Chiara Magri

List of Publications by Year in descending order

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40 papers

2,397 citations

430874 18 h-index 289244 40 g-index

42 all docs 42 docs citations

times ranked

42

4591 citing authors

#	Article	IF	CITATIONS
1	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.	6.2	397
2	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.	6.2	345
3	Meta-analysis of Genome-wide Association Studies for Neuroticism, and the Polygenic Association With Major Depressive Disorder. JAMA Psychiatry, 2015, 72, 642.	11.0	289
4	Phylogeography of Y-Chromosome Haplogroup I Reveals Distinct Domains of Prehistoric Gene Flow in Europe. American Journal of Human Genetics, 2004, 75, 128-137.	6.2	256
5	Saami and Berbers—An Unexpected Mitochondrial DNA Link. American Journal of Human Genetics, 2005, 76, 883-886.	6.2	196
6	Meta-analysis of Genome-Wide Association Studies for Extraversion: Findings from the Genetics of Personality Consortium. Behavior Genetics, 2016, 46, 170-182.	2.1	178
7	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.	2.7	135
8	New Copy Number Variations in Schizophrenia. PLoS ONE, 2010, 5, e13422.	2.5	82
9	From surnames to the history of Y chromosomes: the Sardinian population as a paradigm. European Journal of Human Genetics, 2003, 11, 802-807.	2.8	47
10	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.	6.2	47
11	ROLE OF ALLELIC VARIANTS OF FK506-BINDING PROTEIN 51 (FKBP5) GENE IN THE DEVELOPMENT OF ANXIETY DISORDERS. Depression and Anxiety, 2013, 30, 1170-1176.	4.1	42
12	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.	1.7	35
13	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.	1.7	31
14	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.	2.4	27
15	Proteasome system dysregulation and treatment resistance mechanisms in major depressive disorder. Translational Psychiatry, 2015, 5, e687-e687.	4.8	26
16	Copy number variants in attention-deficit hyperactive disorder. Psychiatric Genetics, 2015, 25, 59-70.	1.1	25
17	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.	2.9	22
18	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.	1.3	20

Mitochondrial DNA haplogroups and age at onset of schizophrenia. American Journal of Medical Cenetics Part B: Neuropsychiatric Cenetics, 2007, 1448, 496-501. RNA Editing and Modifications in Mood Disorders. Genes, 2020, 11, 872. 2.4 18 2.4 A novel homozygous mutation in GAD1 gene described in a schizophrenic patient impairs activity and dimerization of CAB67 enzyme. Scientific Reports, 2018, 8, 15470. 2.3 16 2.4 Analysis of complete mitochondrial genomes of patients with schizophrenia and bipolar disorder. Journal of Human Genetics, 2011, 56, 869-872. 2.3 16 2.3 Association grudy between scpp. 41HTDA/d19 (scpp. 16513) polymorphism and early response to risperidone and clianzapine in schizophrenia patients. Drug Development Research, 2020, 81, 754-761. 2.9 15 2.9 16 2.9 16 2.9 17 me effect of childhood trauma on blood transcriptome expression in major depressive disorder. Journal of Psychiatric Research, 2018, 104, 5054. 2.9 14 2.9 Cenetic determinants of circulating VEOT levels in major depressive disorder and electroconvulsive therapy response. Drug Development Research, 2020, 81, 593-599. 2.9 Exome sequencing in schizophrenic patients with high levels of homozygosity Identifies novel and extremely rare mutations in the CABA[glutamatergic pathways, PLoS ONE, 2017, 12, e0182778. 2.5 14 2.7 Association study and mutational screening of SYNGRI as a candidate susceptibility gene for schizophrenia. Psychiatric Genetics, 2009, 19, 237-243. 2.9 ITG62 mutation combined with delected ring 21 chromosome in a child with leukocyte adhesion deficiency, Journal of Allergy and Citical minumology, 2009, 124, 1356-1358. 2.9 ITG62 mutation combined with delected ring 21 chromosome in a child with leukocyte adhesion deficiency, Journal of Allergy and Citical minumology, 2009, 124, 1356-1358. 3.0 Processing genes and suggests correlations with cell types and bloological variables. BMC GenomeNa, 2018, 19, 963. 3.1 Siddence of an interaction between 615 XRI (s) and 40 CSKI31-615 polymorphisms on levels of Negati	#	Article	IF	CITATIONS
Anovel homozygous mutation in GAD1 gene described in a schizophrenic patient impairs activity and dimerization of GAD67 enzyme. Scientific Reports, 2018, 8, 15470. Analysis of complete mitochondrial genomes of patients with schizophrenia and bipolar disorder. Journal of Human Genetics, 2011, 56, 869-872. Analysis of complete mitochondrial genomes of patients with schizophrenia and bipolar disorder. 2.3 16 The effect of childhood trauma on blood transcriptome expression in major depressive disorder. The effect of childhood trauma on blood transcriptome expression in major depressive disorder. The effect of childhood trauma on blood transcriptome expression in major depressive disorder. Cenetic determinants of circulating VECF levels in major depressive disorder and electroconvulsive therapy response. Drug Development Research, 2020, 81, 593-599. Exome sequencing in schizophrenic patients with high levels of homozygosity identifies novel and extremely rare mutations in the CABAlglutamatergic pathways. PLoS ONE, 2017, 12, e0162776. Association study and mutational screening of SYNGR1 as a candidate susceptibility gene for schizophrenia. Psychiatric Genetics, 2009, 19, 237-243. De novo 1Mb interstitial deletion of Bp22 in a patient with slight mental retardation and speech delay. Molecular Cytogenetics, 2014, 7, 25. TIGB2 mutation combined with deleted ring 21 chromosome in a child with leukocyte adhesion deficiency, Journal of Allergy and Clinical immunology, 2003, 124, 1356-1358. Cenome-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. Evidence of an interaction between (1) FXR1 (1): and (1) CKS1P (1): polymorphisms on levels of Negative Symptoms of Schizophrenia and their response to antipsychotics. European Psychiatry, 2021, 64, e39. SNP array and FISH analysis of a proband with a 22q13.2-22qter duplication shed light on the molecular origin	19	Mitochondrial DNA haplogroups and age at onset of schizophrenia. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2007, 144B, 496-501.	1.7	18
dimerization of GAD67 enzyme. Scientific Reports, 2018, 8, 15470. Analysis of complete mitochondrial genomes of patients with schizophrenia and bipolar disorder. Journal of Human Genetics, 2011, 56, 869-872. Association study between escp <15 HTR2Ax/is clscps rs6313 polymorphism and early response to risperdone and olanzaphie in schizophrenia patients. Drug Development Research, 2020, 81, 754-761. The effect of childhood trauma on blood transcriptome expression in major depressive disorder. Journal of Psychiatric Research, 2018, 104, 50-54. Cenetic determinants of circulating VEGF levels in major depressive disorder and electroconvulsive therapy response. Drug Development Research, 2020, 81, 593-599. 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. Association study and mutational screening of SYNCR1 as a candidate susceptibility gene for schizophrenia. Psychiatric Genetics, 2009, 19, 237-243. De novo 1Mb interstitial deletion of 8p22 in a patient with slight mental retardation and speech delay. Molecular Cytogenetics, 2014, 7, 25. De novo 1Mb interstitial deletion of 8p22 in a patient with slight mental retardation and speech delay. Molecular Cytogenetics, 2014, 7, 25. Tig28 mutation combined with deleted ring 21 chromosome in a child with leukocyte adhesion deficiency, Journal of Allergy and Clinical Immunology, 2009, 124, 1356-1358. 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. Evidence of an interaction between cost XXI clis and cis CSK3Pc (is polymorphisms on levels of Negative Symptoms of Schizophrenia and their response to antipsychotics. European Psychiatry, 2021, 64, 699. SNP array and FISH analysis of a proband with a 22q13.2-2 2qter duplication shed light on the molecular origin of th	20	RNA Editing and Modifications in Mood Disorders. Genes, 2020, 11, 872.	2.4	18
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The effect of childhood trauma on blood transcriptome expression in major depressive disorder. The effect of childhood trauma on blood transcriptome expression in major depressive disorder. 3.1 14 25 Genetic determinants of circulating VECF levels in major depressive disorder and electroconvulsive therapy response. Drug Development Research, 2020, 81, 593-599. 26 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. 27 Association study and mutational screening of SYNCR1 as a candidate susceptibility gene for schizophrenia. Psychiatric Genetics, 2009, 19, 237-243. 28 De novo 1Mb interstitial deletion of 8p.22 in a patient with slight mental retardation and speech delay. Molecular Cytogenetics, 2014, 7, 25. 29 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. 20 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. 30 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. 31 Evidence of an interaction between < FXR1 (I) and < FXR2 (I) polymorphisms on levels of Negative Symptoms of Schizophrenia and their response to antipsychotics. European Psychiatry, 2021, 64, e39. 32 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. 33 "GenotypeColourâ,e": colour visualisation of SNPs and CNVs. BMC Bioinformatics, 2009, 10, 49. 4.1 4 Whole Blood Transcriptome Characterization of 3xTg-AD Mouse and its Modulation by Transcranial Direct	22		2.3	16
Journal of Psychiatric Research, 2018, 104, 50-54. 26 Genetic determinants of circulating VEGF levels in major depressive disorder and electroconvulsive therapy response. Drug Development Research, 2020, 81, 593-599. 27 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. 28 Association study and mutational screening of SYNCR1 as a candidate susceptibility gene for schizophrenia. Psychiatric Genetics, 2009, 19, 237-243. 29 De novo 1Mb interstitial deletion of 8p22 in a patient with slight mental retardation and speech delay. Molecular Cytogenetics, 2014, 7, 25. 29 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. 20 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. 30 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. 31 Evidence of an interaction between <1>FXR1 (i)> and <i> SCRS/3² (i)> polymorphisms on levels of Negative Symptoms of Schizophrenia and their response to antipsychotics. European Psychiatry, 2021, 64, e39. 32 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. 33 "GenotypeColoura, c": colour visualisation of SNPs and CNVs. BMC Bioinformatics, 2009, 10, 49. 34 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. 4.1 4</i>	23		2.9	15
therapy response. Drug Development Research, 2020, 81, 593-599. Exome sequencing in schizophrenic patients with high levels of homozygosity identifies novel and extremely rare mutations in the CABA/glutamatergic pathways. PLoS ONE, 2017, 12, e0182778. Association study and mutational screening of SYNGR1 as a candidate susceptibility gene for schizophrenia. Psychiatric Genetics, 2009, 19, 237-243. De novo 1Mb interstitial deletion of 8p22 in a patient with slight mental retardation and speech delay. Molecular Cytogenetics, 2014, 7, 25. De novo 1Mb interstitial deletion of 8p22 in a patient with slight mental retardation and speech delay. Molecular Cytogenetics, 2014, 7, 25. TIGB2 mutation combined with deleted ring 21 chromosome in a child with leukocyte adhesion deficiency. Journal of Allergy and Clinical Immunology, 2009, 124, 1356-1358. 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. Evidence of an interaction between https://doi.org/10/2018/jop/spolymorphisms on levels of Negative Symptoms of Schizophrenia and their response to antipsychotics. European Psychiatry, 2021, 64, e39. 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. SNP array and FISH analysis of a proband with a 22q13.2-21, 64, 64. Whole Blood Transcriptome Characterization of SNPs and CNVs. BMC Bioinformatics, 2009, 10, 49. 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.	24		3.1	14
extremely rare mutations in the GABA/glutamatergic pathways. PLoS ONE, 2017, 12, e0182778. 23	25	Genetic determinants of circulating VEGF levels in major depressive disorder and electroconvulsive therapy response. Drug Development Research, 2020, 81, 593-599.	2.9	14
De novo 1Mb interstitial deletion of 8p22 in a patient with slight mental retardation and speech delay. De novo 1Mb interstitial deletion of 8p22 in a patient with slight mental retardation and speech delay. Molecular Cytogenetics, 2014, 7, 25. 1TGB2 mutation combined with deleted ring 21 chromosome in a child with leukocyte adhesion deficiency. Journal of Allergy and Clinical Immunology, 2009, 124, 1356-1358. 2.9 11 Cenome-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. Evidence of an interaction between <a> FXR1 < > and <a> SCR3 ² < > polymorphisms on levels of Negative Symptoms of Schizophrenia and their response to antipsychotics. European Psychiatry, 2021, 64, e39. 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. 33 "GenotypeColoura, c": colour visualisation of SNPs and CNVs. BMC Bioinformatics, 2009, 10, 49. 4.1 4 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. 4.1 4	26		2.5	14
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deficiency. Journal of Allergy and Clinical Immunology, 2009, 124, 1356-1358. 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. Evidence of an interaction between <i>FXR1</i> and <i>GSK3β</i> polymorphisms on levels of Negative Symptoms of Schizophrenia and their response to antipsychotics. European Psychiatry, 2021, 64, e39. 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. 33 "GenotypeColourâ,¢": colour visualisation of SNPs and CNVs. BMC Bioinformatics, 2009, 10, 49. 4.1 4 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. 4.1 4	28	De novo 1Mb interstitial deletion of 8p22 in a patient with slight mental retardation and speech delay. Molecular Cytogenetics, 2014, 7, 25.	0.9	12
processing genes and suggests correlations with cell types and biological variables. BMC Genomics, 2018, 19, 963. Evidence of an interaction between <i>FXR1</i> symptoms of Schizophrenia and their response to antipsychotics. European Psychiatry, 2021, 64, e39. 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. 33 "GenotypeColourâ,,¢": colour visualisation of SNPs and CNVs. BMC Bioinformatics, 2009, 10, 49. 4.1 4 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. 4.1 4	29	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.	2.9	11
Symptoms of Schizophrenia and their response to antipsychotics. European Psychiatry, 2021, 64, e39. 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. "GenotypeColourâ,,¢": colour visualisation of SNPs and CNVs. BMC Bioinformatics, 2009, 10, 49. 2.6 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. 4.1 4	30	processing genes and suggests correlations with cell types and biological variables. BMC Genomics,	2.8	8
molecular origin of the rearrangement. BMC Medical Genetics, 2015, 16, 47. "GenotypeColourâ,,¢": colour visualisation of SNPs and CNVs. BMC Bioinformatics, 2009, 10, 49. 2.1 3 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. 4.1 4	31		0.2	6
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. 4.1 4	32	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
Direct Current Stimulation (tDCS). International Journal of Molecular Sciences, 2021, 22, 7629.	33	"GenotypeColourâ,,¢": colour visualisation of SNPs and CNVs. BMC Bioinformatics, 2009, 10, 49.	2.6	4
35 Assessment of haptoglobin alleles in autism spectrum disorders. Scientific Reports, 2020, 10, 7758. 3.3 2	34	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.	4.1	4
	35	Assessment of haptoglobin alleles in autism spectrum disorders. Scientific Reports, 2020, 10, 7758.	3.3	2

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. Overloc

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#	Article	IF	CITATIONS
37	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.	3.3	1
38	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.	2.0	1
39	Alterations observed in the interferon \hat{l}^{\pm} and \hat{l}^2 signaling pathway in MDD patients are marginally influenced by cis-acting alleles. Scientific Reports, 2021, 11, 727.	3.3	1
40	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.	2.8	1