

# Yu Matsuki

## List of Publications by Year in descending order

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Version: 2024-02-01

31  
papers

910  
citations

567144

15  
h-index

477173

29  
g-index

31  
all docs

31  
docs citations

31  
times ranked

1115  
citing authors

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Testing species hypotheses in the mangrove genus <i>Rhizophora</i> from the Western hemisphere and South Pacific islands. <i>Estuarine, Coastal and Shelf Science</i> , 2021, 248, 106948.  | 0.9 | 7         |
| 2  | NET-CAGE characterizes the dynamics and topology of human transcribed cis-regulatory elements. <i>Nature Genetics</i> , 2019, 51, 1369-1379.  | 9.4 | 72        |
| 3  | Testing the effects of plant species loss on multiple ecosystem functions based on extinction scenarios. <i>Basic and Applied Ecology</i> , 2019, 38, 13-22.  | 1.2 | 4         |
| 4  | Genetic structure of <i>Pinus parviflora</i> on Mt. Fuji in relation to the hoarding behavior of the Japanese nutcracker. <i>Ecosphere</i> , 2019, 10, e02694.  | 1.0 | 1         |
| 5  | Pretreatment of Lignocellulosic Biomass with Cattle Rumen Fluid for Methane Production: Fate of Added Rumen Microbes and Indigenous Microbes of Methane Seed Sludge. <i>Microbes and Environments</i> , 2019, 34, 421-428.                                    | 0.7 | 17        |
| 6  | Phylogeographic analysis suggests two origins for the riparian azalea <i>Rhododendron indicum</i> (L.) Sweet. <i>Heredity</i> , 2018, 121, 594-604.   | 1.2 | 16        |
| 7  | Development of microsatellite markers for the endangered orchid <i>Calanthe izu-insularis</i> (Orchidaceae). <i>Genes and Genetic Systems</i> , 2018, 93, 31-35.  | 0.2 | 3         |
| 8  | The phylogeography of the cycad genus <i>Dioon</i> (Zamiaceae) clarifies its Cenozoic expansion and diversification in the Mexican transition zone. <i>Annals of Botany</i> , 2018, 121, 535-548.   | 1.4 | 42        |
| 9  | Considering evolutionary processes in cycad conservation: identification of evolutionarily significant units within <i>Dioon sonorense</i> (Zamiaceae) in northwestern Mexico. <i>Conservation Genetics</i> , 2018, 19, 1069-1081.                            | 0.8 | 15        |
| 10 | Pretreatment of lignocellulosic biomass by cattle rumen fluid for methane production: Bacterial flora and enzyme activity analysis. <i>Journal of Bioscience and Bioengineering</i> , 2017, 123, 489-496.   | 1.1 | 48        |
| 11 | Population genetics information for the regional conservation of a tropical seagrass, <i>Enhalus acoroides</i> , around the Guimaras Strait, Philippines. <i>Conservation Genetics</i> , 2017, 18, 789-798.   | 0.8 | 8         |
| 12 | Inconsistency between morphological traits and ancestry of individuals in the hybrid zone between two <i>Rhododendron japonoheptamerum</i> varieties revealed by a genotyping-by-sequencing approach. <i>Tree Genetics and Genomes</i> , 2017, 13, 1.         | 0.6 | 22        |
| 13 | Chloroplast DNA sequencing and detailed microsatellite genotyping of all remnant populations suggests that only a single genet survives of the critically endangered plant <i>Rehmannia japonica</i> . <i>Journal of Plant Research</i> , 2017, 130, 117-124. | 1.2 | 3         |
| 14 | Population genetic diversity and structure of a dominant tropical seagrass, <i>Cymodocea rotundata</i>, in the Western Pacific region. <i>Marine Ecology</i> , 2016, 37, 786-800.   | 0.4 | 12        |
| 15 | Lack of genetic variation prevents adaptation at the geographic range margin in a damselfly. <i>Molecular Ecology</i> , 2016, 25, 4450-4460.  | 2.0 | 40        |
| 16 | Development of 11 microsatellite markers in <i>Pinus parviflora</i> by the dual-suppression technique and next-generation sequencing. <i>Journal of Forest Research</i> , 2016, 21, 193-196.  | 0.7 | 1         |
| 17 | A baseline for the genetic conservation of tropical seagrasses in the western North Pacific under the influence of the Kuroshio Current: the case of <i>Syringodium isoetifolium</i> . <i>Conservation Genetics</i> , 2016, 17, 103-110.                      | 0.8 | 18        |
| 18 | MIG-seq: an effective PCR-based method for genome-wide single-nucleotide polymorphism genotyping using the next-generation sequencing platform. <i>Scientific Reports</i> , 2015, 5, 16963.   | 1.6 | 244       |

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 19 | Genetic diversity and structure of the tropical seagrass <i>Cymodocea serrulata</i> spanning its central diversity hotspot and range edge. <i>Aquatic Ecology</i> , 2015, 49, 357-372.   | 0.7 | 29        |
| 20 | Isolation and characterization of novel microsatellite markers for <i>Cymodocea serrulata</i> ( <i>Cymodoceaceae</i> ), a seagrass distributed widely in the Indo-Pacific region. <i>Plant Species Biology</i> , 2015, 30, 297-299.          | 0.6 | 6         |
| 21 | High-throughput linkage mapping of Australian white cypress pine ( <i>Callitris glaucophylla</i> ) and map transferability to related species. <i>Tree Genetics and Genomes</i> , 2015, 11, 1.   | 0.6 | 70        |
| 22 | Development of novel microsatellite markers for <i>Cymodocea rotundata</i> Ehrenberg ( <i>Cymodoceaceae</i> ), a pioneer seagrass species widely distributed in the Indo-Pacific. <i>Conservation Genetics Resources</i> , 2014, 6, 135-138. | 0.4 | 8         |
| 23 | The <i>Keuroshio Current</i> influences genetic diversity and population genetic structure of a tropical seagrass, <i>Enhalus acoroides</i> . <i>Molecular Ecology</i> , 2014, 23, 6029-6044.  | 2.0 | 49        |
| 24 | Development of microsatellite markers in a tropical seagrass <i>Syringodium isoetifolium</i> ( <i>Cymodoceaceae</i> ). <i>Conservation Genetics Resources</i> , 2013, 5, 715-717.  | 0.4 | 9         |
| 25 | Development of 10 novel polymorphic microsatellite markers for the Indo-Pacific horned starfish, <i>Protoreaster nodosus</i> . <i>Marine Genomics</i> , 2013, 11, 27-29.   | 0.4 | 1         |
| 26 | Development of novel microsatellite markers in a tropical seagrass, <i>Enhalus acoroides</i> . <i>Conservation Genetics Resources</i> , 2012, 4, 515-517.  | 0.4 | 12        |
| 27 | Development of microsatellite markers for <i>Thalassia hemprichii</i> ( <i>Hydrocharitaceae</i> ), a widely distributed seagrass in the Indo-Pacific. <i>Conservation Genetics Resources</i> , 2012, 4, 1007-1010.                           | 0.4 | 3         |
| 28 | Pollination Efficiencies of Insects Visiting <i>Magnolia obovata</i> , as Determined by Single-Pollen Genotyping. <i>Structure and Function of Mountain Ecosystems in Japan</i> , 2011, , 17-32.   | 0.1 | 1         |
| 29 | Pollination efficiencies of flower-visiting insects as determined by direct genetic analysis of pollen origin. <i>American Journal of Botany</i> , 2008, 95, 925-930.  | 0.8 | 62        |
| 30 | The determination of multiple microsatellite genotypes and DNA sequences from a single pollen grain. <i>Molecular Ecology Notes</i> , 2007, 7, 194-198.  | 1.7 | 51        |
| 31 | Genetic and reproductive consequences of forest fragmentation for populations of <i>Magnolia obovata</i> . <i>Ecological Research</i> , 2007, 22, 382-389.   | 0.7 | 36        |