## Reconstructing the Tree of Life (Fitting Distances by Tree Metrics)

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## Abstract -

We consider the numerical taxonomy problem of fitting an  $S \times S$  distance matrix D with a tree metric T. Here T is a weighted tree spanning S where the path lengths in T induce a metric on S. If there is a tree metric matching D exactly, then it is easily found. If there is no exact match, then for some k, we want to minimize the  $L_k$  norm of the errors, that is, pick T so as to minimize

$$||D - T||_k = \left(\sum_{i,j \in S} |D(i,j) - T(i,j)|^k\right)^{1/k}.$$

This problem was raised in biology in the 1960s for k = 1, 2. The biological interpretation is that T represents a possible evolution behind the species in S matching some measured distances in D. Sometimes, it is required that T is an ultrametric, meaning that all species are at the same distance from the root.

An evolutionary tree induces a hierarchical classification of species and this is not just tied to biology. Medicine, ecology and linguistics are just some of the fields where this concept appears, and it is even an integral part of machine learning and data science. Fundamentally, if we can approximate distances with a tree, then they are much easier to reason about: many questions that are NP-hard for general metrics can be answered in linear time on tree metrics. In fact, humans have appreciated hierarchical classifications at least since Plato and Aristotle (350 BC).

The numerical taxonomy problem is important in practice and many heuristics have been proposed. In this talk we will review the basic algorithmic theory, results and techniques, for the problem, including the most recent result from FOCS'21 [3]. They paint a varied landscape with big differences between different moments, and with some very nice open problems remaining.

- At STOC'93, Farach, Kannan, and Warnow [4] proved that under  $L_{\infty}$ , we can find the optimal ultrametric. Almost all other variants of the problem are APX-hard.
- At SODA'96, Agarwala, Bafna, Farach, Paterson, and Thorup [1] showed that for any norm  $L_k, k \geq 1$ , if the best ultrametric can be  $\alpha$ -approximated, then the best tree metric can be  $3\alpha$ -approximated. In particular, this implied a 3-approximation for tree metrics under  $L_{\infty}$ .
- At FOCS'05, Ailon and Charikar [2] showed that for any  $L_k$ ,  $k \ge 1$ , we can get an approximation factor of  $O(((\log n)(\log \log n))^{1/k})$  for both tree and ultrametrics. Their paper was focused on the  $L_1$  norm, and they wrote "Determining whether an O(1) approximation can be obtained is a fascinating question".
- At FOCS'21, Cohen-Addad, Das, Kipouridis, Parotsidis, and Thorup [3] showed that indeed a constant factor is possible for  $L_1$  for both tree and ultrametrics. This uses the special structure of  $L_1$  in relation to hierarchies.
- The status of  $L_k$  is wide open for  $1 < k < \infty$ . All we know is that the approximation factor is between  $\Omega(1)$  and  $O((\log n)(\log \log n))$ .

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