Sara Palomo-DÃ-ez

List of Publications by Year in descending order

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1478280 1372474 31 123 10 6 citations h-index g-index papers 31 31 31 133 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Metagenomic Research of Infectious Diseases in Archaeological Contexts: Evidence from the Hospital Real de Todos-os-Santos (Portugal). Applied Sciences (Switzerland), 2022, 12, 6096.	1.3	5
2	Genealogy: The Tree Where History Meets Genetics. Genealogy, 2021, 5, 98.	0.4	9
3	Paleogenetic evidence of a Pyrenean Neolithic family: Kinship, physical appearance and biogeography multidisciplinary analysis. Journal of Archaeological Science, 2020, 123, 105226.	1.2	3
4	Spanish allele and haplotype database for 32 X-chromosome Insertion-Deletion polymorphisms. Forensic Science International: Genetics, 2020, 46, 102262.	1.6	8
5	An innovative DNA extraction method: Water versus commercial buffers. Forensic Science International: Genetics Supplement Series, 2019, 7, 282-284.	0.1	1
6	Effect of the activity in secondary transfer of DNA profiles. Forensic Science International: Genetics Supplement Series, 2019, 7, 578-579.	0.1	3
7	Kinship analysis on skeletal ancient remains: The case of "el cerro de la horra―(Burgos, Spain). Forensic Science International: Genetics Supplement Series, 2019, 7, 279-281.	0.1	1
8	Genetic identification of Spanish civil war victims. The state of the art in Catalonia (Northeastern) Tj ETQq0 0 0 r	gBT /Over	loc _锋 10 Tf 50 4
9	X-InDels efficacy evaluation in a critical samples paternity case: A Spanish Civil War case from the memorial of the camposines (Tarragona, Spain). Forensic Science International: Genetics Supplement Series, 2019, 7, 494-495.	0.1	3
10	Dos tumbas individuales calcolÃŧicas en las inmediaciones de los dólmenes de Osorno y Simancas: estudio bioantropológico, ofrenda de perros y â€~postvida' megalÃŧica en el valle medio del Duero. Trabajos De Prehistoria, 2019, 76, 236.	0.2	4
11	Diachronic mtDNA study of the long time occupied archaeological site of Segobriga (Spain) and comparison with nowadays population. Forensic Science International: Genetics Supplement Series, 2019, 7, 859-861.	0.1	O
12	An unusual kinship case from the Spanish Civil War (1936–1939): Ancient versus degraded sample's investigation. Forensic Science International: Genetics Supplement Series, 2019, 7, 690-691.	0.1	1
13	Evaluation of two FTA card elutions with sterile vs distilled water. Forensic Science International: Genetics Supplement Series, 2019, 7, 727-729.	0.1	O
14	Kinship analysis and allelic dropout: a forensic approach on an archaeological case. Annals of Human Biology, 2018, 45, 365-368.	0.4	11
15	"Inhibiting inhibitors― Preliminary results of a new "DNA extraction-amplification―disinhibition technique in critical human samples. Forensic Science International: Genetics Supplement Series, 2017, 6, e197-e199.	0.1	3
16	Genetic characterization and determination of the number of individuals by molecular analysis in a prehistoric finding. Forensic Science International: Genetics Supplement Series, 2017, 6, e487-e489.	0.1	3
17	Comparison of two different DNA extraction methodologies for critical bone or teeth samples. Forensic Science International: Genetics Supplement Series, 2017, 6, e359-e361.	0.1	3
18	An unexpected case in the prehistory of the Iberian Peninsula: Biogeographical origin analysis through mitochondrial DNA. Forensic Science International: Genetics Supplement Series, 2017, 6, e205-e207.	0.1	0

#	Article	IF	Citations
19	Prep-n-Goâ,,¢: A new and fast extraction method for forensic blood samples. Forensic Science International: Genetics Supplement Series, 2017, 6, e265-e266.	0.1	9
20	Sex molecular diagnosis on critical samples: Comparison of different methodologies. Forensic Science International: Genetics Supplement Series, 2017, 6, e385-e387.	0.1	1
21	Presumptive tests: A substitute for Benzidine in blood samples recognition. Forensic Science International: Genetics Supplement Series, 2017, 6, e546-e548.	0.1	3
22	A new strategy for a "direct―amplification of forensic samples. Forensic Science International: Genetics Supplement Series, 2017, 6, e560-e561.	0.1	3
23	Comparison of three comercial kits to the establishment of str genetic profiles on critical samples. Forensic Science International: Genetics Supplement Series, 2017, 6, e200-e202.	0.1	1
24	Biological kinship analysis in extremely critical samples: The case of a Spanish Neolithic necropolis. Forensic Science International: Genetics Supplement Series, 2017, 6, e421-e422.	0.1	1
25	Familiar Kinship? Palaeogenetic and Isotopic Evidence from a Triple Burial of the Cogotas I Archaeological Culture (Bronze Age, Iberian Peninsula). Oxford Journal of Archaeology, 2017, 36, 223-242.	0.3	16
26	Phenotyping the ancient world: The physical appearance and ancestry of very degraded samples from a chalcolithic human remains. Forensic Science International: Genetics Supplement Series, 2017, 6, e484-e486.	0.1	3
27	Looking for a reliable criteria for the establishment of solid STR profiles using ancient critical samples from 3000 to 5000 years ago. Forensic Science International: Genetics Supplement Series, 2015, 5, e78-e80.	0.1	2
28	Kinship analysis in mass graves: evaluation of the Blind Search tool of the Familias 3.0 Software in critical samples. Forensic Science International: Genetics Supplement Series, 2015, 5, e547-e550.	0.1	2
29	Nondestructive extraction DNA method from bones or teeth, true or false?. Forensic Science International: Genetics Supplement Series, 2015, 5, e279-e282.	0.1	16
30	A maternity case with human remains from a XIII–XIV century burial at Uceda, Guadalajara, Central Spain. Forensic Science International: Genetics Supplement Series, 2015, 5, e10-e12.	0.1	2
31	Study of medieval critical samples—a genetic approach to the study of the Mudejar Community. Forensic Science International: Genetics Supplement Series, 2015, 5, e193-e195.	0.1	2