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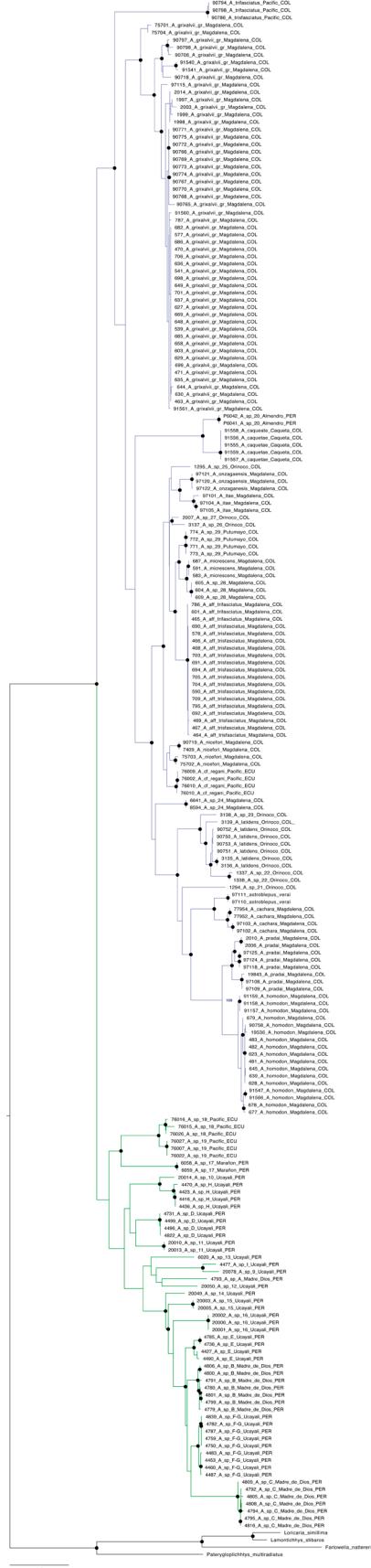
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SUPPLEMENTARY MATERIAL FIGURE S3 |

Maximum likelihood tree inferred by RAxML from the concatenated dataset of mitochondrial genes. Bootstrap values >95% as black circles. Tip names include tissue number and final identification following S1.



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