

Chiara Magri

List of Publications by Year in descending order

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Version: 2024-02-01

40
papers

2,397
citations

430442

18
h-index

288905

40
g-index

42
all docs

42
docs citations

42
times ranked

4591
citing authors

#	ARTICLE	IF	CITATIONS
1	Clinical and Histological Prognostic Factors of Recurrence and Malignant Transformation in a Large Series of Oral Potentially Malignant Disorders. <i>Frontiers in Oncology</i> , 2022, 12, 886404.	1.3	1
2	Investigating an in silico approach for prioritizing antidepressant drug prescription based on drug-induced expression profiles and predicted gene expression. <i>Pharmacogenomics Journal</i> , 2021, 21, 85-93.	0.9	1
3	Alterations observed in the interferon $\hat{1}\pm$ and $\hat{1}^2$ signaling pathway in MDD patients are marginally influenced by cis-acting alleles. <i>Scientific Reports</i> , 2021, 11, 727.	1.6	1
4	Evidence of an interaction between <i>FXR1</i> and <i>GSK3$\hat{1}^2$</i> polymorphisms on levels of Negative Symptoms of Schizophrenia and their response to antipsychotics. <i>European Psychiatry</i> , 2021, 64, e39.	0.1	6
5	Whole Blood Transcriptome Characterization of 3xTg-AD Mouse and Its Modulation by Transcranial Direct Current Stimulation (tDCS). <i>International Journal of Molecular Sciences</i> , 2021, 22, 7629.	1.8	4
6	Clinical validation of a combinatorial PharmAcogeNomic approach in major Depressive disorder: an Observational prospective RANdOmized, participant and rater-blinded, controlled trial (PANDORA) <i>Tj ETQqO 0 0 rgB0, Overlock 10 Tf 50</i>		
7	RNA Editing and Modifications in Mood Disorders. <i>Genes</i> , 2020, 11, 872.	1.0	18
8	Assessment of haptoglobin alleles in autism spectrum disorders. <i>Scientific Reports</i> , 2020, 10, 7758.	1.6	2
9	Association study between <i>HTR2A</i> rs6313 polymorphism and early response to risperidone and olanzapine in schizophrenia patients. <i>Drug Development Research</i> , 2020, 81, 754-761.	1.4	15
10	Genetic determinants of circulating VEGF levels in major depressive disorder and electroconvulsive therapy response. <i>Drug Development Research</i> , 2020, 81, 593-599.	1.4	14
11	Genome-wide analysis of consistently RNA edited sites in human blood reveals interactions with mRNA processing genes and suggests correlations with cell types and biological variables. <i>BMC Genomics</i> , 2018, 19, 963.	1.2	8
12	A novel homozygous mutation in GAD1 gene described in a schizophrenic patient impairs activity and dimerization of GAD67 enzyme. <i>Scientific Reports</i> , 2018, 8, 15470.	1.6	17
13	The effect of childhood trauma on blood transcriptome expression in major depressive disorder. <i>Journal of Psychiatric Research</i> , 2018, 104, 50-54.	1.5	14
14	Compound heterozygosity for a hemizygous rare missense variant (rs141999351) and a large CNV deletion affecting the FSTL5 gene in a patient with schizophrenia. <i>Psychiatry Research</i> , 2017, 258, 598-599.	1.7	1
15	Exome sequencing in schizophrenic patients with high levels of homozygosity identifies novel and extremely rare mutations in the GABA/glutamatergic pathways. <i>PLoS ONE</i> , 2017, 12, e0182778.	1.1	14
16	Meta-analysis of Genome-Wide Association Studies for Extraversion: Findings from the Genetics of Personality Consortium. <i>Behavior Genetics</i> , 2016, 46, 170-182.	1.4	178
17	Meta-analysis of Genome-wide Association Studies for Neuroticism, and the Polygenic Association With Major Depressive Disorder. <i>JAMA Psychiatry</i> , 2015, 72, 642.	6.0	289
18	Proteasome system dysregulation and treatment resistance mechanisms in major depressive disorder. <i>Translational Psychiatry</i> , 2015, 5, e687-e687.	2.4	26

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19	Copy number variants in attention-deficit hyperactive disorder. <i>Psychiatric Genetics</i> , 2015, 25, 59-70.	0.6	25
20	SNP array and FISH analysis of a proband with a 22q13.2-22qter duplication shed light on the molecular origin of the rearrangement. <i>BMC Medical Genetics</i> , 2015, 16, 47.	2.1	5
21	De novo 1Mb interstitial deletion of 8p22 in a patient with slight mental retardation and speech delay. <i>Molecular Cytogenetics</i> , 2014, 7, 25.	0.4	12
22	Combined DOCK8 and CLEC7A mutations causing immunodeficiency in 3 brothers with diarrhea, eczema, and infections. <i>Journal of Allergy and Clinical Immunology</i> , 2013, 131, 594-597.e3.	1.5	22
23	ROLE OF ALLELIC VARIANTS OF FK506-BINDING PROTEIN 51 (FKBP5) GENE IN THE DEVELOPMENT OF ANXIETY DISORDERS. <i>Depression and Anxiety</i> , 2013, 30, 1170-1176.	2.0	42
24	De novo deletion of chromosome 2q24.2 region in a mentally retarded boy with muscular hypotonia. <i>European Journal of Medical Genetics</i> , 2011, 54, 361-364.	0.7	20
25	Analysis of complete mitochondrial genomes of patients with schizophrenia and bipolar disorder. <i>Journal of Human Genetics</i> , 2011, 56, 869-872.	1.1	16
26	New Copy Number Variations in Schizophrenia. <i>PLoS ONE</i> , 2010, 5, e13422.	1.1	82
27	"GenotypeColourâ„“: colour visualisation of SNPs and CNVs. <i>BMC Bioinformatics</i> , 2009, 10, 49.	1.2	4
28	ITGB2 mutation combined with deleted ring 21 chromosome in a child with leukocyte adhesion deficiency. <i>Journal of Allergy and Clinical Immunology</i> , 2009, 124, 1356-1358.	1.5	11
29	Association study and mutational screening of SYNGR1 as a candidate susceptibility gene for schizophrenia. <i>Psychiatric Genetics</i> , 2009, 19, 237-243.	0.6	12
30	Study on GRIA2, GRIA3 and GRIA4 genes highlights a positive association between schizophrenia and GRIA3 in female patients. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2008, 147B, 745-753.	1.1	31
31	Mitochondrial DNA haplogroups and age at onset of schizophrenia. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2007, 144B, 496-501.	1.1	18
32	Glutamate AMPA receptor subunit 1 gene (GRIA1) and DSM-IV-TR schizophrenia: A pilot case-control association study in an Italian sample. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2006, 141B, 287-293.	1.1	35
33	Saami and Berbersâ„“ An Unexpected Mitochondrial DNA Link. <i>American Journal of Human Genetics</i> , 2005, 76, 883-886.	2.6	196
34	Origin, Diffusion, and Differentiation of Y-Chromosome Haplogroups E and J: Inferences on the Neolithization of Europe and Later Migratory Events in the Mediterranean Area. <i>American Journal of Human Genetics</i> , 2004, 74, 1023-1034.	2.6	345
35	Phylogeography of Y-Chromosome Haplogroup I Reveals Distinct Domains of Prehistoric Gene Flow in Europe. <i>American Journal of Human Genetics</i> , 2004, 75, 128-137.	2.6	256
36	The Molecular Dissection of mtDNA Haplogroup H Confirms That the Franco-Cantabrian Glacial Refuge Was a Major Source for the European Gene Pool. <i>American Journal of Human Genetics</i> , 2004, 75, 910-918.	2.6	397

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37	Y-chromosome and mtDNA polymorphisms in Iraq, a crossroad of the early human dispersal and of post-Neolithic migrations. <i>Molecular Phylogenetics and Evolution</i> , 2003, 28, 458-472.	1.2	135
38	From surnames to the history of Y chromosomes: the Sardinian population as a paradigm. <i>European Journal of Human Genetics</i> , 2003, 11, 802-807.	1.4	47
39	Mitochondrial DNA Haplogroups Do Not Play a Role in the Variable Phenotypic Presentation of the A3243G Mutation. <i>American Journal of Human Genetics</i> , 2003, 72, 1005-1012.	2.6	47
40	The 49a,f haplotype 11 is a new marker of the EU19 lineage that traces migrations from northern regions of the black sea. <i>Human Immunology</i> , 2001, 62, 922-932.	1.2	27