

# Carla Santos

## List of Publications by Year in descending order

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Version: 2024-02-01

43  
papers

1,918  
citations

361296

20  
h-index

265120

42  
g-index

43  
all docs

43  
docs citations

43  
times ranked

2121  
citing authors

#	ARTICLE	IF	CITATIONS
1	Taxonomy and Phylogenetic Analysis Reveal One New Genus and Three New Species in <i>Inonotus</i> s.l. (Hymenochaetaceae) from Brazil. <i>Cryptogamie, Mycologie</i> , 2022, 43, .	0.2	5
2	PIMA: A population informative multiplex for the Americas. <i>Forensic Science International: Genetics</i> , 2020, 44, 102200.	1.6	7
3	Additions to neotropical stereoid fungi (Polyporales, Basidiomycota): one new species of <i>Lopharia</i> and one new combination in <i>Phlebiopsis</i> . <i>Mycological Progress</i> , 2020, 19, 31-40.	0.5	9
4	Mycobiota in Chilean chilli <i>Capsicum annuum</i> L. used for production of Merck®. <i>International Journal of Food Microbiology</i> , 2020, 334, 108833.	2.1	11
5	Fungal Endophytic Community Associated with Guarana ( <i>Paullinia cupana</i> Var. <i>Sorbilis</i> ): Diversity Driver by Genotypes in the Centre of Origin. <i>Journal of Fungi</i> (Basel, Switzerland), 2020, 6, 123.	1.5	3
6	Molecular Characterization of <i>Diaporthe</i> Species Associated With Hazelnut Defects. <i>Frontiers in Plant Science</i> , 2020, 11, 611655.	1.7	20
7	<i>Gongronella eborensis</i> sp. nov., from vineyard soil of Alentejo (Portugal). <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2020, 70, 3475-3482.	0.8	6
8	Polyphasic, Including MALDI-TOF MS, Evaluation of Freeze-Drying Long-Term Preservation on <i>Aspergillus</i> (Section <i>Nigri</i> ) Strains. <i>Microorganisms</i> , 2019, 7, 291.	1.6	4
9	ITS rDNA Gene Analysis Versus MALDI-TOF MS For Identification of <i>Neoscytalidium dimidiatum</i> Isolated from Onychomycosis and Dermatomycosis Cases in Medellin (Colombia). <i>Microorganisms</i> , 2019, 7, 306.	1.6	10
10	Ancestry analysis in rural Brazilian populations of African descent. <i>Forensic Science International: Genetics</i> , 2018, 36, 160-166.	1.6	9
11	Inferring biogeographic ancestry with compound markers of slow and fast evolving polymorphisms. <i>European Journal of Human Genetics</i> , 2018, 26, 1697-1707.	1.4	13
12	<i>Penicillium tunisiense</i> sp. nov., a novel species of <i>Penicillium</i> section <i>Ramosa</i> discovered from Tunisian orchard apples. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 3217-3225.	0.8	8
13	Evaluation of the Qiagen 140-SNP forensic identification multiplex for massively parallel sequencing. <i>Forensic Science International: Genetics</i> , 2017, 28, 35-43.	1.6	33
14	Using EuroForMix to analyse complex SNP mixtures, up to six contributors. <i>Forensic Science International: Genetics Supplement Series</i> , 2017, 6, e277-e279.	0.1	4
15	Open source software EuroForMix can be used to analyse complex SNP mixtures. <i>Forensic Science International: Genetics</i> , 2017, 31, 105-110.	1.6	37
16	Inference of Ancestry in Forensic Analysis II: Analysis of Genetic Data. <i>Methods in Molecular Biology</i> , 2016, 1420, 255-285.	0.4	27
17	Inference of Ancestry in Forensic Analysis I: Autosomal Ancestry-Informative Marker Sets. <i>Methods in Molecular Biology</i> , 2016, 1420, 233-253.	0.4	20
18	Inter-laboratory evaluation of the EUROFORGEN Global ancestry-informative SNP panel by massively parallel sequencing using the Ion PGM™. <i>Forensic Science International: Genetics</i> , 2016, 23, 178-189.	1.6	65

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19	The Global AIMs Nano set: A 31-plex SNaPshot assay of ancestry-informative SNPs. <i>Forensic Science International: Genetics</i> , 2016, 22, 81-88.	1.6	57
20	Pacifiplex : an ancestry-informative SNP panel centred on Australia and the Pacific region. <i>Forensic Science International: Genetics</i> , 2016, 20, 71-80.	1.6	60
21	Forensic ancestry analysis with two capillary electrophoresis ancestry informative marker (AIM) panels: Results of a collaborative EDNAP exercise. <i>Forensic Science International: Genetics</i> , 2015, 19, 56-67.	1.6	27
22	Inter-laboratory evaluation of SNP-based forensic identification by massively parallel sequencing using the Ion PGM. <i>Forensic Science International: Genetics</i> , 2015, 17, 110-121.	1.6	105
23	Completion of a worldwide reference panel of samples for an ancestry informative Indel assay. <i>Forensic Science International: Genetics</i> , 2015, 17, 75-80.	1.6	30
24	Studies of East European populations with a 46-plex ancestry-informative indel set. <i>Forensic Science International: Genetics Supplement Series</i> , 2015, 5, e16-e18.	0.1	1
25	The open-source software LRmix can be used to analyse SNP mixtures. <i>Forensic Science International: Genetics Supplement Series</i> , 2015, 5, e50-e51.	0.1	13
26	A SNaPshot of next generation sequencing for forensic SNP analysis. <i>Forensic Science International: Genetics</i> , 2015, 14, 50-60.	1.6	85
27	The Genomic Legacy of the Transatlantic Slave Trade in the Yungas Valley of Bolivia. <i>PLoS ONE</i> , 2015, 10, e0134129.	1.1	8
28	“New turns from old STaRs”: Enhancing the capabilities of forensic short tandem repeat analysis. <i>Electrophoresis</i> , 2014, 35, 3173-3187.	1.3	31
29	Building a forensic ancestry panel from the ground up: The EUROFORGEN Global AIM-SNP set. <i>Forensic Science International: Genetics</i> , 2014, 11, 13-25.	1.6	116
30	The SNPforID 34-plex’s ability to infer level of admixture in individuals. <i>Forensic Science International: Genetics Supplement Series</i> , 2013, 4, e13-e14.	0.1	3
31	Revision of the SNPforID 34-plex forensic ancestry test: Assay enhancements, standard reference sample genotypes and extended population studies. <i>Forensic Science International: Genetics</i> , 2013, 7, 63-74.	1.6	102
32	An assessment of Bayesian and multinomial logistic regression classification systems to analyse admixed individuals. <i>Forensic Science International: Genetics Supplement Series</i> , 2013, 4, e63-e64.	0.1	10
33	Eurasiaplex: A forensic SNP assay for differentiating European and South Asian ancestries. <i>Forensic Science International: Genetics</i> , 2013, 7, 359-366.	1.6	102
34	Development of a novel forensic STR multiplex for ancestry analysis and extended identity testing. <i>Electrophoresis</i> , 2013, 34, 1151-1162.	1.3	34
35	An overview of STRUCTURE: applications, parameter settings, and supporting software. <i>Frontiers in Genetics</i> , 2013, 4, 98.	1.1	432
36	Differentiation of African Components of Ancestry to Stratify Groups in a Case-Control Study of a Brazilian Urban Population. <i>Genetic Testing and Molecular Biomarkers</i> , 2012, 16, 524-530.	0.3	5

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37	Typing short amplicon binary polymorphisms: Supplementary SNP and Indel genetic information in the analysis of highly degraded skeletal remains. <i>Forensic Science International: Genetics</i> , 2012, 6, 469-476.	1.6	60
38	Forensic performance of two insertion-deletion marker assays. <i>International Journal of Legal Medicine</i> , 2012, 126, 725-737.	1.2	70
39	Straightforward Inference of Ancestry and Admixture Proportions through Ancestry-Informative Insertion Deletion Multiplexing. <i>PLoS ONE</i> , 2012, 7, e29684.	1.1	211
40	Characterization of U.S. population samples using a 34plex ancestry informative SNP multiplex. <i>Forensic Science International: Genetics Supplement Series</i> , 2011, 3, e182-e183.	0.1	2
41	Population data for 38 autosomal insertion/deletion (InDels) and 50 SNPS polymorphisms in Argentinean population. <i>Forensic Science International: Genetics Supplement Series</i> , 2011, 3, e419-e420.	0.1	1
42	A study of East Timor variability using the SNPforID 52-plex SNP panel. <i>Forensic Science International: Genetics</i> , 2011, 5, e25-e26.	1.6	13
43	Pharmacogenetics of OATP Transporters Reveals That SLCO1B1 c.388A>G Variant Is Determinant of Increased Atorvastatin Response. <i>International Journal of Molecular Sciences</i> , 2011, 12, 5815-5827.	1.8	49