

Constanza Duque

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10770276/publications.pdf>

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10
papers

1,091
citations

933447

10
h-index

1372567

10
g-index

10
all docs

10
docs citations

10
times ranked

1457
citing authors

#	ARTICLE	IF	CITATIONS
1	A Genomewide Admixture Map for Latino Populations. American Journal of Human Genetics, 2007, 80, 1024-1036.	6.2	265
2	Strong Amerind/White Sex Bias and a Possible Sephardic Contribution among the Founders of a Population in Northwest Colombia. American Journal of Human Genetics, 2000, 67, 1287-1295.	6.2	198
3	Strong Amerind/White Sex Bias and a Possible Sephardic Contribution among the Founders of a Population in Northwest Colombia. American Journal of Human Genetics, 2000, 67, 1287-1295.	6.2	157
4	Genetic make up and structure of Colombian populations by means of uniparental and biparental DNA markers. American Journal of Physical Anthropology, 2010, 143, 13-20.	2.1	140
5	Autosomal, mtDNA, and Y-Chromosome Diversity in Amerinds: Pre- and Post-Columbian Patterns of Gene Flow in South America. American Journal of Human Genetics, 2000, 67, 1277-1286.	6.2	124
6	African ancestry is associated with risk of asthma and high total serum IgE in a population from the Caribbean Coast of Colombia. Human Genetics, 2009, 125, 565-579.	3.8	62
7	Autosomal, mtDNA, and Y-Chromosome Diversity in Amerinds: Pre- and Post-Columbian Patterns of Gene Flow in South America. American Journal of Human Genetics, 2000, 67, 1277-1286.	6.2	59
8	Amerind Ancestry, Socioeconomic Status and the Genetics of Type 2 Diabetes in a Colombian Population. PLoS ONE, 2012, 7, e33570.	2.5	47
9	Transmission distortion of BDNF variants to bipolar disorder type I patients from a south american population isolate,. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2006, 141B, 435-439.	1.7	21
10	Unexpected inverse correlation between Native American ancestry and Asian American variants of HPV16 in admixed Colombian cervical cancer cases. Infection, Genetics and Evolution, 2014, 28, 339-348.	2.3	18