Filippos A Aravanopoulos

List of Publications by Year in descending order

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79 papers 2,038 citations

20 h-index 276875 41 g-index

82 all docs 82 docs citations

82 times ranked 3686 citing authors

#	Article	IF	Citations
1	The GenTree Leaf Collection: Inter―and intraspecific leaf variation in seven forest tree species in Europe. Global Ecology and Biogeography, 2021, 30, 590-597.	5.8	11
2	The GenTree Platform: growth traits and tree-level environmental data in 12 European forest tree species. GigaScience, $2021,10,10$	6.4	3
3	Phenotypic, Genetic, and Epigenetic Variation among Diverse Sweet Cherry Gene Pools. Agronomy, 2021, 11, 680.	3.0	6
4	Genetic Characterisation of Chestnut Cultivars in Crete. Forests, 2021, 12, 1659.	2.1	3
5	The GenTree Dendroecological Collection, tree-ring and wood density data from seven tree species across Europe. Scientific Data, 2020, 7, 1.	5.3	830
6	Evolutionary rate and genetic load in an emblematic Mediterranean tree following an ancient and prolonged population collapse. Molecular Ecology, 2020, 29, 4797-4811.	3.9	15
7	Genetics to the rescue: managing forests sustainably in a changing world. Tree Genetics and Genomes, 2020, $16,1.$	1.6	11
8	Whole genome re-sequencing of sweet cherry (Prunus avium L.) yields insights into genomic diversity of a fruit species. Horticulture Research, 2020, 7, 60.	6.3	27
9	A draft genome of sweet cherry (<i>Prunus avium</i> L.) reveals genomeâ€wide and local effects of domestication. Plant Journal, 2020, 103, 1420-1432.	5.7	23
10	Dataset of Targeted Metabolite Analysis for Five Taxanes of Hellenic Taxus baccata L. Populations. Data, 2020, 5, 22.	2.3	4
11	Genomics Opportunities and Breeding Strategies Towards Improvement of Climate-Smart Traits and Disease Resistance Against Pathogens in Sweet Cherry. , 2020, , 385-404.		2
12	Genetic Analysis by nuSSR Markers of Silver Birch (Betula pendula Roth) Populations in Their Southern European Distribution Range. Frontiers in Plant Science, 2020, 11, 310.	3.6	13
13	Assessment of Genetic Diversity and Population Genetic Structure of Norway Spruce (Picea abies (L.)) Tj ETQq1 1 (Resources. Forests, 2019, 10, 258.	0.784314 2.1	rgBT /Overlo 32
14	Conserving Biodiversity and Plant Genetic Resources: The Hellenic Legal Framework. Advances in Global Change Research, 2019, , 141-148.	1.6	0
15	State of Biodiversity and Forest Genetic Resources in Greece in Relation to Conservation. Advances in Global Change Research, 2019, , 73-83.	1.6	2
16	$\hat{l}^{\intercal M}$ ntra-species grafting induces epigenetic and metabolic changes accompanied by alterations in fruit size and shape of Cucurbita pepo L Plant Growth Regulation, 2019, 87, 93-108.	3.4	17
17	From Nature Conservation to Dynamic Genetic Conservation: The Hellenic Case. Advances in Global Change Research, 2019, , 207-216.	1.6	O
18	Overview on Forest Genetic Monitoring (FGM) Including Case Studies on FGM for Two Species from Greece. Advances in Global Change Research, 2019, , 401-407.	1.6	0

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19	Climate Change Impacts on the Genetics of Post-Fire Regeneration and Reproductive Phenology. Advances in Global Change Research, 2019, , 449-457.	1.6	O
20	Ex Situ Conservation of Forest Genetic Resources in Greece. Advances in Global Change Research, 2019, , 291-301.	1.6	1
21	Obituary - Professor Kostas Panetsos (1930-2018). Silvae Genetica, 2019, 68, 79-80.	0.8	O
22	Crossing the Mid-Aegean Trench: vicariant evolution of the Eastern pine processionary moth, Thaumetopoea wilkinsoni (Lepidoptera: Notodontidae), in Crete. Biological Journal of the Linnean Society, 2018, 124, 228-236.	1.6	4
23	Genetic diversity of Thymus sibthorpii Bentham in mountainous natural grasslands of Northern Greece as related to local factors and plant community structure. Industrial Crops and Products, 2018, 111, 651-659.	5.2	13
24	Mediterranean Islands Hosting Marginal and Peripheral Forest Tree Populations: The Case of Pinus brutia Ten. in Cyprus. Forests, 2018, 9, 514.	2.1	6
25	Phenotypic and molecular characterization of apple (Malus $ ilde{A}-$ domestica Borkh) genetic resources in Greece. Scientia Agricola, 2018, 75, 509-518.	1.2	13
26	Towards sweet cherry (Prunus avium L.) breeding: phenotyping evaluation of newly developed hybrids. Euphytica, 2018, 214, 1.	1.2	5
27	The Interplay between Forest Management Practices, Genetic Monitoring, and Other Long-Term Monitoring Systems. Forests, 2018, 9, 133.	2.1	17
28	Do Silviculture and Forest Management Affect the Genetic Diversity and Structure of Long-Impacted Forest Tree Populations?. Forests, 2018, 9, 355.	2.1	26
29	Linkage and QTL mapping in Cupressus sempervirens L. provides the first detailed genetic map of the species and identifies a QTLassociated with crown form. Tree Genetics and Genomes, 2017, 13, 1.	1.6	2
30	Adaptive response of Pinus monticola driven by positive selection upon resistance gene analogs (RGAs) of the TIR-NBS-LRR subfamily. IForest, 2017, 10, 237-241.	1.4	2
31	Evidence of extensive positive selection acting on cherry (Prunus avium L.) resistance gene analogs (RGAs). Australian Journal of Crop Science, 2016, 10, 1324-1329.	0.3	7
32	Evolution-based approach needed for the conservation and silviculture of peripheral forest tree populations. Forest Ecology and Management, 2016, 375, 66-75.	3.2	97
33	Forest genetic monitoring: an overview of concepts and definitions. Environmental Monitoring and Assessment, 2016, 188, 493.	2.7	33
34	Conservation and Monitoring of Tree Genetic Resources in Temperate Forests. Current Forestry Reports, 2016, 2, 119-129.	7.4	38
35	Morpho-physiological diversity in the collection of sour cherry (Prunus cerasus) cultivars of the Fruit Genebank in Naoussa, Greece using multivariate analysis. Scientia Horticulturae, 2016, 207, 225-232.	3.6	15
36	In silico analysis of the LRR receptor-like serine threonine kinases subfamily in Morus notabilis. Plant OMICS, 2016, 9, 319-326.	0.4	4

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37	First Report of an Arbuscular Mycorrhizal Fungus Funneliformis mosseae Associated with Thuja plicata in an Ectomycorrhizal Forest in Greece. International Journal of Phytopathology, 2016, 5, 53-53.	0.5	O
38	Global DNA methylation changes in Cucurbitaceae inter-species grafting. Crop Breeding and Applied Biotechnology, 2015, 15, 112-116.	0.4	33
39	Determination of epigenetic inheritance, genetic inheritance, and estimation of genome DNA methylation in a full-sib family of Cupressus sempervirens L Gene, 2015, 562, 180-187.	2.2	21
40	Population and Conservation Genomics in Forest and Fruit Trees. Advances in Botanical Research, 2015, , 125-155.	1.1	8
41	Beyond population genetics: natural epigenetic variation in wild cherry (Prunus avium). Tree Genetics and Genomes, 2015, 11 , 1 .	1.6	24
42	Diversity of morpho-physiological traits in worldwide sweet cherry cultivars of GeneBank collection using multivariate analysis. Scientia Horticulturae, 2015, 197, 381-391.	3.6	25
43	Mediterranean basin Ficus carica L.: from genetic diversity and structure to authentication of a Protected Designation of Origin cultivar using microsatellite markers. Trees - Structure and Function, 2015, 29, 1959-1971.	1.9	16
44	Selection Strategy for Chestnut (<i>Castanea sativa</i> Mill.) Families Originating from Contrasting European Populations. Open Journal of Forestry, 2015, 05, 489-499.	0.3	1
45	Microsatellite high-resolution melting (SSR-HRM) analysis for identification of sweet cherry rootstocks in Greece. Plant Genetic Resources: Characterisation and Utilisation, 2014, 12, 160-163.	0.8	4
46	Genetic effects of forest management practices: Global synthesis and perspectives. Forest Ecology and Management, 2014, 333, 52-65.	3.2	102
47	Global to local genetic diversity indicators of evolutionary potential in tree species within and outside forests. Forest Ecology and Management, 2014, 333, 35-51.	3.2	57
48	Micropropagation of Elite Genotypes of Castanea Sativa (MILL.). Journal of Advances in Biotechnology, 2014, 3, 200-209.	0.1	1
49	Genetic differentiation and gene flow between wild and cultivated <i>Prunus avium </i> : An analysis of molecular genetic evidence at a regional scale. Plant Biosystems, 2013, 147, 678-685.	1.6	11
50	Taxonomic Identification of Mediterranean Pines and Their Hybrids Based on the High Resolution Melting (HRM) and trnL Approaches: From Cytoplasmic Inheritance to Timber Tracing. PLoS ONE, 2013, 8, e60945.	2.5	30
51	Genome and population dynamics under selection and neutrality: an example of S-allele diversity in wild cherry (Prunus avium L.). Tree Genetics and Genomes, 2012, 8, 1181-1190.	1.6	13
52	Genetic monitoring in natural perennial plant populations. Botany, 2011, 89, 75-81.	1.0	38
53	Linkage mapping of the Mediterranean cypress, Cupressus sempervirens, based on molecular and morphological markers. Genetics and Molecular Research, 2011, 10, 1891-1909.	0.2	9
54	Is the genetic diversity of small scattered forest tree populations at the southern limits of their range more prone to stochastic events? A wild cherry case study by microsatellite-based markers. Tree Genetics and Genomes, 2011, 7, 1299-1313.	1.6	27

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55	Interfertile oaks in an island environment: I. High nuclear genetic differentiation and high degree of chloroplast DNA sharing between Q. alnifolia and Q. coccifera in Cyprus. A multipopulation study. European Journal of Forest Research, 2011, 130, 543-555.	2.5	20
56	Interfertile oaks in an island environment. II. Limited hybridization between Quercus alnifolia Poech and Q. coccifera L. in a mixed stand. European Journal of Forest Research, 2011, 130, 623-635.	2.5	25
57	Molecular identification of Greek olive(Olea europaea) cultivars based on microsatellite loci. Genetics and Molecular Research, 2010, 9, 1865-1876.	0.2	30
58	Assessing inter- and intra-cultivar variation in Greek <i>Prunus avium</i> by SSR markers. Plant Genetic Resources: Characterisation and Utilisation, 2010, 8, 242-248.	0.8	11
59	DNA fingerprinting of elite Greek wild cherry (Prunus avium L.) genotypes using microsatellite markers. Forestry, 2010, 83, 527-533.	2.3	20
60	Breeding of fast growing forest tree species for biomass production in Greece. Biomass and Bioenergy, 2010, 34, 1531-1537.	5.7	38
61	A comparative fluctuating asymmetry study between two walnut (Juglans regia L.) populations may contribute as an early signal for bio-monitoring. IForest, 2010, 3, 150-152.	1.4	3
62	Detecting interspecific and geographic differentiation patterns in two interfertile oak species (Quercus petraea (Matt.) Liebl. and Q. robur L.) using small sets of microsatellite markers. Forest Ecology and Management, 2010, 259, 2026-2035.	3.2	68
63	EVALUATION OF SELECTED EUROPEAN CHESTNUT (CASTANEA SATIVA) PROVENANCES - I: INTER-PROVENANCE GENETIC VARIATION. Acta Horticulturae, 2010, , 203-213.	0.2	4
64	EVALUATION OF SELECTED EUROPEAN CHESTNUT (CASTANEA SATIVA) PROVENANCES - II: INTRA-PROVENANCE FAMILY VARIATION. Acta Horticulturae, 2010, , 215-224.	0.2	4
65	GENETIC DIVERSITY OF INTENSIVELY AND EXTENSIVELY MANAGED CHESTNUT (CASTANEA SATIVA) ORCHARDS IN GREECE. Acta Horticulturae, 2010, , 121-126.	0.2	0
66	Conservation of Nuclear SSR Loci Reveals High Affinity of Quercus infectoria ssp. veneris A. Kern (Fagaceae) to Section Robur. Plant Molecular Biology Reporter, 2008, 26, 133-141.	1.8	6
67	Artificial hybridization between Robinia pseudoacacia L. and R. pseudoacacia var. monophylla Carr Forestry, 2008, 81, 91-101.	2.3	7
68	Morphological Differentiation and Hybridization between Quercus alnifolia Poech and Quercus coccifera L. (Fagaceae) in Cyprus. Silvae Genetica, 2007, 56, 271-277.	0.8	17
69	Genetic structure of Pinus brutia stands exposed to wild fires. Plant Ecology, 2004, 171, 175-183.	1.6	8
70	Absence of association between heterozygosity and biomass production in Salix exigua Nutt. Theoretical and Applied Genetics, 2000, 100, 1203-1208.	3.6	5
71	Geneti c diversity of superior Salix clones selected for intensive forestry plantations. Biomass and Bioenergy, 1999, 16, 249-255.	5.7	28
72	Heterozygosity and biomass production in Salix eriocephala. Heredity, 1998, 81, 396-403.	2.6	13

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73	Inheritance and Linkage of Isozyme Loci in the Basket Willow (Salix viminalis L.). Journal of Heredity, 1997, 88, 144-150.	2.4	2
74	Genetic relationship between Salix exigua and other North American willows (Salix L.): Evidence from allozyme variation. Biochemical Systematics and Ecology, 1995, 23, 767-771.	1.3	5
75	DNA fingerprinting willows(SalixL.) using polymerase chain reaction with the M13 universal primer. Scandinavian Journal of Forest Research, 1995, 10, 27-31.	1.4	6
76	Inheritance of Isozymes in Salix eriocephala Michx. Journal of Heredity, 1994, 85, 381-388.	2.4	4
77	Molecular Markers as Probes in Screening Genetic Diversity and Gene Pool Sustainability in Salix Intensive Forestry. Journal of Sustainable Forestry, 1994, 1, 1-7.	1.4	2
78	Growth-allometry relations in <i>Salix</i> species and families, having different tree form and being under different mating design. Forestry Chronicle, 1993, 69, 717-720.	0.6	2
79	ConservePlants: An integrated approach to conservation of threatened plants for the 21st Century. Research Ideas and Outcomes, 0, 7, .	1.0	6