Chiara Magri

List of Publications by Year
in descending order

40 papers	2,397 citations	489802 18 h-index	325983 40 g-index
42 all docs	42 docs citations	42 times ranked	5151 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. Frontiers in Oncology, 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. Pharmacogenomics Journal, 2021, 21, 85-93.	0.9	1
3	Alterations observed in the interferon Î± and Î² signaling pathway in MDD patients are marginally influenced by cis-acting alleles. Scientific Reports, 2021, 11, 727.	1.6	1
4	Evidence of an interaction between FXR1 and GSK3Î² polymorphisms on levels of Negative Symptoms of Schizophrenia and their response to antipsychotics. European Psychiatry, 2021, 64, e39.	0.1	6
5	Whole Blood Transcriptome Characterization of 3xTg-AD Mouse and Its Modulation by Transcranial Direct Current Stimulation (tDCS). International Journal of Molecular Sciences, 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) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	5.7	1
7	RNA Editing and Modifications in Mood Disorders. Genes, 2020, 11, 872.	1.0	18
8	Assessment of haptoglobin alleles in autism spectrum disorders. Scientific Reports, 2020, 10, 7758.	1.6	2
9	Association study between <scp>HTR2A</scp> rs6313 polymorphism and early response to risperidone and olanzapine in schizophrenia patients. Drug Development Research, 2020, 81, 754-761.	1.4	15
10	Genetic determinants of circulating VEGF levels in major depressive disorder and electroconvulsive therapy response. Drug Development Research, 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. BMC Genomics, 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. Scientific Reports, 2018, 8, 15470.	1.6	17
13	The effect of childhood trauma on blood transcriptome expression in major depressive disorder. Journal of Psychiatric Research, 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. Psychiatry Research, 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. PLoS ONE, 2017, 12, e0182778.	1.1	14
16	Meta-analysis of Genome-Wide Association Studies for Extraversion: Findings from the Genetics of Personality Consortium. Behavior Genetics, 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. JAMA Psychiatry, 2015, 72, 642.	6.0	289
18	Proteasome system dysregulation and treatment resistance mechanisms in major depressive disorder. Translational Psychiatry, 2015, 5, e687-e687.	2.4	26

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19	Copy number variants in attention-deficit hyperactive disorder. Psychiatric Genetics, 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. BMC Medical Genetics, 2015, 16, 47.	2.1	5
21	De novo 1Mb interstitial deletion of 8p22 in a patient with slight mental retardation and speech delay. Molecular Cytogenetics, 2014, 7, 25.	0.4	12
22	Combined DOCK8 and CLEC7A mutations causing immunodeficiency in 3 brothers with diarrhea, eczema, and infections. Journal of Allergy and Clinical Immunology, 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. Depression and Anxiety, 2013, 30, 1170-1176.	2.0	42
24	De novo deletion of chromosome 2q24.2 region in a mentally retarded boy with muscular hypotonia. European Journal of Medical Genetics, 2011, 54, 361-364.	0.7	20
25	Analysis of complete mitochondrial genomes of patients with schizophrenia and bipolar disorder. Journal of Human Genetics, 2011, 56, 869-872.	1.1	16
26	New Copy Number Variations in Schizophrenia. PLoS ONE, 2010, 5, e13422.	1.1	82
27	"GenotypeColourâ„¢": colour visualisation of SNPs and CNVs. BMC Bioinformatics, 2009, 10, 49.	1.2	4
28	ITGB2 mutation combined with deleted ring 21 chromosome in a child with leukocyte adhesion deficiency. Journal of Allergy and Clinical Immunology, 2009, 124, 1356-1358.	1.5	11
29	Association study and mutational screening of SYNGR1 as a candidate susceptibility gene for schizophrenia. Psychiatric Genetics, 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. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2008, 147B, 745-753.	1.1	31
31	Mitochondrial DNA haplogroups and age at onset of schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 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. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2006, 141B, 287-293.	1.1	35
33	Saami and Berbersâ€"An Unexpected Mitochondrial DNA Link. American Journal of Human Genetics, 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. American Journal of Human Genetics, 2004, 74, 1023-1034.	2.6	345
35	Phylogeography of Y-Chromosome Haplogroup I Reveals Distinct Domains of Prehistoric Gene Flow in Europe. American Journal of Human Genetics, 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. American Journal of Human Genetics, 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. Molecular Phylogenetics and Evolution, 2003, 28, 458-472.	1.2	135
38	From surnames to the history of Y chromosomes: the Sardinian population as a paradigm. European Journal of Human Genetics, 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. American Journal of Human Genetics, 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. Human Immunology, 2001, 62, 922-932.	1.2	27