

CORRECTION

Correction: Phylogenetic Relationships of American Willows (*Salix* L., Salicaceae)

Aurélien Lauron-Moreau, Frédéric E. Pitre, George W. Argus, Michel Labrecque, Luc Brouillet

Previous revision versions of [Fig 2](#), [Fig 3](#), and [S1 Fig](#) were incorrectly published. Please view the correct [Fig 2](#), [Fig 3](#), and [S1 Fig](#) here.



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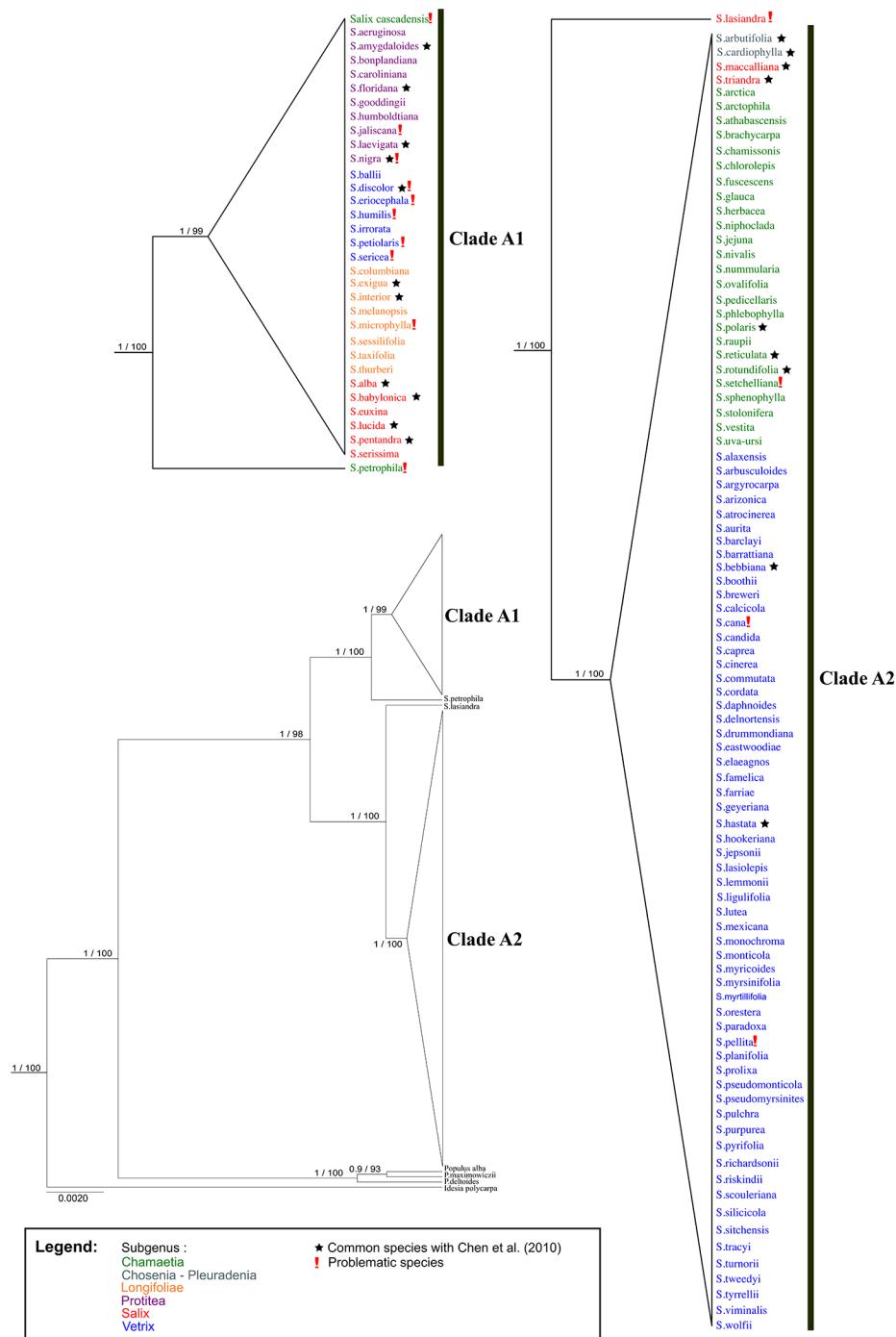


Fig 2. BEAST gene tree of *matK* and *rbcL*. Branch support is Bayesian posterior probabilities and ML bootstrap values; subgenera are identified using colors; *Idesia* and *Populus* are outgroups.

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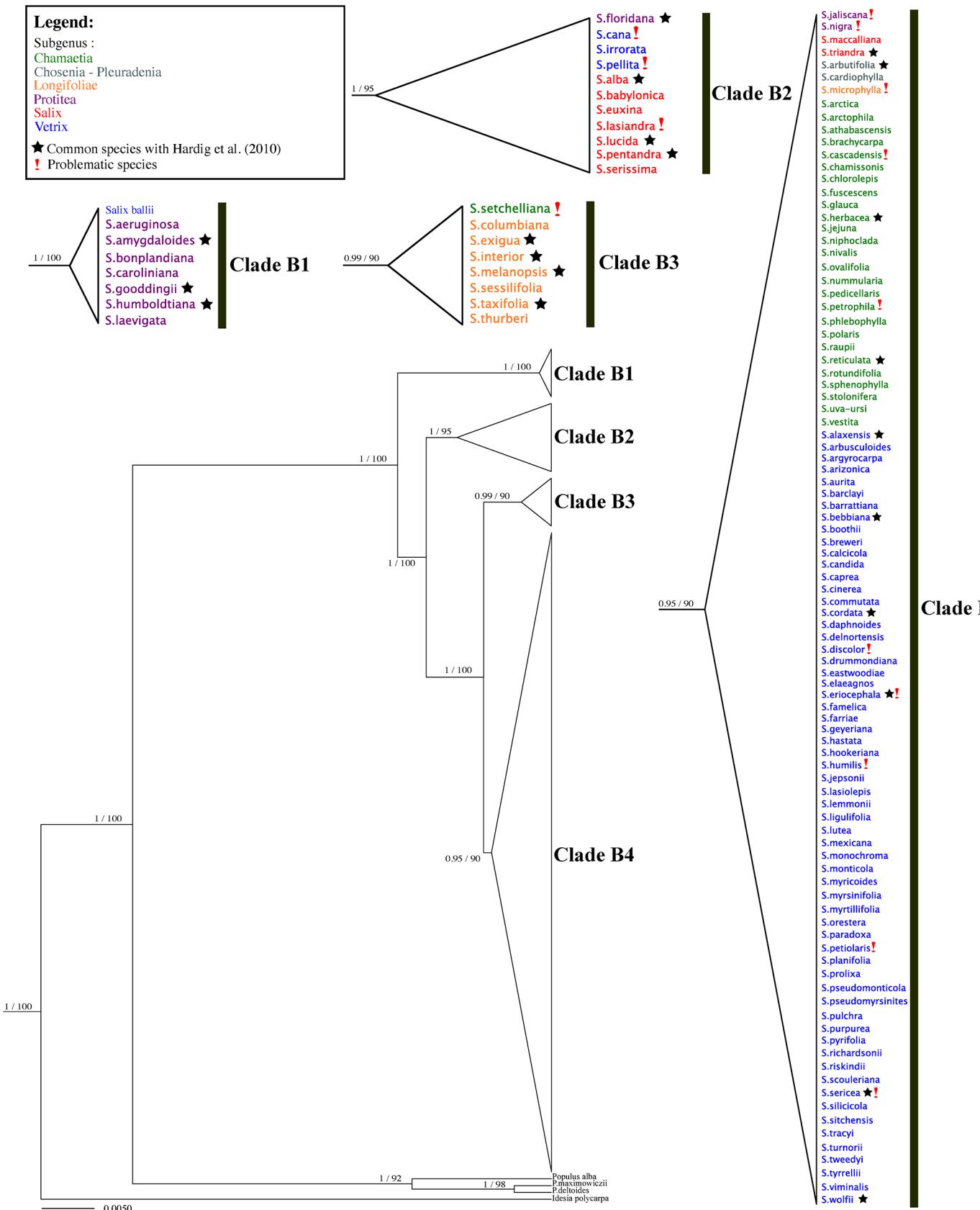


Fig 3. BEAST gene tree of ITS. Branch support is Bayesian posterior probabilities and ML bootstrap values; subgenera are identified using colors; *Idesia* and *Populus* are outgroups.

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Supporting Information

S1 Fig. BEAST species tree generated with *ITS*, *matK* and *rbcL*, constrained to fit the subgenera of Argus (2010). Branch support is Bayesian posterior probabilities and ML bootstrap values.
(EPS)

Reference

1. Lauron-Moreau A, Pitre FE, Argus GW, Labrecque M, Brouillet L (2015) Phylogenetic Relationships of American Willows (*Salix* L., Salicaceae). PLoS ONE 10(4): e0121965. doi:[10.1371/journal.pone.0121965](https://doi.org/10.1371/journal.pone.0121965) PMID: [25880993](#)