

TABLE S1 Mantel modeling tests performed with asymmetric matrices: Between Recent migration rates (RMR) and dispersal inferred by Oceanographic modeling. Pearson correlation values (r). Significance values (p), showed in bold for $p < 0.05$. Molecular marker: SNP or Msat (microsatellite) and Reference of data source are shown. ¹ Atlantic Ocean Basin; ² Pacific Ocean Basin. The information was not shown when at least one of the matrices presented all values equal to zero.

| a) <i>Avicennia germinans</i> | | |
|-----------------------------------|--------|--|
| Mantel RMR | | |
| Oceanography r (p) | Marker | Reference |
| -0.13 (0.84) | Msat | Hodel et al., 2016 |
| -0.17 (0.89) | Msat | Kennedy et al., 2020 |
| 0.97 (<10⁻⁵) | Msat | Mori et al., 2015 |
| -0.13 (0.85) | Msat | Ochoa-Zavala et al., 2019 ¹ |
| 0.24 (0.78) | SNP | Cruz et al., 2020 |
| b) <i>Laguncularia racemosa</i> | | |
| -0.17 (0.41) | SNP | Hodel et al., 2018 |
| c) <i>Rhizophora mangle</i> | | |
| 0.14 (0.50) | Msat | Cisneros de la Cruz et al., 2018 |
| -0.30 (0.50) | Msat | Francisco et al., 2018 |
| 0.95 (0.03) | Msat | Hodel et al., 2016 |
| -0.11 (0.56) | Msat | Hodel et al., 2017 |
| 0.23 (0.45) | Msat | Kennedy et al., 2016 |
| -0.17 (0.73) | Msat | Kennedy et al., 2017 |
| 0.99 (<10⁻⁵) | Msat | Takayama et al., 2013 ¹ |
| 0.42 (0.14) | Msat | Takayama et al., 2013 ² |
| -0.13 (0.88) | SNP | Hodel et al., 2017 |
| 0.13 (0.42) | SNP | Hodel et al., 2018 |
| -0.25 (0.50) | SNP | Madeira et al., 2023 |

References

- Cisneros-de la Cruz, D. J., Martínez-Castillo, J., Herrera-Silveira, J., Yáñez-Espinosa, L., Ortiz-García, M., Us-Santamaria, R., & Andrade, J. L. (2018). Short-distance barriers affect genetic variability of *Rhizophora mangle* L. in the Yucatan Peninsula. *Ecology and Evolution*, 8(22), 11083–11099.
<https://doi.org/10.1002/ece3.4575>
- Cruz, M. V., Mori, G. M., Oh, D. H., Dassanayake, M., Zucchi, M. I., Oliveira, R. S., & Souza, A. P. de. (2020). Molecular responses to freshwater limitation in the

mangrove tree *Avicennia germinans* (Acanthaceae). *Molecular Ecology*, 29(2), 344–362. <https://doi.org/10.1111/mec.15330>

Francisco, P. M., Mori, G. M., Alves, F. M., Tambarussi, E. V., & de Souza, A. P. (2018). Population genetic structure, introgression, and hybridization in the genus *Rhizophora* along the Brazilian coast. *Ecology and Evolution*, 8(6), 3491–3504. <https://doi.org/10.1002/ece3.3900>

Hodel, R. G. J., Chen, S., Payton, A. C., Mcdaniel, S. F., Soltis, P., & Soltis, D. E. (2017). Adding loci improves phylogeographic resolution in red mangroves despite increased missing data : comparing microsatellites and RAD-Seq and investigating loci filtering. *Scientific Reports*, November, 1–14. <https://doi.org/10.1038/s41598-017-16810-7>

Hodel, R. G. J., De Souza Cortez, M. B., Soltis, P. S., & Soltis, D. E. (2016). Comparative phylogeography of black mangroves (*Avicennia germinans*) and red mangroves (*Rhizophora mangle*) in Florida: Testing the maritime discontinuity in coastal plants. *American Journal of Botany*, 103(4), 730–739. <https://doi.org/10.3732/ajb.1500260>

Hodel, R. G. J., Knowles, L. L., McDaniel, S. F., Payton, A. C., Dunaway, J. F., Soltis, P. S., & Soltis, D. E. (2018). Terrestrial species adapted to sea dispersal: Differences in propagule dispersal of two Caribbean mangroves. *Molecular Ecology*, 27(22), 4612–4626. <https://doi.org/10.1111/mec.14894>

Kennedy, J. P., Garavelli, L., Truelove, N. K., Devlin, D. J., Box, S. J., Ch, L. M., & Feller, I. C. (2017). *Contrasting genetic effects of red mangrove (Rhizophora mangle L .) range expansion along West and East Florida*. 335–347. <https://doi.org/10.1111/jbi.12813>

Kennedy, J. P., Pil, M. W., Proffi, C. E., Boeger, W. A., Stanford, A. M., & Devlin, D. J. (2016). *Postglacial expansion pathways of red mangrove , Rhizophora mangle , in the Caribbean Basin and Florida 1*. 103(2), 260–276. <https://doi.org/10.3732/ajb.1500183>

Kennedy, J. P., Preziosi, R. F., Rowntree, J. K., & Feller, I. C. (2020). Is the central-marginal hypothesis a general rule? Evidence from three distributions of an expanding mangrove species, *Avicennia germinans* (L.) L. *Molecular Ecology*,

29(4), 704–719. <https://doi.org/10.1111/mec.15365>

- Madeira, A. G., Tsuda, Y., Nagano, Y., Iwasaki, T., Zucchi, M. I., Kajita, T., & Mori, G. M. (2023). The role of oceanic currents in the dispersal and connectivity of the mangrove *Rhizophora mangle* on the Southwest Atlantic region. *Molecular Ecology Resources*. <https://doi.org/10.1111/1755-0998.13807>
- Mori, G. M., Zucchi, M. I., & Souza, A. P. (2015). Multiple-geographic-scale genetic structure of two mangrove tree species: The roles of mating system, hybridization, limited dispersal and extrinsic factors. *PLoS ONE*, *10*(2), 1–23. <https://doi.org/10.1371/journal.pone.0118710>
- Ochoa-Zavala, M., Jaramillo-Correa, J. P., Piñero, D., Nettel-Hernanz, A., & Núñez-Farfán, J. (2019). Contrasting colonization patterns of black mangrove (*Avicennia germinans* (L.) L.) gene pools along the Mexican coasts. *Journal of Biogeography*, *46*(5), 884–898. <https://doi.org/10.1111/jbi.13536>
- Takayama, K., Tamura, M., Tateishi, Y., Webb, E. L., & Kajita, T. (2013). Strong genetic structure over the American continents and transoceanic dispersal in the mangrove genus *Rhizophora* (Rhizophoraceae) revealed by broad-scale nuclear and chloroplast DNA analysis. *American Journal of Botany*, *100*(6), 1191–1201. <https://doi.org/10.3732/ajb.1200567>