Esther Esteban

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/3549894/publications.pdf

Version: 2024-02-01

67	636	759233	752698
papers	citations	12 h-index	20 g-index

69	69	69	875
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Androgen receptor CAG and GGC polymorphisms in Mediterraneans: repeat dynamics and population relationships. Journal of Human Genetics, 2006, 51, 129-136.	2.3	42
2	Population relationships in the Mediterranean revealed by autosomal genetic data (Alu and) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	2.1	36
3	Mixed origin of the current Tunisian population from the analysis of Alu and Alu/STR compound systems. Journal of Human Genetics, 2010, 55, 827-833.	2.3	33
4	New insights into the genetic history of Tunisians: Data from Alu insertion and apolipoprotein E gene polymorphisms. Annals of Human Biology, 2008, 35, 22-33.	1.0	23
5	Glutamine effects on heat shock protein 70 and interleukines 6 and 10: Randomized trial of glutamine supplementation versus standard parenteral nutrition in critically ill children. Clinical Nutrition, 2016, 35, 34-40.	5.0	23
6	Genetic relationships between southeastern Spain and Morocco: New data on ABO, RH, MNSs, and DUFFY polymorphisms. , 1999, 11, 745-752.		22
7	Classical polymorphisms in Berbers from Moyen Atlas (Morocco): genetics, geography, and historical evidence in the Mediterranean peoples. Annals of Human Biology, 2002, 29, 473-487.	1.0	22
8	The X chromosome Alu insertions as a tool for human population genetics: data from European and African human groups. European Journal of Human Genetics, 2007, 15, 578-583.	2.8	19
9	Apolipoprotein gene polymorphisms and plasma levels in healthy Tunisians and patients with coronary artery disease. Lipids in Health and Disease, 2008, 7, 46.	3.0	19
10	Genetic diversity in Northern Spain (Basque Country and Cantabria): GM and KM variation related to demographic histories. European Journal of Human Genetics, 1998, 6, 315-324.	2.8	18
11	STR genetic diversity in a Mediterranean population from the south of the Iberian Peninsula. Annals of Human Biology, 2010, 37, 254-267.	1.0	18
12	Results From Screening Immigrants of Lowâ€Income Countries: Data From a Public Primary Health Care. Journal of Travel Medicine, 2014, 21, 92-98.	3.0	18
13	Alu insertions in the Iberian Peninsula and north west Africa--genetic boundaries or melting pot?. Collegium Antropologicum, 2003, 27, 491-500.	0.2	18
14	Sex differences in children with severe health conditions: Causes of admission and mortality in a Pediatric Intensive Care Unit. American Journal of Human Biology, 2015, 27, 613-619.	1.6	17
15	Genetic study of the population of Tenerife (Canary Islands, Spain): Protein markers and review of classical polymorphisms. , 1997, 102, 337-349.		16
16	Lack of association between eNOS gene polymorphisms and ischemic heart disease in the Spanish population. , 2003, 116A, 243-248.		12
17	Autosomal and X chromosome Alu insertions in Bolivian Aymaras and Quechuas: Two languages and one genetic pool. American Journal of Human Biology, 2010, 22, 154-162.	1.6	12
18	Usefulness of autosomal STR polymorphisms beyond forensic purposes: data on Arabic- and Berber-speaking populations from central Morocco. Annals of Human Biology, 2012, 39, 297-304.	1.0	12

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19	Measuring fitness heritability: Life history traits versus morphological traits in humans. American Journal of Physical Anthropology, 2017, 164, 321-330.	2.1	12
20	Genetic relationships among Berbers and South Spaniards based on CD4 microsatellite/Alu haplotypes. Annals of Human Biology, 2004, 31, 202-212.	1.0	11
21	Genetic Change in the Polynesian Population of Easter Island: Evidence from Alu Insertion Polymorphisms. Annals of Human Genetics, 2006, 70, 829-840.	0.8	11
22	The Mediterranean Sea as a barrier to gene flow: evidence from variation in and around the F7 and F12 genomic regions. BMC Evolutionary Biology, 2010, 10, 84.	3.2	10
23	Genetic Differentiation and Origin of the Jordanian Population: An Analysis ofAluInsertion Polymorphisms. Genetic Testing and Molecular Biomarkers, 2012, 16, 324-329.	0.7	10
24	Genetic Risk Score of NOS Gene Variants Associated with Myocardial Infarction Correlates with Coronary Incidence across Europe. PLoS ONE, 2014, 9, e96504.	2.5	9
25	How many populations set foot through the Patagonian door? Genetic composition of the current population of BahÃa Blanca (Argentina) based on data from 19 Alu polymorphisms. American Journal of Human Biology, 2007, 19, 827-835.	1.6	8
26	The ins and outs of population relationships in west-Mediterranean islands: data from autosomal Alu polymorphisms and Alu/STR compound systems. Journal of Human Genetics, 2007, 52, 999-1010.	2.3	8
27	Genetic differences among North African Berber and Arab-speaking populations revealed by Y-STR diversity. Annals of Human Biology, 2011, 38, 228-236.	1.0	8
28	Apolipoprotein E/C1/C4/C2 Gene Cluster Diversity in Two Native Andean Populations: Aymaras and Quechuas. Annals of Human Genetics, 2012, 76, 283-295.	0.8	8
29	Human Diversity in Jordan: PolymorphicAluInsertions in General Jordanian and Bedouin Groups. Human Biology, 2014, 86, 131-138.	0.2	8
30	Potential Signals of Natural Selection in the Top Risk Loci for Coronary Artery Disease: 9p21 and 10q11. PLoS ONE, 2015, 10, e0134840.	2.5	8
31	Genetic position of Bahrain natives among wider Middle East populations according to Alu insertion polymorphisms. Annals of Human Biology, 2013, 40, 35-40.	1.0	7
32	Genetic diversity of CYP3A4 and CYP3A5 polymorphisms in North African populations from Morocco and Tunisia. International Journal of Biological Markers, 2015, 30, 148-151.	1.8	7
33	Pharmacogenetics of ugt genes in North African populations. Pharmacogenomics Journal, 2018, 18, 609-612.	2.0	7
34	Gm and Km alleles in two Spanish Pyrenean populations (Andorra and Pallars Sobira): a review of Gm variation in the Western Mediterranean basin. Annals of Human Genetics, 2001, 65, 537-548.	0.8	6
35	Dermatoglyphic characterization of Berbers from Morocco: qualitative and quantitative digital and palm data. Annals of Human Biology, 2002, 29, 442-456.	1.0	6
36	Lack of association between methylenetetrahydrofolate reductase (MTHFR) C677T and ischaemic heart disease (IHD): family-based association study in a Spanish population. Clinical Genetics, 2002, 62, 235-239.	2.0	6

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37	Allele-allele interaction within the F13A1 gene: A risk factor for Ischaemic Heart Disease in Spanish population. Thrombosis Research, 2010, 126, e241-e245.	1.7	6
38	Molecular variation in endothelial nitric oxide synthase gene (eNOS) in western Mediterranean populations. Collegium Antropologicum, 2003, 27, 117-24.	0.2	6
39	A training plan to implement lung ultrasound for diagnosing pneumonia in children. Pediatric Research, 2022, 92, 1115-1121.	2.3	6
40	Apolipoprotein molecular variation in Moroccan Berbers: pentanucleotide (TTTTA)n repeat in the LPA gene and APOE-C1-C2 gene cluster. Clinical Genetics, 2002, 62, 240-244.	2.0	5
41	An unexpected wide population variation of the G1733A polymorphism of the androgen receptor gene: Data on the Mediterranean region. American Journal of Human Biology, 2005, 17, 690-695.	1.6	5
42	Different Evolutionary Histories of the Coagulation Factor VII Gene in Human Populations?. Annals of Human Genetics, 2010, 74, 34-45.	0.8	5
43	Research of the origin of a particular Tunisian group using a physical marker and Alu insertion polymorphisms. Genetics and Molecular Biology, 2011, 34, 371-376.	1.3	5
44	Ethnic composition and genetic differentiation of the Libyan population: insights onAlupolymorphisms. Annals of Human Biology, 2014, 41, 229-237.	1.0	5
45	Genetic Differentiation of North-East Argentina Populations Based on 30 Binary X Chromosome Markers. Frontiers in Genetics, 2018, 9, 208.	2.3	5
46	An unexpected world population variation of MCT1 polymorphism 1470T > A involved in lactate transport. European Journal of Sport Science, 2018, 18, 1376-1382.	2.7	5
47	Prognostic value of biomarkers after cardiopulmonary bypass in pediatrics: The prospective PANCAP study. PLoS ONE, 2019, 14, e0215690.	2.5	5
48	Deviceâ€associated multidrugâ€resistant bacteria surveillance in critically ill children: 10Âyears of experience. Acta Paediatrica, International Journal of Paediatrics, 2021, 110, 203-209.	1.5	5
49	Variation of Rhesus Haplotype Frequencies in North Africans and in Worldwide Population Analyses. International Journal of Human Genetics, 2015, 15, 21-31.	0.1	4
50	Analysis of Genomic Regions Associated With Coronary Artery Disease Reveals Continent-Specific Single Nucleotide Polymorphisms in North African Populations. Journal of Epidemiology, 2016, 26, 264-271.	2.4	4
51	Dinucleotide (CA)n tandem repeats on the human X-chromosome and the history of the Mediterranean populations. Annals of Human Biology, 2018, 45, 72-76.	1.0	4
52	UDP-glucuronosyltransferase genetic variation in North African populations: a comparison with African and European data. Annals of Human Biology, 2018, 45, 516-523.	1.0	4
53	Molecular variation at functional genes and the history of human populations--data on candidate genes for cardiovascular risk in the Mediterranean. Collegium Antropologicum, 2003, 27, 523-36.	0.2	4
54	Genetic variability in the Guahibo population from venezuela. American Journal of Human Biology, 2002, 14, 21-28.	1.6	3

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55	Population variability in some genes involving the haemostatic system: data on the general population of Corsica (France), Sardinia and Sicily (Italy). Genetics and Molecular Biology, 2004, 27, 139-146.	1.3	3
56	E65ÂK polymorphism in KCNMB1 gene is not associated with ischaemic heart disease in Spanish patients. Journal of Human Genetics, 2005, 50, 604-606.	2.3	3
57	Polymorphism FXII 46C>T and cardiovascular risk: additional data from Spanish and Tunisian patients. BMC Research Notes, 2009, 2, 154.	1.4	3
58	Close genetic relationships in vast territories: autosomal and X chromosome Alu diversity in Yakuts from Siberia. Anthropologischer Anzeiger, 2013, 70, 309-317.	0.4	3
59	Morbidity and mortality risk factors of pertussis in pediatrics. Journal of Infection, 2017, 74, 97-100.	3.3	3
60	A Commentary on Genetic affinity and admixture of northern Thai People along their migration route in Northern Thailand: evidence from autosomal STR loci. Journal of Human Genetics, 2011, 56, 99-100.	2.3	1
61	Population structure from NOS genes correlates with geographical differences in coronary incidence across Europe. American Journal of Physical Anthropology, 2016, 161, 634-645.	2.1	1
62	Human Diversity in Jordan: Polymorphic Alu Insertions in General Jordanian and Bedouin Groups. Human Biology, 2014, 86, 131.	0.2	1
63	Distinctive genetic signatures of <i>Alu</i>/STR compound systems revealed by analyses of Mediterranean and Middle East populations. Anthropological Science, 2014, 122, 81-88.	0.4	1
64	Comparative Study of 10 X-STR Markers in Populations of Northeast Argentina. Human Biology, 2019, 91, 9.	0.2	1
65	Biomedical Insights of Human Genetic Diversity in Complex Diseases. BioMed Research International, 2015, 2015, 1-2.	1.9	0
66	Religion and fertility patterns: comparison of life history traits in Catholics and Protestants, Hallstatt (Austria) 1733â€“1908. Journal of Biosocial Science, 2021, 53, 305-318.	1.2	0
67	Genetics, geography, and culture: The population of S. Pietro Island (Sardinia, Italy). American Journal of Physical Anthropology, 1996, 100, 461-471.	2.1	0