

Integrating phylogenies with chronology to assemble the Tree of Life Supplementary Material

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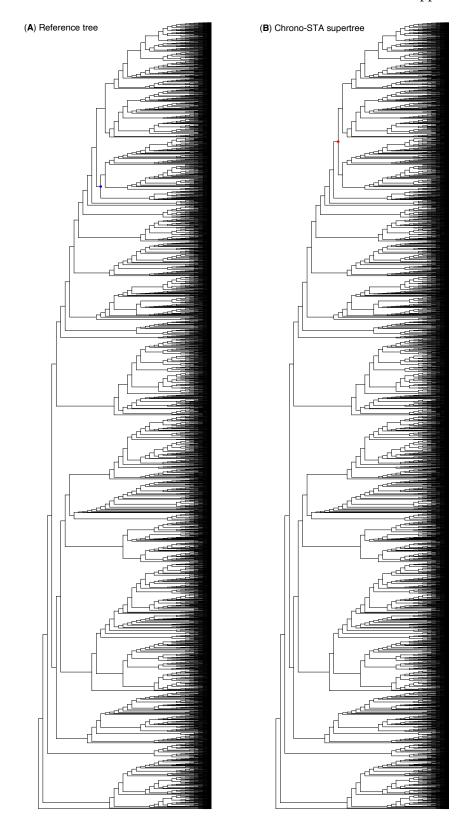
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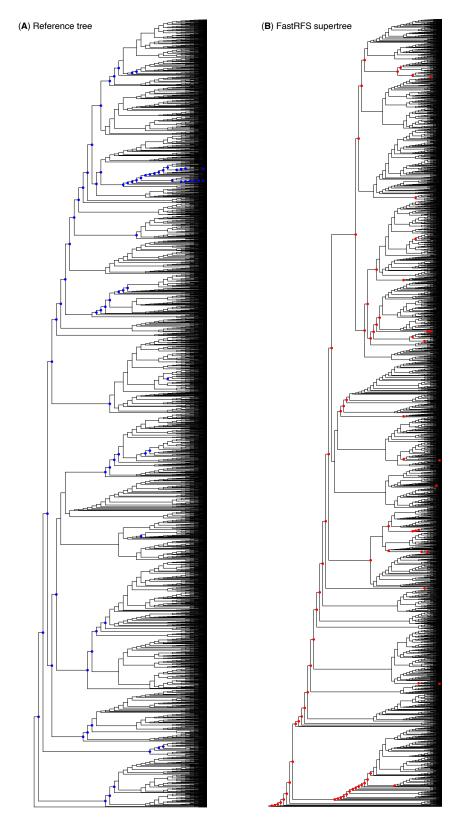
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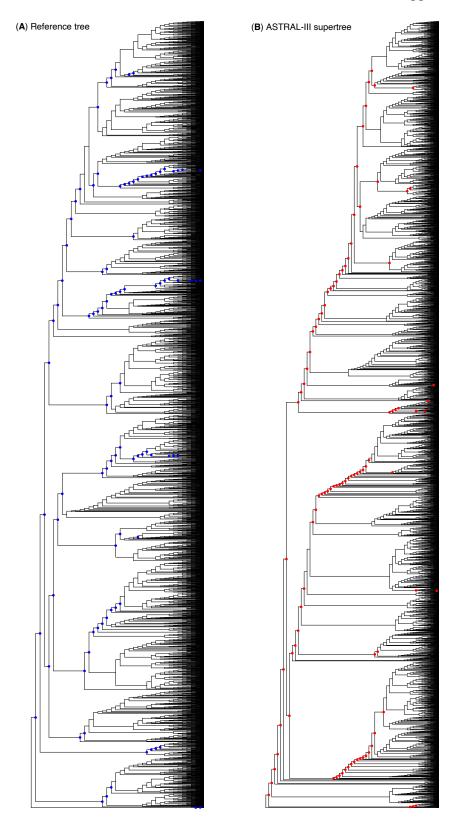
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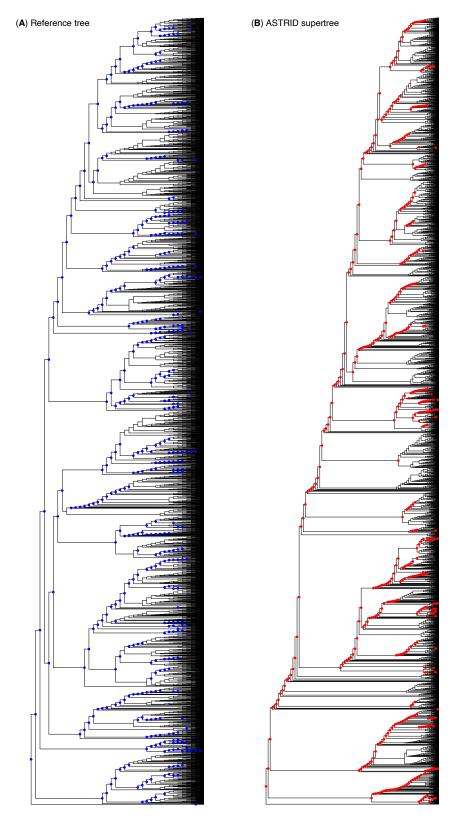
Supplementary Figure 1. Comparison of (A) the mammal timetree (Álvarez-Carretero *et al.*, 2022). consisting of 4,705 species (reference tree) and (B) the Chrono-STA supertree. Red dots represent clades absent in (A), and blue dots represent clades absent in (B).



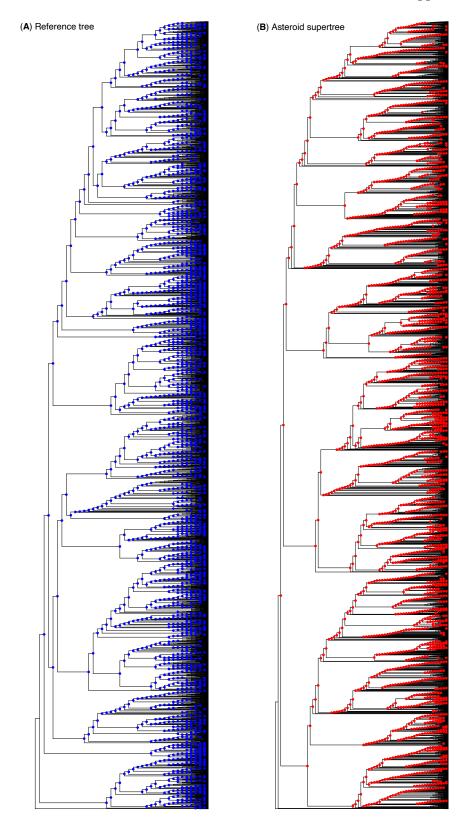
Supplementary Figure 2. Comparison of (A) the mammal timetree (Álvarez-Carretero *et al.*, 2022). consisting of 4,705 species (reference tree) and (B) the FastRFS supertree. Red dots represent clades absent in (A), and blue dots represent clades absent in (B).



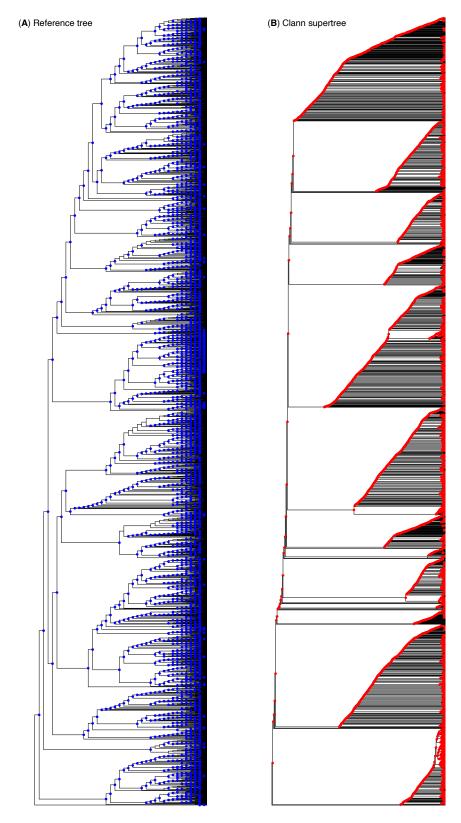
Supplementary Figure 3. Comparison of (A) the mammal timetree (Álvarez-Carretero *et al.*, 2022). consisting of 4,705 species (reference tree) and (B) the ASTRAL-III supertree. Red dots represent clades absent in (A), and blue dots represent clades absent in (B).



Supplementary Figure 4. Comparison of (A) the mammal timetree (Álvarez-Carretero *et al.*, 2022). consisting of 4,705 species (reference tree) and (B) the ASTRID supertree. Red dots represent clades absent in (A), and blue dots represent clades absent in (B).



Supplementary Figure 5. Comparison of (A) the mammal timetree (Álvarez-Carretero *et al.*, 2022). consisting of 4,705 species (reference tree) and (B) the Asteroid supertree. Red dots represent clades absent in (A), and blue dots represent clades absent in (B).



Supplementary Figure 6. Comparison of (A) the mammal timetree (Álvarez-Carretero *et al.*, 2022). consisting of 4,705 species (reference tree) and (B) the Clann supertree. Red dots represent clades absent in (A), and blue dots represent clades absent in (B).