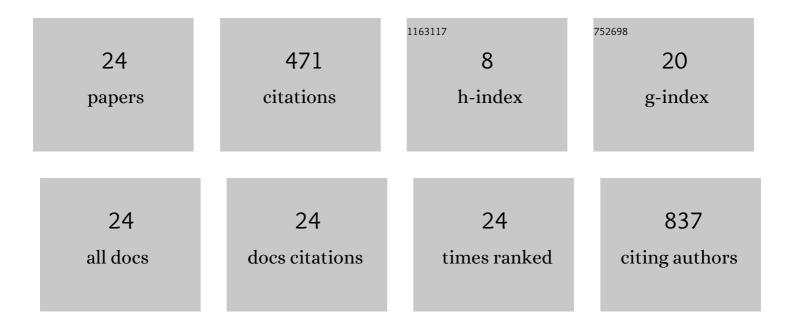
## Júlia Maria Pavan Soler

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2480844/publications.pdf

Version: 2024-02-01



#	Article	IF	CITATIONS
1	Ovarian follicular dynamics in nelore breed () cattle. Theriogenology, 1997, 47, 1489-1505.	2.1	146
2	Brazilian urban population genetic structure reveals a high degree of admixture. European Journal of Human Genetics, 2012, 20, 111-116.	2.8	95
3	Erbium, chromium:yttrium scandium gallium garnet laser for caries removal: influence on bonding of a self-etching adhesive system. Lasers in Medical Science, 2008, 23, 435-441.	2.1	42
4	Assessing functional status after intensive care unit stay: the Barthel Index and the Katz Index. International Journal for Quality in Health Care, 2018, 30, 265-270.	1.8	33
5	Cohort profile: the Baependi Heart Study—a family-based, highly admixed cohort study in a rural Brazilian town. BMJ Open, 2016, 6, e011598.	1.9	32
6	Hemochromatosis gene variants in patients with cardiomyopathy. American Journal of Cardiology, 2001, 88, 388-391.	1.6	25
7	Strategy and model building in the fourth dimension: a null model for genotype × age interaction as a Gaussian stationary stochastic process. BMC Genetics, 2003, 4, S34.	2.7	25
8	Congenic strains provide evidence that four mapped loci in chromosomes 2, 4, and 16 influence hypertension in the SHR. Physiological Genomics, 2009, 37, 52-57.	2.3	16
9	Mineralization of stylohyoid ligament complex in patients with temporomandibular disorders and asymptomatic individuals: a comparative study. Journal of Oral Rehabilitation, 2003, 30, 909-913.	3.0	8
10	Influence of polymorphisms in toll-like receptors (TLRs) on malaria susceptibility in low-endemic area of the Atlantic Forest, São Paulo, Brazil. Acta Tropica, 2018, 182, 309-316.	2.0	8
11	Longitudinal familial analysis of blood pressure involving parametric (co)variance functions. BMC Genetics, 2003, 4, S87.	2.7	7
12	Genetic Analysis of Age-at-Onset for Cardiovascular Risk Factors in a Brazilian Family Study. Human Heredity, 2009, 68, 131-138.	0.8	5
13	Polyamines in tomato plants grown during an incidence of tospovirus exposure. European Journal of Plant Pathology, 2014, 140, 701-709.	1.7	5
14	Heritability and Sex-Specific Genetic Effects of Self-Reported Physical Activity in a Brazilian Highly Admixed Population. Human Heredity, 2019, 84, 151-158.	0.8	5
15	Optimal covariance adjustment in growth curve models. Computational Statistics and Data Analysis, 2000, 33, 101-110.	1.2	4
16	Gene by environment QTL mapping through multiple trait analyses in blood pressure salt-sensitivity: identification of a novel QTL in rat chromosome 5. BMC Medical Genetics, 2006, 7, 47.	2.1	4
17	Using Item Response Theory to Model Multiple Phenotypes and Their Joint Heritability in Family Data. Genetic Epidemiology, 2014, 38, 152-161.	1.3	4
18	Meta-analysis of aspirin for the prevention of preeclampsia: do the main randomized controlled trials support an association between low-dose aspirin and a reduced risk of developing preeclampsia?. Clinics, 2006, 61, 179-182.	1.5	4

#	Article	IF	CITATIONS
19	Bayesian Variable Selection in Multilevel Item Response Theory Models with Application in Genomics. Genetic Epidemiology, 2016, 40, 253-263.	1.3	2
20	Using the theory of added-variable plot for linear mixed models to decompose genetic effects in family data. Statistical Applications in Genetics and Molecular Biology, 2014, 13, 359-78.	0.6	1
21	Causal Inference and Structure Learning of Genotype–Phenotype Networks Using Genetic Variation. , 2016, , 89-143.		Ο
22	Variance-Preserving Estimation of Intensity Values Obtained From Omics Experiments. Frontiers in Genetics, 2019, 10, 855.	2.3	0
23	Learning genetic and environmental graphical models from family data. Statistics in Medicine, 2020, 39, 2403-2422.	1.6	0
24	Factors Associated with Obesity Adjusting for Self-Reported Skin Color-Ethnicity Versus Genomic Ancestry in Admixed Brazilian Population. Current Developments in Nutrition, 2022, 6, 1082.	0.3	0