

Discovering the Repeatome of Five Species Belonging to the Asteraceae Family: A Computational Study

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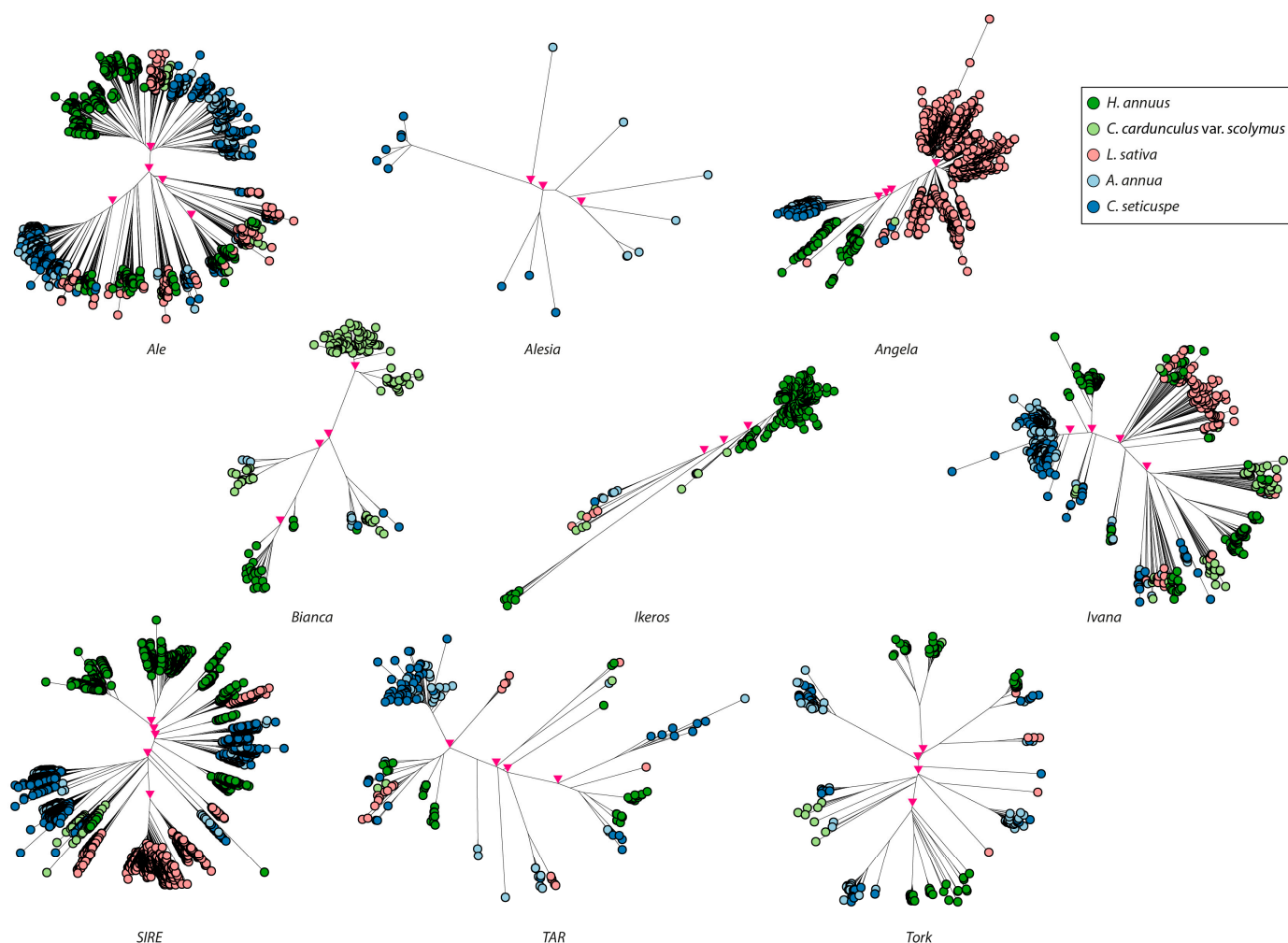


Figure S1. Phylogenetic trees of all RT sequences of *Copia* elements retrieved in the five Asteraceae species, separated according to the lineage. The main nodes (bootstrap values > 0.6) are marked with pink triangles. RT = reverse transcriptase.

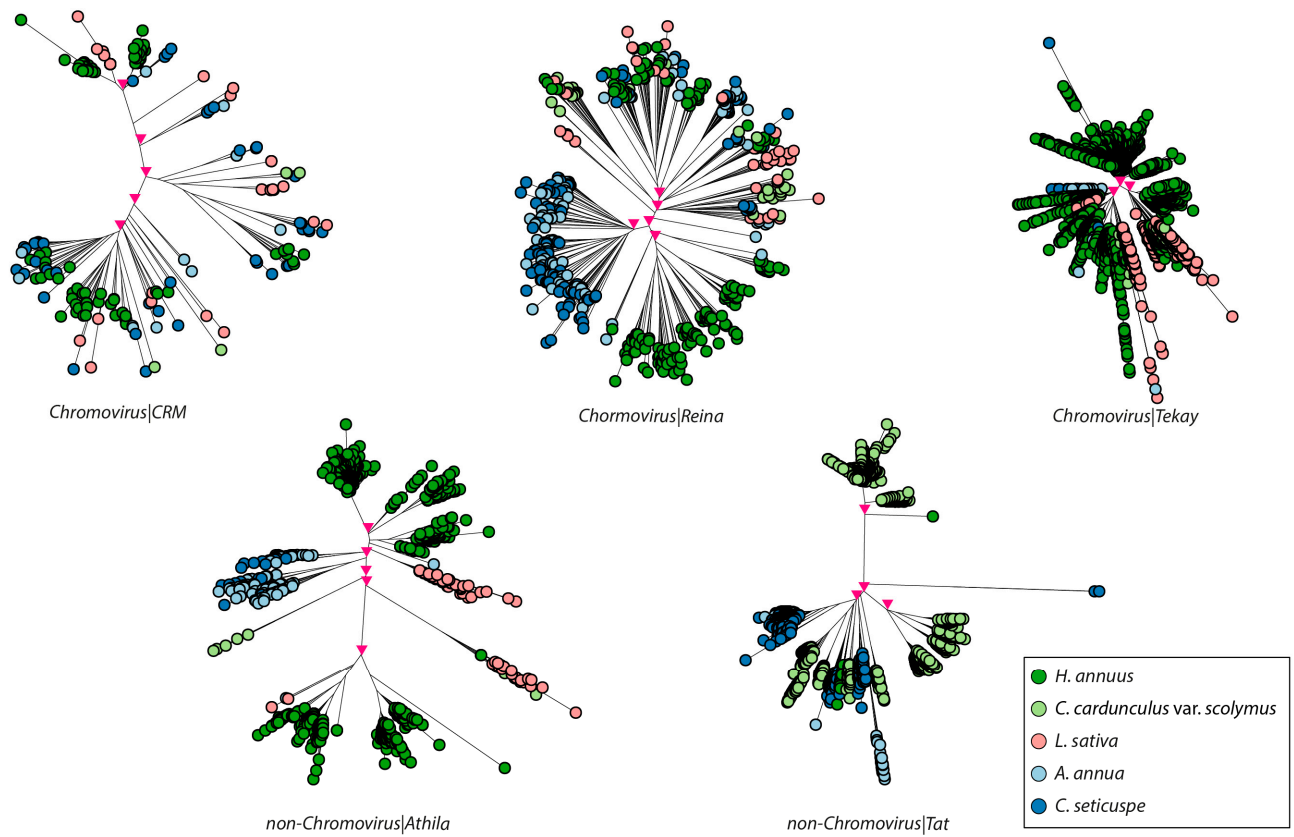


Figure S2. Phylogenetic trees of all RT sequences of *Gypsy* elements retrieved in the five Asteraceae species, separated according to the lineage. The main nodes (bootstrap values > 0.6) are marked with pink triangles. RT = reverse transcriptase.