## Preview: PACE 2026

Alexander Leonhardt<sup>1</sup>, Manuel Penschuck<sup>2</sup>, Mathias Weller<sup>3</sup>

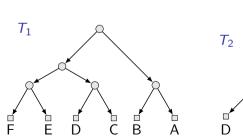
<sup>1</sup>Goethe-Universität Frankfurt, Germany <sup>2</sup>University of Southern Denmark, Odense, Denmark <sup>3</sup>CNRS, Université Gustave Eiffel, Paris, France https://pacechallenge.org/

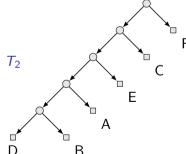
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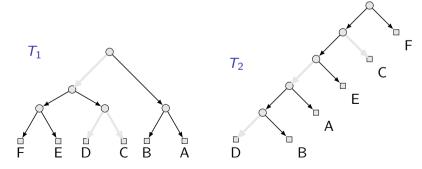
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- An agreement forest of t phylogenetic trees  $T_1, T_2, \ldots, T_t$  is any forest of phylogenetic trees that can be obtained from each  $T_i$  by removing directed edges (+ "cleanup")





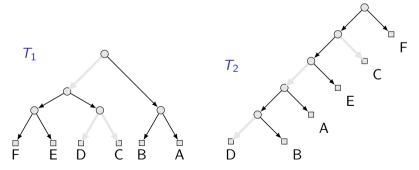
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**Note**: No other parameterization explored!

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#### **Generated Data**

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mostly follows previous PACE-instances

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good luck and an enjoyable competition –