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A "candidate-interactome" aggregate analysis of genome-wide association data in multiple sclerosis

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31	Increased CD8+ T cell responses to apoptotic T cell-associated antigens in multiple sclerosis. Journal of Neuroinflammation, 2013 , 10, 94	10.1	20
30	Susceptibility genes are enriched in those of the herpes simplex virus 1/host interactome in psychiatric and neurological disorders. <i>Pathogens and Disease</i> , 2013 , 69, 240-61	4.2	27
29	Noise in multiple sclerosis: unwanted and necessary. <i>Annals of Clinical and Translational Neurology</i> , 2014 , 1, 502-11	5.3	6
28	Gene-environment interaction in autoimmune disease. <i>Expert Reviews in Molecular Medicine</i> , 2014 , 16, e4	6.7	22
27	Thrombotic processes in multiple sclerosis as manifestation of innate immune activation. <i>Frontiers in Neurology</i> , 2014 , 5, 119	4.1	13
26	High frequency of antiphospholipid antibodies in relapse of multiple sclerosis: a possible indicator of inflammatory-thrombotic processes. <i>Neurological Sciences</i> , 2014 , 35, 1737-41	3.5	14
25	The Norwegian Multiple Sclerosis Registry and Biobank. <i>Acta Neurologica Scandinavica</i> , 2015 , 132, 24-8	3.8	21
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23	EBNA2 binds to genomic intervals associated with multiple sclerosis and overlaps with vitamin D receptor occupancy. <i>PLoS ONE</i> , 2015 , 10, e0119605	3.7	30
22	Twin studies in multiple sclerosis: A meta-estimation of heritability and environmentality. <i>Multiple Sclerosis Journal</i> , 2015 , 21, 1404-13	5	28
21	Epstein-Barr virus genetic variants are associated with multiple sclerosis. <i>Neurology</i> , 2015 , 84, 1362-8	6.5	29
20	IFN-land multiple sclerosis: from etiology to therapy and back. <i>Cytokine and Growth Factor Reviews</i> , 2015 , 26, 221-8	17.9	20
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17	The relationships among host transcriptional responses reveal distinct signatures underlying viral infection-disease associations. <i>Molecular BioSystems</i> , 2016 , 12, 653-65		
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15	EBV Infection and Vitamin D in Multiple Sclerosis Patients. 2017 , 9-20		1

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14	A Perspective of Coagulation Dysfunction in Multiple Sclerosis and in Experimental Allergic Encephalomyelitis. <i>Frontiers in Neurology</i> , 2018 , 9, 1175	4.1	12
13	Evaluation of variation in genes of the arylhydrocarbon receptor pathway for an association with multiple sclerosis. <i>Journal of Neuroimmunology</i> , 2019 , 334, 576979	3.5	6
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9	Reworking GWAS Data to Understand the Role of Nongenetic Factors in MS Etiopathogenesis. <i>Genes</i> , 2020 , 11,	4.2	2
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7	Association of Epstein-Barr virus latently expressed genes with multiple sclerosis. <i>Multiple Sclerosis and Related Disorders</i> , 2021 , 52, 103008	4	1
6	Multiple Sclerosis and SARS-CoV-2: Has the Interplay Started?. Frontiers in Immunology, 2021, 12, 75533	3 8.4	7
5	Genetic, transcriptome, proteomic and epidemiological evidence for blood brain barrier disruption and polymicrobial brain invasion as determinant factors in Alzheimer disease.		2
4	Significant enrichment of Herpesvirus interactors in GWAS data suggests causal inferences for the association between Epstein Barr virus and multiple sclerosis.		1
3	Prevention of MS Requires Intervention on the Causes of the Disease: Reconciling Genes, Epigenetics, and Epstein Barr Virus <i>Frontiers in Neurology</i> , 2022 , 13, 817677	4.1	0
2	Multiple Sclerosis and Autoimmunity: A Veiled Relationship. Cureus, 2022,	1.2	
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