Sandra D'Alfonso

List of Publications by Year
in descending order

164	15,245	28274	19749
papers	citations	55 h-index	117 g-index

173	173	173	20158
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	The impact of lifetime coffee and tea loads on Multiple Sclerosis severity. Clinical Nutrition ESPEN, 2022, 47, 199-205.	1.2	0
2	Characterization of the p.L145F and p.S135N Mutations in SOD1: Impact on the Metabolism of Fibroblasts Derived from Amyotrophic Lateral Sclerosis Patients. Antioxidants, 2022, 11, 815.	5.1	3
3	A multi-step genomic approach prioritized TBKBP1 gene as relevant for multiple sclerosis susceptibility. Journal of Neurology, 2022, 269, 4510-4522.	3.6	2
4	Metal(loid)s role in the pathogenesis of amyotrophic lateral sclerosis: Environmental, epidemiological, and genetic data. Environmental Research, 2021, 192, 110292.	7.5	16
5	Pathogenic Huntingtin Repeat Expansions in Patients with Frontotemporal Dementia and Amyotrophic Lateral Sclerosis. Neuron, 2021, 109, 448-460.e4.	8.1	56
6	Mutational Analysis of Known ALS Genes in an Italian Population-Based Cohort. Neurology, 2021, 96, e600-e609.	1.1	23
7	Expanding the genetic spectrum of primary familial brain calcification due to SLC20A2 mutations: a case series. Neurogenetics, 2021, 22, 65-70.	1.4	4
8	Genomic and functional evaluation of TNFSF14 in multiple sclerosis susceptibility. Journal of Genetics and Genomics, 2021, 48, 497-507.	3.9	3
9	Association of Variants in the SPTLC1 Gene With Juvenile Amyotrophic Lateral Sclerosis. JAMA Neurology, 2021, 78, 1236.	9.0	46
10	Targeted Next-Generation Sequencing for the Identification of Genetic Predictors of Radiation-Induced Late Skin Toxicity in Breast Cancer Patients: A Preliminary Study. Journal of Personalized Medicine, 2021, 11, 967.	2.5	3
11	C9ORF72 Repeat Expansion Affects the Proteome of Primary Skin Fibroblasts in ALS. International Journal of Molecular Sciences, 2021, 22, 10385.	4.1	6
12	Validation of an Algorithm to Detect Multiple Sclerosis Cases in Administrative Health Databases in Piedmont (Italy): An Application to the Estimate of Prevalence by Age and Urbanization Level. Neuroepidemiology, 2021, 55, 119-125.	2.3	5
13	An Investigation of the Role of Common and Rare Variants in a Large Italian Multiplex Family of Multiple Sclerosis Patients. Genes, 2021, 12, 1607.	2.4	4
14	Etiological research in pediatric multiple sclerosis: A tool to assess environmental exposures (PEDiatric Italian Genetic and enviRonment ExposurE Questionnaire). Multiple Sclerosis Journal - Experimental, Translational and Clinical, 2021, 7, 205521732110590.	1.0	1
15	Contribution of Rare and Low-Frequency Variants to Multiple Sclerosis Susceptibility in the Italian Continental Population. Frontiers in Genetics, 2021, 12, 800262.	2.3	3
16	Common and rare variant association analyses in amyotrophic lateral sclerosis identify 15 risk loci with distinct genetic architectures and neuron-specific biology. Nature Genetics, 2021, 53, 1636-1648.	21.4	223
17	Deletions in VANGL1 are a risk factor for antibody-mediated kidney disease. Cell Reports Medicine, 2021, 2, 100475.	6.5	2
18	The first case of the TARDBP p.G294V mutation in a homozygous state: is a single pathogenic allele sufficient to cause ALS?. Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration, 2020, 21, 273-279.	1.7	10

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19	Regional spreading of symptoms at diagnosis as a prognostic marker in amyotrophic lateral sclerosis: a population-based study. Journal of Neurology, Neurosurgery and Psychiatry, 2020, 91, 291-297.	1.9	18
20	Generation of an induced pluripotent stem cell line, CSSi011-A (6534), from an Amyotrophic lateral sclerosis patient with heterozygous L145F mutation in SOD1 gene. Stem Cell Research, 2020, 47, 101924.	0.7	2
21	Vitamin D Supplementation Modulates ICOS+ and ICOSâ^' Regulatory T Cell in Siblings of Children With Type 1 Diabetes. Journal of Clinical Endocrinology and Metabolism, 2020, 105, e4767-e4777.	3.6	9
22	ALS phenotype is influenced by age, sex, and genetics. Neurology, 2020, 94, e802-e810.	1.1	99
23	Analysis of the GCG repeat length in NIPA1 gene in C9orf72-mediated ALS in a large Italian ALS cohort. Neurological Sciences, 2019, 40, 2537-2540.	1.9	7
24	The Impact of Lifetime Alcohol and Cigarette Smoking Loads on Multiple Sclerosis Severity. Frontiers in Neurology, 2019, 10, 866.	2.4	14
25	Multiple sclerosis genomic map implicates peripheral immune cells and microglia in susceptibility. Science, 2019, 365, .	12.6	710
26	Shared polygenic risk and causal inferences in amyotrophic lateral sclerosis. Annals of Neurology, 2019, 85, 470-481.	5.3	118
27	Cognitive impairment across ALS clinical stages in a population-based cohort. Neurology, 2019, 93, e984-e994.	1.1	115
28	A case of late-onset OCD developing PLS and FTD. Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration, 2018, 19, 463-465.	1.7	5
29	Characterization of the c9orf72 GC-rich low complexity sequence in two cohorts of Italian and Turkish ALS cases. Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration, 2018, 19, 426-431.	1.7	2
30	HLA alleles modulate EBV viral load in multiple sclerosis. Journal of Translational Medicine, 2018, 16, 80.	4.4	44
31	Genome-wide Analyses Identify KIF5A as a Novel ALS Gene. Neuron, 2018, 97, 1268-1283.e6.	8.1	517
32	Epstein-Barr virus-associated immune reconstitution inflammatory syndrome as possible cause of fulminant multiple sclerosis relapse after natalizumab interruption. Journal of Neuroimmunology, 2018, 319, 9-12.	2.3	21
33	Low-Frequency and Rare-Coding Variation Contributes to Multiple Sclerosis Risk. Cell, 2018, 175, 1679-1687.e7.	28.9	115
34	The multistep hypothesis of ALS revisited. Neurology, 2018, 91, e635-e642.	1.1	146
35	A Census of Tandemly Repeated Polymorphic Loci in Genic Regions Through the Comparative Integration of Human Genome Assemblies. Frontiers in Genetics, 2018, 9, 155.	2.3	9
36	The Length of SNCA Rep1 Microsatellite May Influence Cognitive Evolution in Parkinsonâ€™s Disease. Frontiers in Neurology, 2018, 9, 213.	2.4	21

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37	Overexpression of the Cytokine BAFF and Autoimmunity Risk. New England Journal of Medicine, 2017, 376, 1615-1626.	27.0	301
38	Transancestral mapping and genetic load in systemic lupus erythematosus. Nature Communications, 2017, 8, 16021.	12.8	314
39	TBK1 is associated with ALS and ALS-FTD in Sardinian patients. Neurobiology of Aging, 2016, 43, 180.e1-180.e5.	3.1	40
40	Multiple sclerosis risk loci and disease severity in 7,125 individuals from 10 studies. Neurology: Genetics, 2016, 2, e87.	1.9	76
41	Genome-wide association analyses identify new risk variants and the genetic architecture of amyotrophic lateral sclerosis. Nature Genetics, 2016, 48, 1043-1048.	21.4	494
42	Next Generation Sequencing of Pooled Samples: Guideline for Variantsâ€™ Filtering. Scientific Reports, 2016, 6, 33735.	3.3	81
43	Association of a Locus in theCAMTA1Gene With Survival in Patients With Sporadic Amyotrophic Lateral Sclerosis. JAMA Neurology, 2016, 73, 812.	9.0	57
44	Power estimation for non-standardized multisite studies. NeuroImage, 2016, 134, 281-294.	4.2	36
45	ATNX2 is not a regulatory gene in Italian amyotrophic lateral sclerosis patients with C9ORF72 GGGGCC expansion. Neurobiology of Aging, 2016, 39, 218.e5-218.e8.	3.1	6
46	Genetic variants are major determinants of CSF antibody levels in multiple sclerosis. Brain, 2015, 138, 632-643.	7.6	54
47	PXKlocus in systemic lupus erythematosus: fine mapping and functional analysis reveals novel susceptibility geneABHD6. Annals of the Rheumatic Diseases, 2015, 74, e14-e14.	0.9	24
48	Chitotriosidase and lysosomal enzymes as potential biomarkers of disease progression in amyotrophic lateral sclerosis: A survey clinic-based study. Journal of the Neurological Sciences, 2015, 348, 245-250.	0.6	45
49	Mitochondrial DNA sequence variation in multiple sclerosis. Neurology, 2015, 85, 325-330.	1.1	60
50	Coeliac disease mimicking Amyotrophic Lateral Sclerosis. Amyotrophic Lateral Sclerosis and Frontotemporal Degeneration, 2015, 16, 277-279.	1.7	4
51	TUBA4A gene analysis in sporadic amyotrophic lateral sclerosis: identification of novel mutations. Journal of Neurology, 2015, 262, 1376-1378.	3.6	44
52	Epstein-Barr virus genetic variants are associated with multiple sclerosis. Neurology, 2015, 84, 1362-1368.	1.1	44
53	Inverse correlation of genetic risk score with age at onset in bout-onset and progressive-onset multiple sclerosis. Multiple Sclerosis Journal, 2015, 21, 1463-1467.	3.0	13
54	HLA Genotyping: Methods for the Identification of the HLA-DQ2,-DQ8 Heterodimers Implicated in Celiac Disease (CD) Susceptibility. Methods in Molecular Biology, 2015, 1326, 79-92.	0.9	5

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55	Class II HLA interactions modulate genetic risk for multiple sclerosis. Nature Genetics, 2015, 47, 1107-1113.	21.4	312
56	The burden of multiple sclerosis variants in continental Italians and Sardinians. Multiple Sclerosis Journal, 2015, 21, 1385-1395.	3.0	10
57	ATXN2 polyQ intermediate repeats are a modifier of ALS survival. Neurology, 2015, 84, 251-258.	1.1	52
58	Functional variations modulating PRKCA expression and alternative splicing predispose to multiple sclerosis. Human Molecular Genetics, 2014, 23, 6746-6761.	2.9	32
59	Early onset frontotemporal dementia with psychiatric presentation due to the C9ORF72 hexanucleotide repeat expansion: a case report. BMC Neurology, 2014, 14, 228.	1.8	13
60	Analysis of the KIFAP3 gene in amyotrophic lateral sclerosis: a multicenter survival study. Neurobiology of Aging, 2014, 35, 2420.e13-2420.e14.	3.1	16
61	A genome-wide association meta-analysis identifies a novel locus at 17q11.2 associated with sporadic amyotrophic lateral sclerosis. Human Molecular Genetics, 2014, 23, 2220-2231.	2.9	123
62	Multiple sclerosis progression is not associated with birth timing in Italy. Journal of the Neurological Sciences, 2014, 346, 194-196.	0.6	1
63	Genetic burden of common variants in progressive and bout-onset multiple sclerosis. Multiple Sclerosis Journal, 2014, 20, 802-811.	3.0	11
64	Exome-wide Rare Variant Analysis Identifies TUBA4A Mutations Associated with Familial ALS. Neuron, 2014, 84, 324-331.	8.1	308
65	Familial clustering in Italian progressive-onset and bout-onset multiple sclerosis. Neurological Sciences, 2014, 35, 789-791.	1.9	8
66	No evidence for a role of rare CYP27B1 functional variations in multiple sclerosis. Annals of Neurology, 2013, 73, 433-437.	5.3	31
67	Analysis of immune-related loci identifies 48 new susceptibility variants for multiple sclerosis. Nature Genetics, 2013, 45, 1353-1360.	21.4	1,213
68	Screening of the PFN1 gene in sporadic amyotrophic lateral sclerosis and in frontotemporal dementia. Neurobiology of Aging, 2013, 34, 1517.e9-1517.e10.	3.1	35
69	Analysis of hnRNPA1, A2/B1, and A3 genes in patients with amyotrophic lateral sclerosis. Neurobiology of Aging, 2013, 34, 2695.e11-2695.e12.	3.1	30
70	A â€œCandidate-Interactomeâ€•Aggregate Analysis of Genome-Wide Association Data in Multiple Sclerosis. PLoS ONE, 2013, 8, e63300.	2.5	66
71	Network-Based Multiple Sclerosis Pathway Analysis with GWAS Data from 15,000 Cases and 30,000 Controls. American Journal of Human Genetics, 2013, 92, 854-865.	6.2	164
72	Fine-Mapping the Genetic Association of the Major Histocompatibility Complex in Multiple Sclerosis: HLA and Non-HLA Effects. PLoS Genetics, 2013, 9, e1003926.	3.5	250

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73	Ubiquilin 2mutations in Italian patients with amyotrophic lateral sclerosis and frontotemporal dementia. Journal of Neurology, Neurosurgery and Psychiatry, 2013, 84, 183-187.	1.9	74
74	A systemic sclerosis and systemic lupus erythematosus pan-meta-GWAS reveals new shared susceptibility loci. Human Molecular Genetics, 2013, 22, 4021-4029.	2.9	104
75	Rare Variants in theTREX1Gene and Susceptibility to Autoimmune Diseases. BioMed Research International, 2013, 2013, 1-6.	1.9	34
76	Association of Genetic Markers with CSF Oligoclonal Bands in Multiple Sclerosis Patients. PLoS ONE, 2013, 8, e64408.	2.5	27
77	Genetic and physical interaction of the B-cell systemic lupus erythematosus-associated genes BANK1 and BLK. Annals of the Rheumatic Diseases, 2012, 71, 136-142.	0.9	67
78	Analysis of autosomal genes reveals geneâ€“sex interactions and higher total genetic risk in men with systemic lupus erythematosus. Annals of the Rheumatic Diseases, 2012, 71, 694-699.	0.9	87
79	Fine mapping and conditional analysis identify a new mutation in the autoimmunity susceptibility gene BLK that leads to reduced half-life of the BLK protein. Annals of the Rheumatic Diseases, 2012, 71, 1219-1226.	0.9	33
80	The Impact of Osteopontin Gene Variations on Multiple Sclerosis Development and Progression. Clinical and Developmental Immunology, 2012, 2012, 1-6.	3.3	31
81	Extensive genetics of ALS. Neurology, 2012, 79, 1983-1989.	1.1	145
82	Association study of IRAK-M and SIGIRR genes with SLE in a large European-descent population. Lupus, 2012, 21, 1166-1171.	1.6	11
83	C9ORF72 repeat expansion in a large Italian ALS cohort: evidence of a founder effect. Neurobiology of Aging, 2012, 33, 2528.e7-2528.e14.	3.1	74
84	Novel association of acid phosphatase locus 1*C allele with systemic lupus erythematosus. Human Immunology, 2012, 73, 107-110.	2.4	9
85	The -346T polymorphism of the SH2D1A gene is a risk factor for development of autoimmunity/lymphoproliferation in males with defective Fas function. Human Immunology, 2012, 73, 585-592.	2.4	9
86	Polymorphisms in the genes coding for iron binding and transporting proteins are associated with disability, severity, and early progression in multiple sclerosis. BMC Medical Genetics, 2012, 13, 70.	2.1	42
87	Genetic association of miRNA-146a with systemic lupus erythematosus in Europeans through decreased expression of the gene. Genes and Immunity, 2012, 13, 268-274.	4.1	132
88	Genetic risk and a primary role for cell-mediated immune mechanisms in multiple sclerosis. Nature, 2011, 476, 214-219.	27.8	2,400
89	Association of osteopontin regulatory polymorphisms with systemic sclerosis. Human Immunology, 2011, 72, 930-934.	2.4	32
90	No association of DPP6 with amyotrophic lateral sclerosis in an Italian population. Neurobiology of Aging, 2011, 32, 966-967.	3.1	28

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91	A novel peripherin gene (PRPH) mutation identified in one sporadic amyotrophic lateral sclerosis patient. Neurobiology of Aging, 2011, 32, 552.e1-552.e6.	3.1	49
92	Multiple Polymorphisms Affect Expression and Function of the Neuropeptide S Receptor (NPSR1). PLoS ONE, 2011, 6, e29523.	2.5	30
93	VPS54 genetic analysis in ALS Italian cohort. European Journal of Neurology, 2011, 18, e41-e42.	3.3	6
94	ATXN-2 CAG repeat expansions are interrupted in ALS patients. Human Genetics, 2011, 130, 575-580.	3.8	52
95	Association of HLA class I markers with multiple sclerosis in the Italian and UK population: evidence of two independent protective effects. Journal of Medical Genetics, 2011, 48, 485-492.	3.2	9
96	Association of the CBLB gene with multiple sclerosis: new evidence from a replication study in an Italian population. Journal of Medical Genetics, 2011, 48, 210-211.	3.2	9
97	Novel optineurin mutations in patients with familial and sporadic amyotrophic lateral sclerosis. Journal of Neurology, Neurosurgery and Psychiatry, 2011, 82, 1239-1243.	1.9	86
98	The Genetic Association of Variants in CD6, TNFRSF1A and IRF8 to Multiple Sclerosis: A Multicenter Case-Control Study. PLoS ONE, 2011, 6, e18813.	2.5	63
99	Association of Systemic Lupus Erythematosus Clinical Features with European Population Genetic Substructure. PLoS ONE, 2011, 6, e29033.	2.5	14
100	No evidence of association of the rare nsSNP rs35667974 in IFIH1 with multiple sclerosis. Journal of Neuroimmunology, 2010, 221, 112-114.	2.3	4
101	A 3â€²â€untranslated region variant is associated with impaired expression of CD226 in T and natural killer T cells and is associated with susceptibility to systemic lupus erythematosus. Arthritis and Rheumatism, 2010, 62, 3404-3414.	6.7	48
102	HLA-class I markers and multiple sclerosis susceptibility in the Italian population. Genes and Immunity, 2010, 11, 173-180.	4.1	51
103	IL12A, MPHOSPH9/CDK2AP1 and RGS1 are novel multiple sclerosis susceptibility loci. Genes and Immunity, 2010, 11, 397-405.	4.1	70
104	A non-synonymous SNP within membrane metalloendopeptidase-like 1 (MMEL1) is associated with multiple sclerosis. Genes and Immunity, 2010, 11, 660-664.	4.1	25
105	Lack of support for association between the KIF1B rs10492972[C] variant and multiple sclerosis. Nature Genetics, 2010, 42, 469-470.	21.4	23
106	Immunoproteasome LMP2 60HH Variant Alters MBP Epitope Generation and Reduces the Risk to Develop Multiple Sclerosis in Italian Female Population. PLoS ONE, 2010, 5, e9287.	2.5	56
107	Promoter Insertion/Deletion in the IRF5 Gene Is Highly Associated with Susceptibility to Systemic Lupus Erythematosus in Distinct Populations, But Exerts a Modest Effect on Gene Expression in Peripheral Blood Mononuclear Cells. Journal of Rheumatology, 2010, 37, 574-578.	2.0	32
108	Mutations of FUS gene in sporadic amyotrophic lateral sclerosis. Journal of Medical Genetics, 2010, 47, 190-194.	3.2	152

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109	The Osteopontin Gene +1239A/C Single Nucleotide Polymorphism is Associated with Type 1 Diabetes Mellitus in the Italian Population. International Journal of Immunopathology and Pharmacology, 2010, 23, 263-269.	2.1	21
110	STAT4 associates with systemic lupus erythematosus through two independent effects that correlate with gene expression and act additively with IRF5 to increase risk. Annals of the Rheumatic Diseases, 2009, 68, 1746-1753.	0.9	138
111	Identification of a new putative functional IL18 gene variant through an association study in systemic lupus erythematosus. Human Molecular Genetics, 2009, 18, 3739-3748.	2.9	54
112	High frequency ofTARDBPgene mutations in Italian patients with amyotrophic lateral sclerosis. Human Mutation, 2009, 30, 688-694.	2.5	184
113	Replication of the TNFSF4 (OX40L) promoter region association with systemic lupus erythematosus. Genes and Immunity, 2009, 10, 248-253.	4.1	41
114	Mutations in the lamin B1 gene are not present in multiple sclerosis. European Journal of Neurology, 2009, 16, 544-546.	3.3	4
115	Replication of recently identified systemic lupus erythematosus genetic associations: a caseâ€“control study. Arthritis Research and Therapy, 2009, 11, R69.	3.5	131
116	Kallikrein genes are associated with lupus and glomerular basement membraneâ€“specific antibodyâ€“induced nephritis in mice and humans. Journal of Clinical Investigation, 2009, 119, 911-923.	8.2	114
117	Variations of the perforin gene in patients with multiple sclerosis. Genes and Immunity, 2008, 9, 438-444.	4.1	39
118	Functional variants in the B-cell gene BANK1 are associated with systemic lupus erythematosus. Nature Genetics, 2008, 40, 211-216.	21.4	436
119	A sequence variation in the MOG gene is involved in multiple sclerosis susceptibility in Italy. Genes and Immunity, 2008, 9, 7-15.	4.1	20
120	Refining genetic associations in multiple sclerosis. Lancet Neurology, The, 2008, 7, 567-569.	10.2	90
121	Gender-specific influence of the chromosome 16 chemokine gene cluster on the susceptibility to Multiple Sclerosis. Journal of the Neurological Sciences, 2008, 267, 86-90.	0.6	30
122	A loss-of-function variant of PTPN22 is associated with reduced risk of systemic lupus erythematosus. Human Molecular Genetics, 2008, 18, 569-579.	2.9	106
123	Variations in the coding and regulatory sequences of the angiogenin (ANG) gene are not associated to ALS (amyotrophic lateral sclerosis) in the Italian population. Journal of the Neurological Sciences, 2007, 258, 123-127.	0.6	37
124	Neuropeptide S Receptor 1 Gene Polymorphism Is Associated With Susceptibility to Inflammatory Bowel Disease. Gastroenterology, 2007, 133, 808-817.	1.3	87
125	Bias in association studies of systemic lupus erythematosus susceptibility due to geographical variation in the frequency of a programmed cell death 1 polymorphism across Europe. Genes and Immunity, 2007, 8, 138-146.	4.1	33
126	ICOS gene haplotypes correlate with IL10 secretion and multiple sclerosis evolution. Journal of Neuroimmunology, 2007, 186, 193-198.	2.3	24

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127	SOD1 gene mutations in Italian patients with Sporadic Amyotrophic Lateral Sclerosis (ALS). Neuromuscular Disorders, 2006, 16, 800-804.	0.6	40
128	Concordance, disease progression, and heritability of coeliac disease in Italian twins. Gut, 2006, 55, 803-808.	12.1	155
129	Osteopontin gene haplotypes correlate with multiple sclerosis development and progression. Journal of Neuroimmunology, 2005, 163, 172-178.	2.3	66
130	Two single-nucleotide polymorphisms in the 5? and 3? ends of the osteopontin gene contribute to susceptibility to systemic lupus erythematosus. Arthritis and Rheumatism, 2005, 52, 539-547.	6.7	94
131	HLAâ€“multiple sclerosis association in Continental Italy and correlation with disease prevalence in Europe. Journal of Neuroimmunology, 2004, 150, 178-185.	2.3	66
132	A whole genome screen for linkage disequilibrium in multiple sclerosis performed in a continental Italian population. Journal of Neuroimmunology, 2003, 143, 97-100.	2.3	17
133	Prolactin and prolactin receptor gene polymorphisms in multiple sclerosis and systemic lupus erythematosus. Human Immunology, 2003, 64, 274-284.	2.4	34
134	IL12B Polymorphism and Type 1 Diabetes in the Italian Population: A Case-Control Study. Diabetes, 2002, 51, 1649-1650.	0.6	24
135	Genetics of Multiple Sclerosis. Molecular Diagnosis and Therapy, 2002, 2, 37-58.	3.3	37
136	Identification of single nucleotide variations in the coding and regulatory regions of the myelin-associated glycoprotein gene and study of their association with multiple sclerosis. Journal of Neuroimmunology, 2002, 126, 196-204.	2.3	11
137	The IL12B gene does not confer susceptibility to coeliac disease. Tissue Antigens, 2002, 59, 70-72.	1.0	17
138	Cytokine gene polymorphism in human disease: on-line databases, Supplement 2. Genes and Immunity, 2002, 3, 313-330.	4.1	225
139	Association tests with systemic lupus erythematosus (SLE) of IL10 markers indicate a direct involvement of a CA repeat in the 5â€² regulatory region. Genes and Immunity, 2002, 3, 454-463.	4.1	36
140	A genome screen for multiple sclerosis in Sardinian multiplex families. European Journal of Human Genetics, 2001, 9, 621-626.	2.8	95
141	Cytokine gene polymorphism in human disease: on-line databases, Supplement 1. Genes and Immunity, 2001, 2, 61-70.	4.1	248
142	A genome screen for multiple sclerosis in Italian families. Genes and Immunity, 2001, 2, 205-210.	4.1	70
143	MICA and MICB microsatellite alleles in HLA extended haplotypes. International Journal of Immunogenetics, 2001, 28, 523-530.	1.2	22
144	Systemic lupus erythematosus candidate genes in the Italian population: Evidence for a significant association with interleukin-10. Arthritis and Rheumatism, 2000, 43, 120-128.	6.7	103

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145	New polymorphisms in the IL-10 promoter region. Genes and Immunity, 2000, 1, 231-233.	4.1	98
146	HLA class I in acute promyelocytic leukemia (APL): possible correlation with clinical outcome. Leukemia, 2000, 14, 393-398.	7.2	13
147	Association study of a new polymorphism in the PECAM-1 gene in multiple sclerosis. Journal of Neuroimmunology, 2000, 104, 174-178.	2.3	19
148	Linkage analysis of multiple sclerosis with candidate region markers in Sardinian and Continental Italian families. European Journal of Human Genetics, 1999, 7, 377-385.	2.8	38
149	Cytokine gene polymorphism in human disease: on-line databases. Genes and Immunity, 1999, 1, 3-19.	4.1	560
150	Nonparametric linkage analysis of alcohol dependence with chromosome 1 and 7 markers. Genetic Epidemiology, 1999, 17, S127-S131.	1.3	4
151	The natural history of an HLA haplotype and its recombinants. Immunogenetics, 1998, 48, 8-15.	2.4	10
152	Fas Gene Polymorphisms Are Not Associated With Systemic Lupus Erythematosus, Multiple Sclerosis And Hiv Infection. Disease Markers, 1998, 13, 221-225.	1.3	20
153	An intragenic polymorphism in the human tumor necrosis factor alpha (TNFA) chain-encoding gene. Immunogenetics, 1996, 44, 321-322.	2.4	32
154	Association between polymorphisms in the TNF region and systemic lupus erythematosus in the Italian population. Tissue Antigens, 1996, 47, 551-555.	1.0	68
155	Functional Analysis of a New Polymorphism in the Human TNF Î± Gene Promoter. Scandinavian Journal of Immunology, 1995, 42, 501-504.	2.7	161
156	HLA supratypes in an Italian population. Immunogenetics, 1994, 39, 114-20.	2.4	12
157	A polymorphic variation in a putative regulation box of the TNFA promoter region. Immunogenetics, 1994, 39, 150-4.	2.4	317
158	A practical approach to HLA-DR genomic typing by heteroduplex analysis and a selective cleavage at position 86. Human Immunology, 1994, 40, 41-50.	2.4	4
159	Altered biosynthesis of tumour necrosis factor (TNF) alpha is involved in postburn hypertrophic scars. Burns, 1994, 20, 118-121.	1.9	19
160	Gametic association of HSP70â€1 promoter region alleles and their inclusion in extended HLA haplotypes. Tissue Antigens, 1993, 42, 62-66.	1.0	9
161	QUANTITATIVE EXPRESSION OF HLA CLASS I MOLECULES IN ACUTE NON-LYMPHOBLASTIC LEUKAEMIA CELLS. International Journal of Immunogenetics, 1993, 20, 165-173.	1.2	3
162	Distribution of Tumor Necrosis Factor Alleles (Ncol RFLP) and their Relationship to HLA Haplotypes in an Italian Population. Human Heredity, 1993, 43, 103-110.	0.8	11

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163	Genome-Wide Analyses Identify KIF5A as a Novel ALS Gene. SSRN Electronic Journal, 0, , .	0.4	4
164	The Impact of Life-Time Alcohol and Smoking Load on Multiple Sclerosis Severity. SSRN Electronic Journal, 0, , .	0.4	0