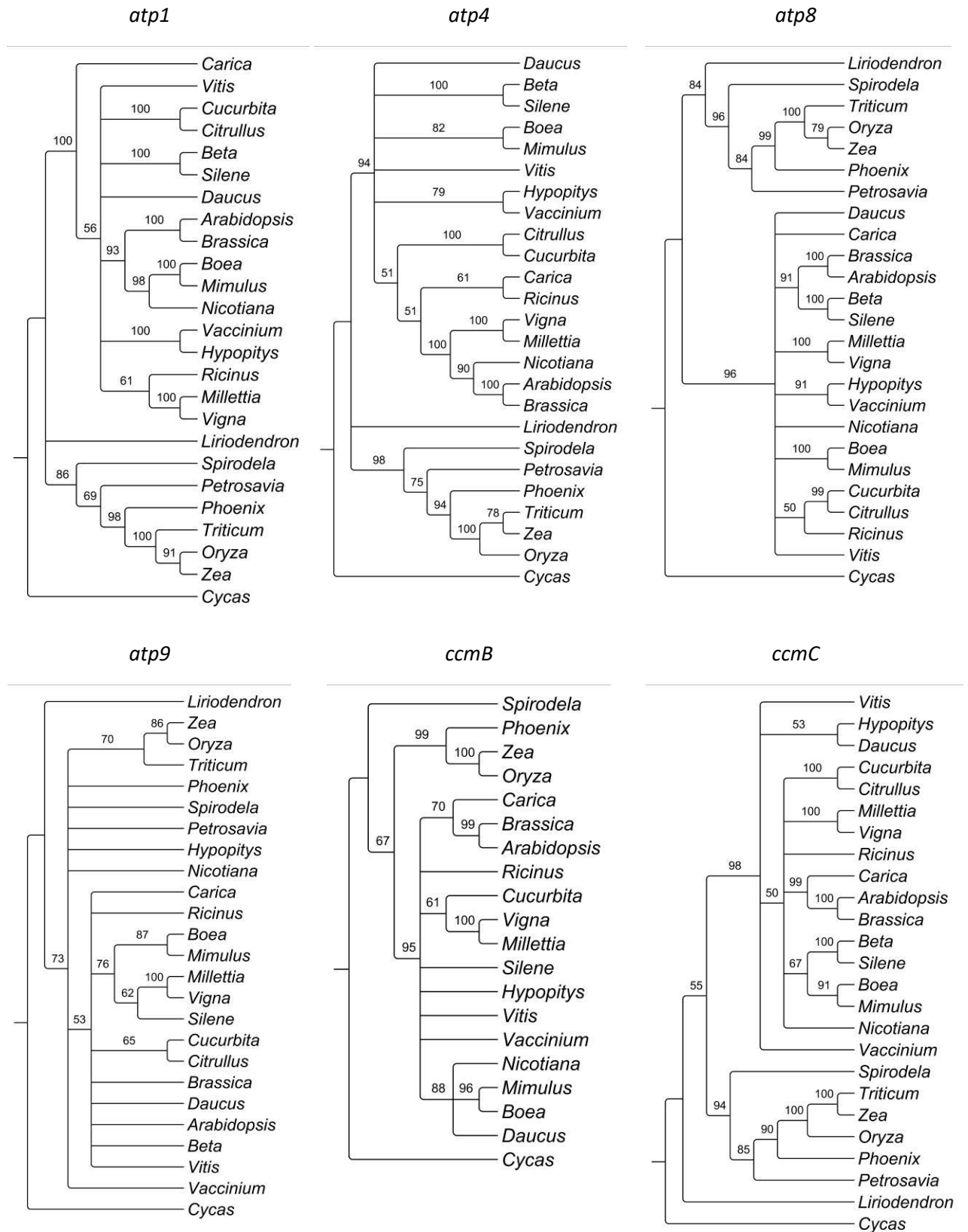
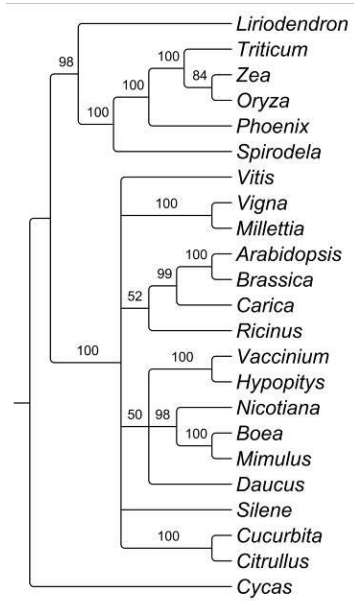


Supplementary figure 5.

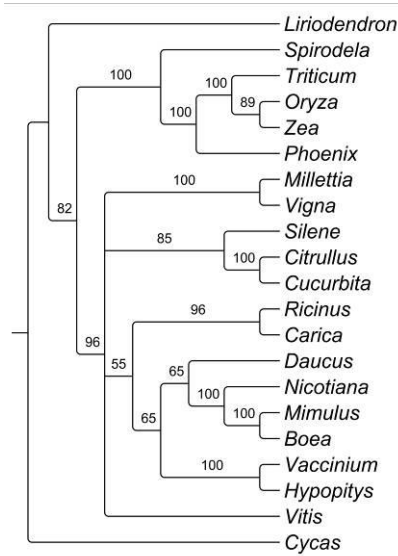
Phylogenetic trees inferred from ML analysis of single mitochondrial genes. Protein coding genes that were annotated in mitochondrial genomes of *Hypopitys monotropa*, *Vaccinium macrocarpon* and all main evolutionary lineages of flowering plants are included in the analysis.



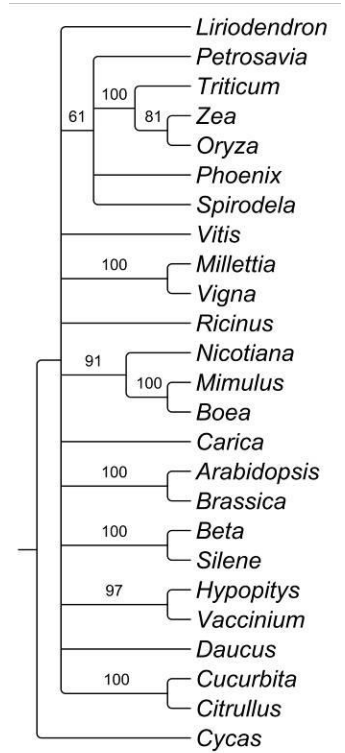
ccmFc



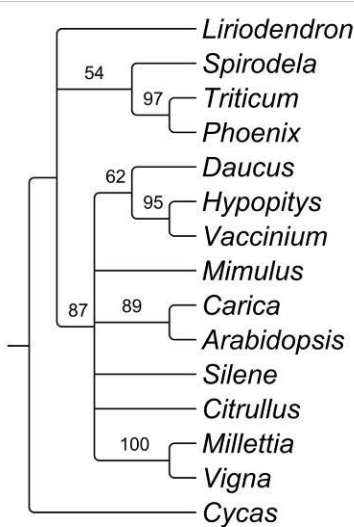
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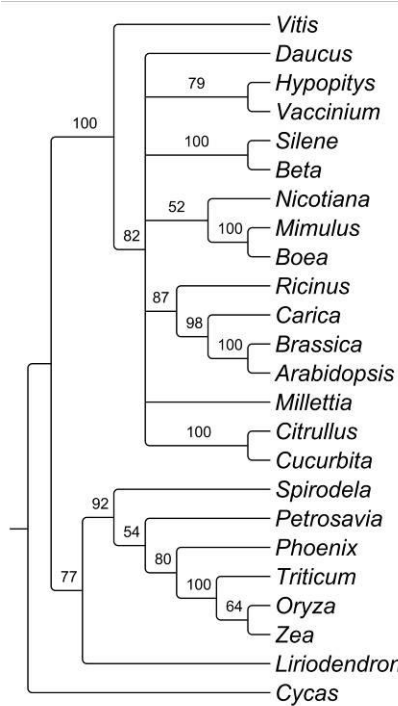
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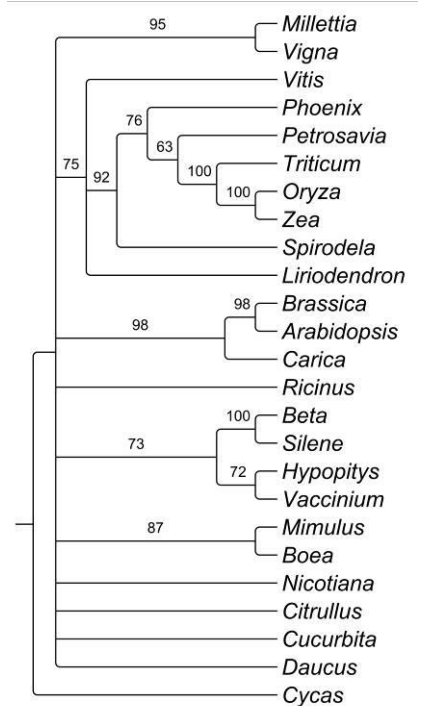
cox1



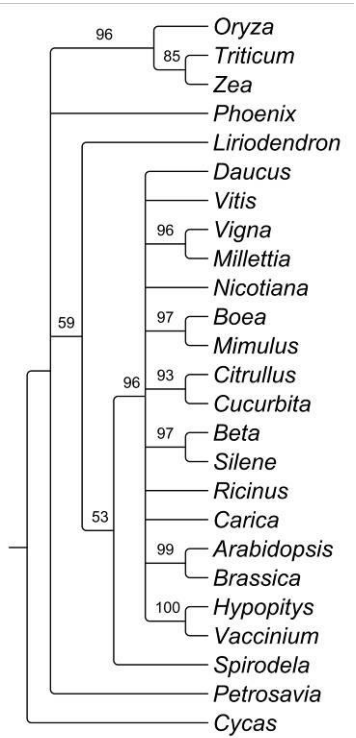
cox2



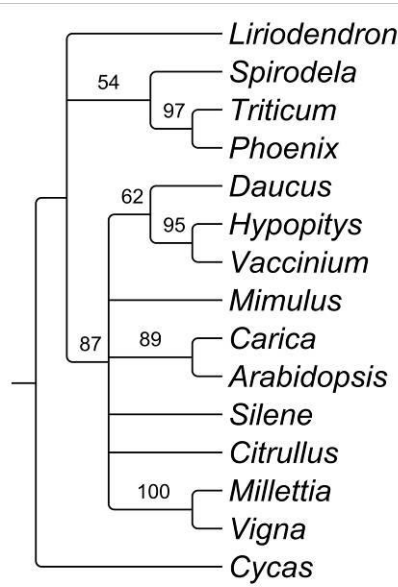
cox3



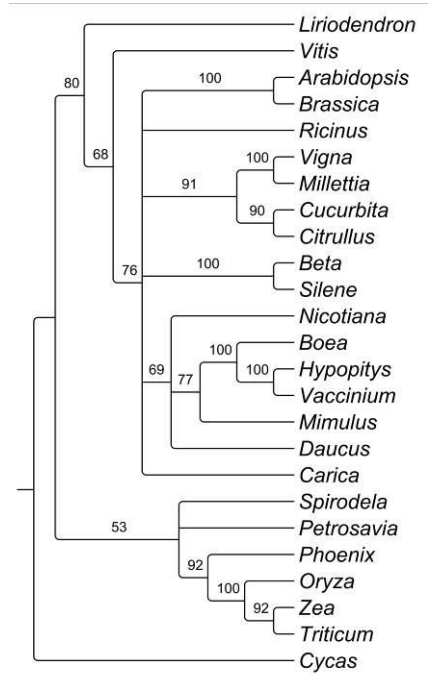
matR



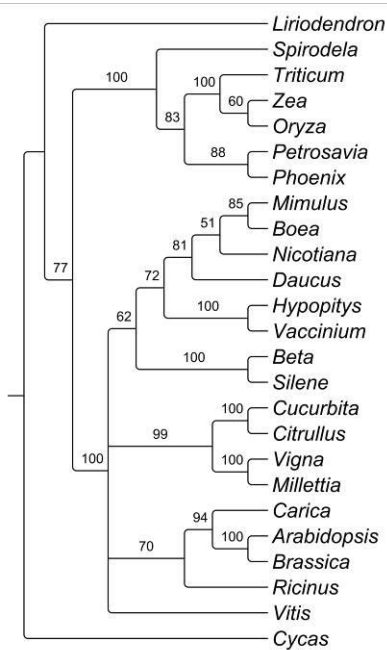
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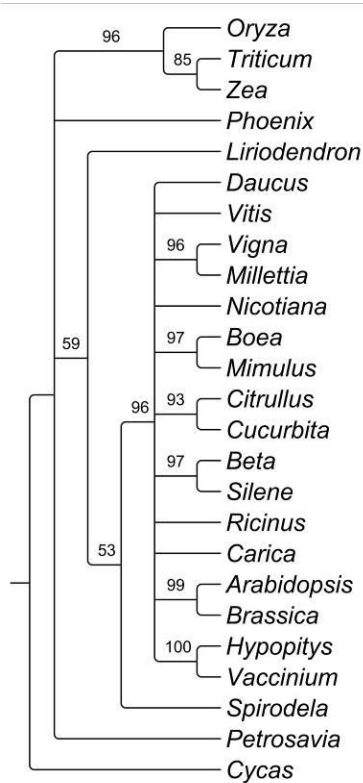
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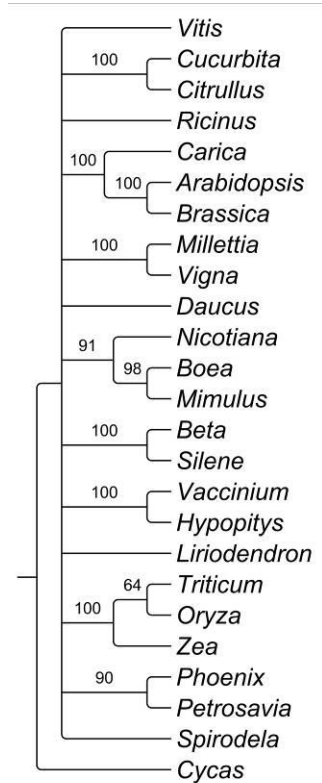
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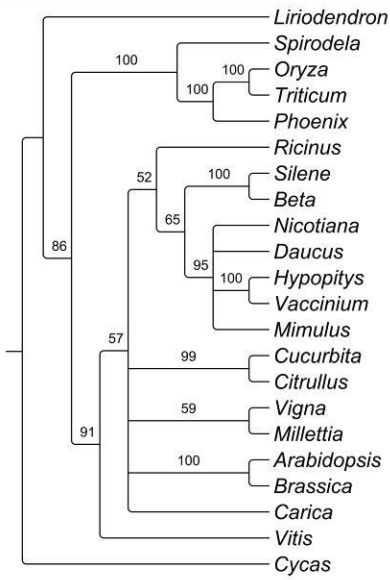


nad4

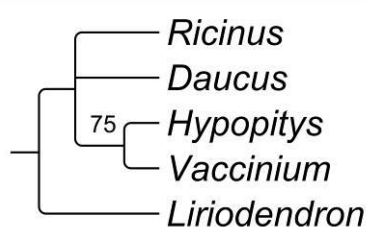




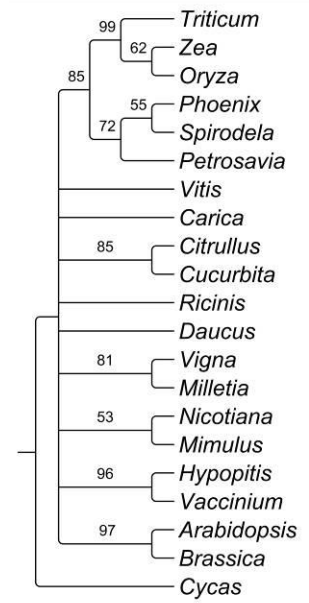
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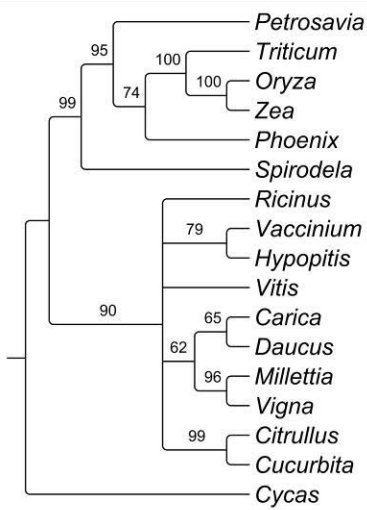
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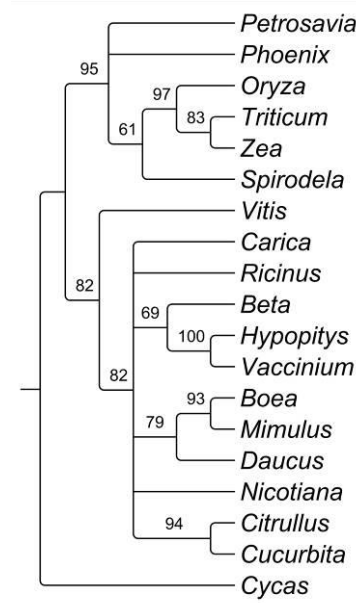
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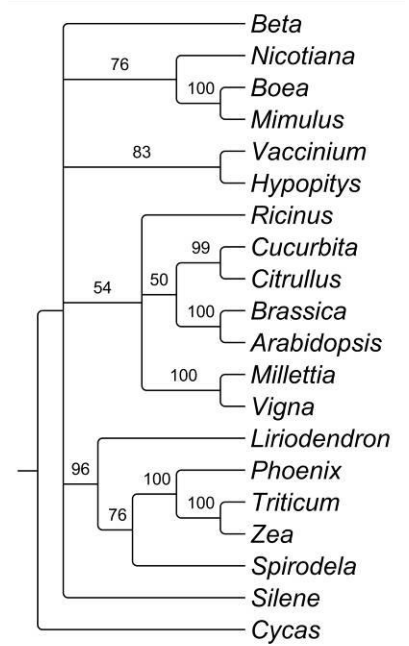
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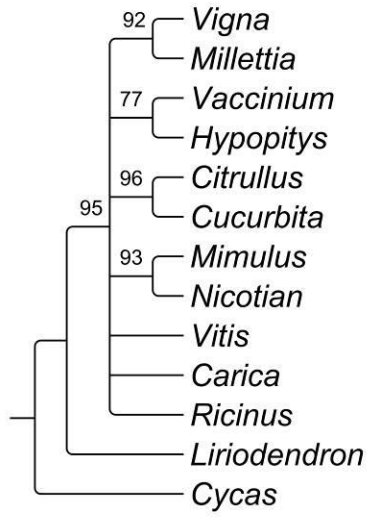
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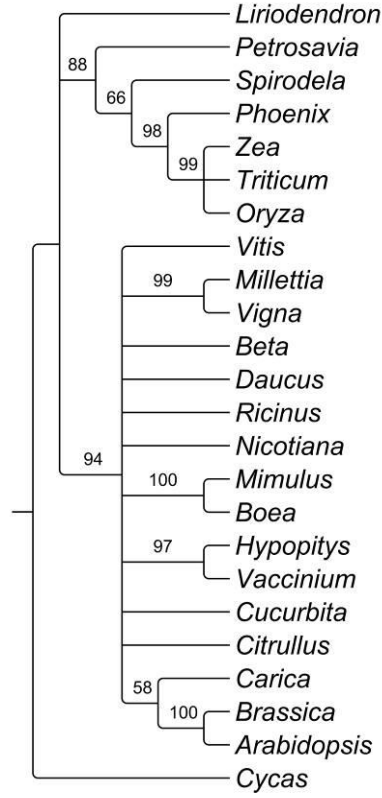
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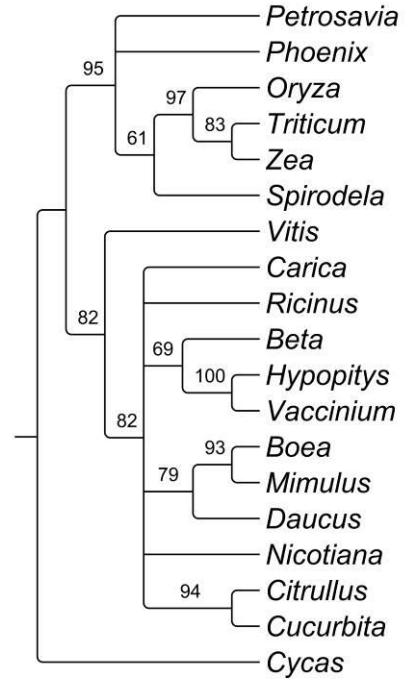
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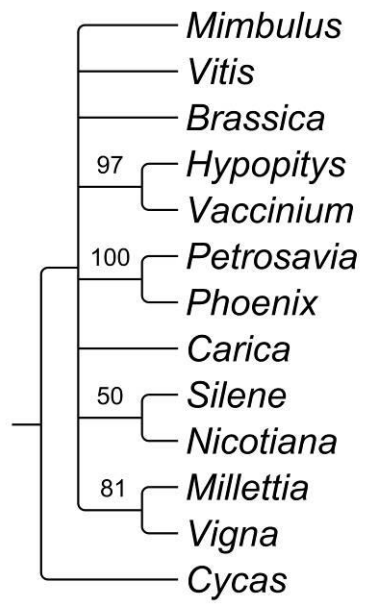
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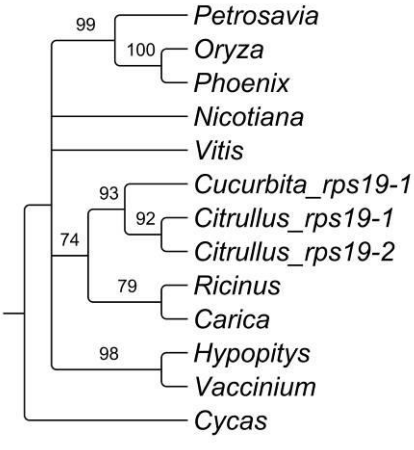
rps13



rps14



rps19



sdh4

