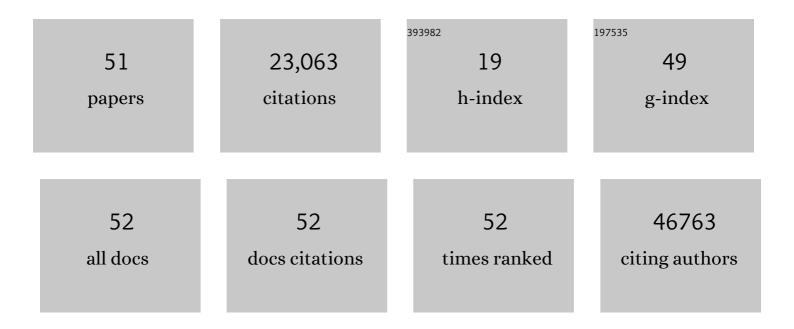
## Marc Via

## List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Molecular and Brain Volume Changes Following Aerobic Exercise, Cognitive and Combined Training in Physically Inactive Healthy Late-Middle-Aged Adults: The Projecte Moviment Randomized Controlled Trial. Frontiers in Human Neuroscience, 2022, 16, 854175.	1.0	8
2	Sex-Specific Protective Effects of APOE ε2 on Cognitive Performance. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2021, 76, 41-49.	1.7	9
3	Exercise and Fitness Neuroprotective Effects: Molecular, Brain Volume and Psychological Correlates and Their Mediating Role in Healthy Late-Middle-Aged Women and Men. Frontiers in Aging Neuroscience, 2021, 13, 615247.	1.7	14
4	Largeâ€scale collaboration in ENIGMAâ€EEG: A perspective on the metaâ€analytic approach to link neurological and psychiatric liability genes to electrophysiological brain activity. Brain and Behavior, 2021, 11, e02188.	1.0	18
5	The Barcelona-Asymptomatic Intracranial Atherosclerosis (AsIA) study: Subclinical cervico-cerebral stenosis and middle cerebral artery pulsatility index as predictors of long-term incident cognitive impairment. Atherosclerosis, 2020, 312, 104-109.	0.4	6
6	Effects and Mechanisms of Cognitive, Aerobic Exercise, and Combined Training on Cognition, Health, and Brain Outcomes in Physically Inactive Older Adults: The Projecte Moviment Protocol. Frontiers in Aging Neuroscience, 2019, 11, 216.	1.7	23
7	Association of <i>GWAS</i> Top Genes With Late-Onset Alzheimer's Disease in Colombian Population. American Journal of Alzheimer's Disease and Other Dementias, 2017, 32, 27-35.	0.9	25
8	Genetic Ancestry and Susceptibility to Late-Onset Alzheimer Disease (LOAD) in the Admixed Colombian Population. Alzheimer Disease and Associated Disorders, 2017, 31, 225-231.	0.6	10
9	COMT and DRD2/ANKK-1 gene-gene interaction account for resetting of gamma neural oscillations to auditory stimulus-driven attention. PLoS ONE, 2017, 12, e0172362.	1.1	7
10	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
11	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.	1.1	4
12	Involvement of the Serotonin Transporter Gene in Accurate Subcortical Speech Encoding. Journal of Neuroscience, 2016, 36, 10782-10790.	1.7	16
13	Pacifiplex : an ancestry-informative SNP panel centred on Australia and the Pacific region. Forensic Science International: Genetics, 2016, 20, 71-80.	1.6	60
14	Potential Signals of Natural Selection in the Top Risk Loci for Coronary Artery Disease: 9p21 and 10q11. PLoS ONE, 2015, 10, e0134840.	1.1	8
15	A global reference for human genetic variation. Nature, 2015, 526, 68-74.	13.7	13,998
16	Genetic Risk Score of NOS Gene Variants Associated with Myocardial Infarction Correlates with Coronary Incidence across Europe. PLoS ONE, 2014, 9, e96504.	1.1	9
17	Human Diversity in Jordan: PolymorphicAluInsertions in General Jordanian and Bedouin Groups. Human Biology, 2014, 86, 131-138.	0.4	8
18	The genetics of Mexico recapitulates Native American substructure and affects biomedical traits. Science, 2014, 344, 1280-1285.	6.0	420

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19	Human Diversity in Jordan: Polymorphic <em>Alu</em> Insertions in General Jordanian and Bedouin Groups. Human Biology, 2014, 86, 131.	0.4	1
20	Role of interactions in pharmacogenetic studies: leukotrienes in asthma. Pharmacogenomics, 2013, 14, 923-929.	0.6	4
21	Reconstructing Native American Migrations from Whole-Genome and Whole-Exome Data. PLoS Genetics, 2013, 9, e1004023.	1.5	185
22	Development of a Panel of Genome-Wide Ancestry Informative Markers to Study Admixture Throughout the Americas. PLoS Genetics, 2012, 8, e1002554.	1.5	212
23	Heterogeneity in Genetic Admixture across Different Regions of Argentina. PLoS ONE, 2012, 7, e34695.	1.1	117
24	Admixture mapping identifies a locus on 6q25 associated with breast cancer risk in US Latinas. Human Molecular Genetics, 2012, 21, 1907-1917.	1.4	60
25	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.	0.4	12
26	Apolipoprotein E/C1/C4/C2 Gene Cluster Diversity in Two Native Andean Populations: Aymaras and Quechuas. Annals of Human Genetics, 2012, 76, 283-295.	0.3	8
27	An integrated map of genetic variation from 1,092 human genomes. Nature, 2012, 491, 56-65.	13.7	7,199
28	Cosmopolitan and ethnic-specific replication of genetic risk factors for asthma in 2 Latino populations. Journal of Allergy and Clinical Immunology, 2011, 128, 37-43.e12.	1.5	34
29	History Shaped the Geographic Distribution of Genomic Admixture on the Island of Puerto Rico. PLoS ONE, 2011, 6, e16513.	1.1	87
30	Population relationships in the Mediterranean revealed by autosomal genetic data ( <i>Alu</i> and) Tj ETQq0 0 0	rgBT /Ove	rlock 10 Tf 5
31	The role of LTA4H and ALOX5AP genes in the risk for asthma in Latinos. Clinical and Experimental Allergy, 2010, 40, 582-589.	1.4	31
32	ALOX5AP and LTA4H polymorphisms modify augmentation of bronchodilator responsiveness by leukotriene modifiers in Latinos. Journal of Allergy and Clinical Immunology, 2010, 126, 853-858.	1.5	19
33	Allele-allele interaction within the F13A1 gene: A risk factor for Ischaemic Heart Disease in Spanish population. Thrombosis Research, 2010, 126, e241-e245.	0.8	6
34	The 1000 Genomes Project: new opportunities for research and social challenges. Genome Medicine, 2010, 2, 3.	3.6	122
35	Recent advances of genetic ancestry testing in biomedical research and direct to consumer testing. Clinical Genetics, 2009, 76, 225-235.	1.0	31

Ancestry-related assortative mating in Latino populations. Genome Biology, 2009, 10, R132. 13.9 89

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37	Augmentation of bronchodilator responsiveness by leukotriene modifiers in Puerto Rican and Mexican children. Annals of Allergy, Asthma and Immunology, 2009, 102, 510-517.	0.5	10
38	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.	0.8	8
39	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.	1.4	19
40	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.	1.1	8
41	Genetic Change in the Polynesian Population of Easter Island: Evidence from Alu Insertion Polymorphisms. Annals of Human Genetics, 2006, 70, 829-840.	0.3	11
42	Androgen receptor CAG and GGC polymorphisms in Mediterraneans: repeat dynamics and population relationships. Journal of Human Genetics, 2006, 51, 129-136.	1.1	42
43	Variability of candidate genes for cardiovascular risk in the Mediterranean. , 2006, , .		0
44	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.	0.8	5
45	E65ÂK polymorphism in KCNMB1 gene is not associated with ischaemic heart disease in Spanish patients. Journal of Human Genetics, 2005, 50, 604-606.	1.1	3
46	Genetic relationships among Berbers and South Spaniards based on CD4 microsatellite/Alu haplotypes. Annals of Human Biology, 2004, 31, 202-212.	0.4	11
47	Poorer cognitive performance in humans with mild cognitive impairment carrying the T variant of the Glu/Asp NOS3 polymorphism. Neuroscience Letters, 2004, 358, 5-8.	1.0	6
48	Lack of association between eNOS gene polymorphisms and ischemic heart disease in the Spanish population. , 2003, 116A, 243-248.		12
49	Molecular variation in endothelial nitric oxide synthase gene (eNOS) in western Mediterranean populations. Collegium Antropologicum, 2003, 27, 117-24.	0.1	6
50	Alu insertions in the Iberian Peninsula and north west Africa–genetic boundaries or melting pot?. Collegium Antropologicum, 2003, 27, 491-500.	0.1	18
51	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.	1.0	6