation possible for level-0?



The Exact Algorithm

- Loop through all subsets of the leaves from small to large
- Compute an optimal network for these leaves based on previously computed optimal networks for smaller leaf-sets



Case 1: root is not in a cycle





Case 2: root is in a cycle

Try each tripartition of L into X, Y and Z



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Problem

What happens with triplets that cross the tripartition?





Solution

- Combine networks for X U Z and Y U Z.
- Where Z has to be below an "n.c.r.-arc".
- Definition: an arc is an *n.c.r.-arc* if it is not reachable from any vertex in a cycle.



Solution







- Leo van Iersel and Steven Kelk, Constructing the Simplest Possible Phylogenetic Network from Triplets, submitted.
- Leo van Iersel, Steven Kelk and Matthias Mnich, Uniqueness, Intractability and Exact Algorithms: Reflections on Level-k Phylogenetic Networks, submitted.
- Leo van Iersel, Judith Keijsper, Steven Kelk, Leen Stougie, Ferry Hagen and Teun Boekhout, Constructing Level-2 Phylogenetic Networks from Triplets, in proceedings of RECOMB 2008.
- All papers (and some implementations): http://www.win.tue.nl/~liersel