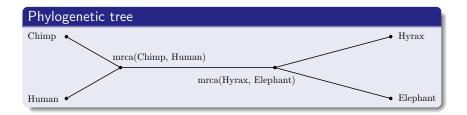
Next generation phylogenetic inference @Evolution2016: Nearest neighbors of phylogenetic time-trees

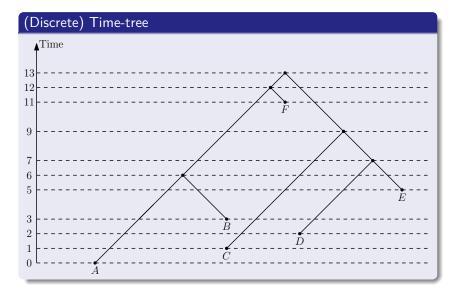
### Alex Gavryushkin (joint work with Chris Whidden and Erick Matsen)

June 20, 2016

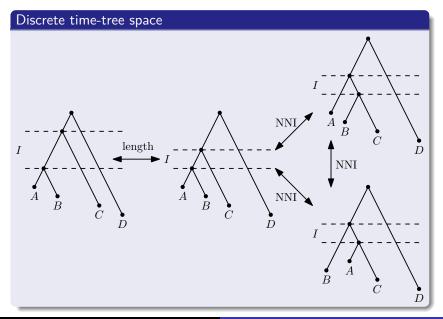


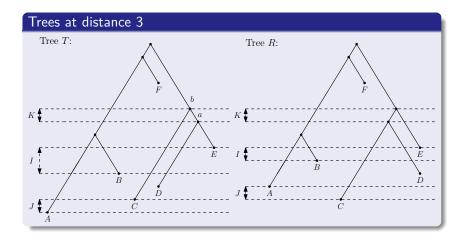




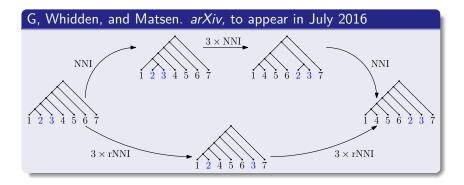


### Main definition

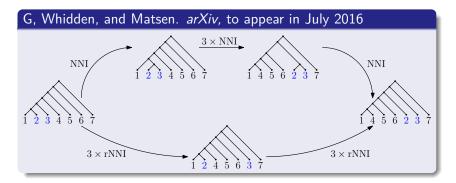




## Looks promising



# Looks promising



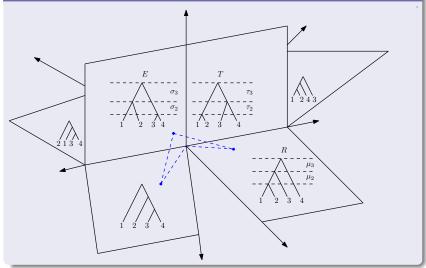
#### And challenging!

What it the complexity of this graph?

- Over 25 years to solve for NNI
- Over 7 erroneous papers published

## Promising indeed





### Mathematical insight

- Geometry of discrete time-trees
  - Diameter
  - Micro- and macro-geometry
- Random walking
  - Bottle necks
  - Curvature

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### Surprise!

A better bound for the size of r-neighborhoods in NNI.

## Take home message

#### Our results

- Time-trees and classical phylogenetic trees have different geometric and algorithmic properties.
- Often, geometric and algorithmic results for classical trees do not scale to time-trees.
- Natural and efficient data structures

## Take home message

#### Our results

- Time-trees and classical phylogenetic trees have different geometric and algorithmic properties.
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- Natural and efficient data structures

#### Future directions

- Applications
  - MCMC proposals, local tree rearrangements
  - Convergence
  - Valleys and terraces
- Math/computing:
  - Challenging problems that matter
  - Connections to other areas of math
  - Amenable to various search algorithms



Alex Gavryushkin and Alexei Drummond The space of ultrametric phylogenetic trees Journal of Theoretical Biology, Vol. 402, 197–208, 2016



Alex Gavryushkin, Chris Whidden, and Frederick A. Matsen IV Combinatorics of discrete time-trees: algorithmic insights and open problems To appear on the arXiv, July 2016



https://github.com/gavruskin/tauGeodesic

https://github.com/gavruskin/tTauCurvature