

Low genetic diversity and limited gene flow in a dominant
(*Rhizophora stylosa*) at its northern biogeographical limit
Sakishima islands of the Japanese archipelago as revealed by
analysis

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Development and Characterization of Polymorphic Microsatellite Markers (SSRs) for an Endemic Plant, <i>Pseudolarix amabilis</i> (Nelson) Rehd. (Pinaceae). <i>Molecules</i> , 2015, 20, 2685-2692.	1.7	14
2	Deciphering the taxonomical controversies of <i>Rhizophora</i> hybrids using AFLP, plastid and nuclear markers. <i>Aquatic Botany</i> , 2015, 125, 48-56.	0.8	6
3	Analysis of the mating system, reproductive characteristics, and spatial genetic structure in a natural mangrove tree (<i>Bruguiera gymnorrhiza</i>) population at its northern biogeographic limit in the southern Japanese archipelago. <i>Journal of Forest Research</i> , 2015, 20, 293-300.	0.7	10
4	Species distribution and introgressive hybridization of two <i>Avicennia</i> species from the Western Hemisphere unveiled by phylogeographic patterns. <i>BMC Evolutionary Biology</i> , 2015, 15, 61.	3.2	23
5	Development and characterization of microsatellite markers for <i>Ulmus chenmoui</i> (Ulmaceae), an endangered tree endemic to eastern China. <i>Genetics and Molecular Research</i> , 2016, 15, .	0.3	2
6	Comparative Analysis of the Pattern of Population Genetic Diversity in Three Indo-West Pacific <i>Rhizophora</i> Mangrove Species. <i>Frontiers in Plant Science</i> , 2016, 7, 1434.	1.7	45
7	Microsatellite markers for the critically endangered elm species <i>Ulmus gaussonii</i> (Ulmaceae). <i>Genes and Genetic Systems</i> , 2016, 91, 11-14.	0.2	2
8	Extremely low genetic diversity of the northern limit populations of <i>Nypa fruticans</i> (Arecaceae) on Iriomote Island, Japan. <i>Conservation Genetics</i> , 2016, 17, 221-228.	0.8	10
9	Hidden founders? Strong bottlenecks and fine-scale genetic structure in mangrove populations of the Cameroon Estuary complex. <i>Hydrobiologia</i> , 2017, 803, 189-207.	1.0	21
10	Marginal distribution and high heterozygosity of asexual <i>Caloglossa vieillardii</i> (Delesseriaceae, Tj ETQq1 1 0.784314 rgBT / Overlock 10	1.0	12
11	Vicariance and Oceanic Barriers Drive Contemporary Genetic Structure of Widespread Mangrove Species <i>Sonneratia alba</i> J. Sm in the Indo-West Pacific. <i>Forests</i> , 2017, 8, 483.	0.9	23
12	Mangrove Biogeography of the Indo-Pacific. <i>Tasks for Vegetation Science</i> , 2019, , 379-400.	0.6	18
13	A general framework for propagule dispersal in mangroves. <i>Biological Reviews</i> , 2019, 94, 1547-1575.	4.7	88
14	High genetic diversity in a "recent outbreak" spider mite, <i>Tetranychus pueraricola</i> , in mainland China. <i>Experimental and Applied Acarology</i> , 2019, 78, 15-27.	0.7	4
15	<i>Avicennia marina</i> maintains genetic structure whereas <i>Rhizophora stylosa</i> connects mangroves in a flooded, former inner sea (Vietnam). <i>Estuarine, Coastal and Shelf Science</i> , 2019, 222, 195-204.	0.9	11
16	Geography alone cannot explain <i>Tetranychus truncatus</i> (Acari: Tetranychidae) population abundance and genetic diversity in the context of the center-periphery hypothesis. <i>Heredity</i> , 2020, 124, 383-396.	1.2	9
17	De novo transcriptome assembly and population genetic analyses of an important coastal shrub, <i>Apocynum venetum</i> L. <i>BMC Plant Biology</i> , 2020, 20, 408.	1.6	10
18	Spatial Structure and Genetic Variation of a Mangrove Species (<i>Avicennia marina</i> (Forssk.) Vierh) in the Farasan Archipelago. <i>Forests</i> , 2020, 11, 1287.	0.9	13

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19	Prominent genetic structure across native and introduced ranges of <i>Pluchea indica</i> , a mangrove associate, as revealed by microsatellite markers. <i>Journal of Plant Ecology</i> , 2020, 13, 341-353.	1.2	2
20	Loss of natural coastline influences species diversity of anemonefish and host anemones in the Ryukyu Archipelago. <i>Aquatic Conservation: Marine and Freshwater Ecosystems</i> , 2021, 31, 15-27.	0.9	5
21	Deciphering the global phylogeography of a coastal shrub (<i>Scaevola taccada</i>) reveals the influence of multiple forces on contemporary population structure. <i>Journal of Systematics and Evolution</i> , 2022, 60, 809-823.	1.6	2
22	Genetic diversity, genetic structure, and demographic history of <i>Cinnamomum chago</i> , a plant species with extremely small populations in China. <i>Global Ecology and Conservation</i> , 2021, 31, e01808.	1.0	1
23	Increases in Genetic Diversity of Weedy Rice Associated with Ambient Temperatures and Limited Gene Flow. <i>Biology</i> , 2021, 10, 71.	1.3	4
24	Dispersal limitation of the mangrove <i>Avicennia marina</i> at its South African range limit in strong contrast to connectivity in its core East African region. <i>Marine Ecology - Progress Series</i> , 2016, 545, 123-134.	0.9	32
25	Extremely Stochastic Connectivity of Island Mangroves. <i>Frontiers in Marine Science</i> , 0, 9, .	1.2	0
26	Genetic diversity and population structure of <i>Bruguiera cylindrica</i> along coastal areas in Thailand. <i>Aquatic Botany</i> , 2022, 183, 103575.	0.8	3
27	Genetic diversity and structure of kola tree (<i>Cola nitida</i>) clones germplasm in Côte d'Ivoire using Single Nucleotide Polymorphism markers. <i>Biodiversitas</i> , 2022, 23, .	0.2	1
28	Assessment of the Genetic Diversity and Population Structure of <i>Rhizophora mucronata</i> along Coastal Areas in Thailand. <i>Biology</i> , 2023, 12, 484.	1.3	1