

# Talin Haritunians

## List of Publications by Year in descending order

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Version: 2024-02-01

53  
papers

15,562  
citations

159585

30  
h-index

175258

52  
g-index

58  
all docs

58  
docs citations

58  
times ranked

27128  
citing authors

#	ARTICLE	IF	CITATIONS
1	Host-microbe interactions have shaped the genetic architecture of inflammatory bowel disease. <i>Nature</i> , 2012, 491, 119-124.	27.8	4,038
2	Association analyses of 249,796 individuals reveal 18 new loci associated with body mass index. <i>Nature Genetics</i> , 2010, 42, 937-948.	21.4	2,634
3	Genome-wide meta-analysis increases to 71 the number of confirmed Crohn's disease susceptibility loci. <i>Nature Genetics</i> , 2010, 42, 1118-1125.	21.4	2,284
4	Hundreds of variants clustered in genomic loci and biological pathways affect human height. <i>Nature</i> , 2010, 467, 832-838.	27.8	1,789
5	Meta-analysis identifies 13 new loci associated with waist-hip ratio and reveals sexual dimorphism in the genetic basis of fat distribution. <i>Nature Genetics</i> , 2010, 42, 949-960.	21.4	836
6	Deep resequencing of GWAS loci identifies independent rare variants associated with inflammatory bowel disease. <i>Nature Genetics</i> , 2011, 43, 1066-1073.	21.4	698
7	Fine-mapping inflammatory bowel disease loci to single-variant resolution. <i>Nature</i> , 2017, 547, 173-178.	27.8	473
8	A common missense variant in NUDT15 confers susceptibility to thiopurine-induced leukopenia. <i>Nature Genetics</i> , 2014, 46, 1017-1020.	21.4	438
9	Functional variants in the <i>LRRK2</i> gene confer shared effects on risk for Crohn's disease and Parkinson's disease. <i>Science Translational Medicine</i> , 2018, 10, .	12.4	273
10	Best Practices and Joint Calling of the HumanExome BeadChip: The CHARGE Consortium. <i>PLoS ONE</i> , 2013, 8, e68095.	2.5	219
11	Reprogramming of gut microbiome energy metabolism by the <i>FUT2</i> Crohn's disease risk polymorphism. <i>ISME Journal</i> , 2014, 8, 2193-2206.	9.8	182
12	Genetic Variants Synthesize to Produce Paneth Cell Phenotypes That Define Subtypes of Crohn's Disease. <i>Gastroenterology</i> , 2014, 146, 200-209.	1.3	155
13	Genetic predictors of medically refractory ulcerative colitis. <i>Inflammatory Bowel Diseases</i> , 2010, 16, 1830-1840.	1.9	135
14	Genome-Wide Association Study Identifies African-Specific Susceptibility Loci in African Americans With Inflammatory Bowel Disease. <i>Gastroenterology</i> , 2017, 152, 206-217.e2.	1.3	120
15	A Pleiotropic Missense Variant in SLC39A8 Is Associated With Crohn's Disease and Human Gut Microbiome Composition. <i>Gastroenterology</i> , 2016, 151, 724-732.	1.3	109
16	Genetic Variants Associated With Quantitative Glucose Homeostasis Traits Translate to Type 2 Diabetes in Mexican Americans: The GUARDIAN (Genetics Underlying Diabetes in Hispanics) Consortium. <i>Diabetes</i> , 2015, 64, 1853-1866.	0.6	77
17	A meta-analysis of 120 246 individuals identifies 18 new loci for fibrinogen concentration. <i>Human Molecular Genetics</i> , 2016, 25, 358-370.	2.9	73
18	Cucurbitacin B induces differentiation, cell cycle arrest, and actin cytoskeletal alterations in myeloid leukemia cells. <i>Leukemia Research</i> , 2008, 32, 1366-1373.	0.8	70

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19	Western diet induces Paneth cell defects through microbiome alterations and farnesoid X receptor and type I interferon activation. <i>Cell Host and Microbe</i> , 2021, 29, 988-1001.e6.	11.0	69
20	Insights into the genetic epidemiology of Crohn's and rare diseases in the Ashkenazi Jewish population. <i>PLoS Genetics</i> , 2018, 14, e1007329.	3.5	66
21	Characterization of Genetic Loci That Affect Susceptibility to Inflammatory Bowel Diseases in African Americans. <i>Gastroenterology</i> , 2015, 149, 1575-1586.	1.3	65
22	A Frameshift in CSF2RB Predominant Among Ashkenazi Jews Increases Risk for Crohn's Disease and Reduces Monocyte Signaling via GM-CSF. <i>Gastroenterology</i> , 2016, 151, 710-723.e2.	1.3	51
23	A protein-truncating R179X variant in RNF186 confers protection against ulcerative colitis. <i>Nature Communications</i> , 2016, 7, 12342.	12.8	50
24	Ocular Manifestations in Inflammatory Bowel Disease Are Associated with Other Extra-intestinal Manifestations, Gender, and Genes Implicated in Other Immune-related Traits. <i>Journal of Crohn's and Colitis</i> , 2016, 10, 43-49.	1.3	50
25	LRRK2 but not ATG16L1 is associated with Paneth cell defect in Japanese Crohn's disease patients. <i>JCI Insight</i> , 2017, 2, e91917.	5.0	46
26	Altered Intestinal ACE2 Levels Are Associated With Inflammation, Severe Disease, and Response to Anti-Cytokine Therapy in Inflammatory Bowel Disease. <i>Gastroenterology</i> , 2021, 160, 809-822.e7.	1.3	45
27	Inflamed Ulcerative Colitis Regions Associated With MRGPRX2-Mediated Mast Cell Degranulation and Cell Activation Modules, Defining a New Therapeutic Target. <i>Gastroenterology</i> , 2021, 160, 1709-1724.	1.3	43
28	Identification of Ten Additional Susceptibility Loci for Ulcerative Colitis Through ImmunoChip Analysis in Koreans. <i>Inflammatory Bowel Diseases</i> , 2016, 22, 13-19.	1.9	40
29	ImmunoChip Meta-Analysis of Inflammatory Bowel Disease Identifies Three Novel Loci and Four Novel Associations in Previously Reported Loci. <i>Journal of Crohn's and Colitis</i> , 2018, 12, 730-741.	1.3	38
30	Genetic Markers Predict Primary Nonresponse and Durable Response to Anti-Tumor Necrosis Factor Therapy in Ulcerative Colitis. <i>Inflammatory Bowel Diseases</i> , 2018, 24, 1840-1848.	1.9	34
31	Pleiotropic ZIP8 A391T implicates abnormal manganese homeostasis in complex human disease. <i>JCI Insight</i> , 2020, 5, .	5.0	34
32	Characterization of candidate genes in inflammatory bowel disease-associated risk loci. <i>JCI Insight</i> , 2016, 1, e87899.	5.0	30
33	Variants in ZNF365 isoform D are associated with Crohn's disease. <i>Gut</i> , 2011, 60, 1060-1067.	12.1	27
34	Whole-genome sequencing of African Americans implicates differential genetic architecture in inflammatory bowel disease. <i>American Journal of Human Genetics</i> , 2021, 108, 431-445.	6.2	21
35	Amino acid position 37 of HLA-DRB1 affects susceptibility to Crohn's disease in Asians. <i>Human Molecular Genetics</i> , 2018, 27, 3901-3910.	2.9	19
36	Transethnic analysis of the human leukocyte antigen region for ulcerative colitis reveals not only shared but also ethnicity-specific disease associations. <i>Human Molecular Genetics</i> , 2021, 30, 356-369.	2.9	19

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37	Linkage Analysis of High-density SNPs Confirms Keratoconus Locus at 5q Chromosomal Region. <i>Ophthalmic Genetics</i> , 2016, 37, 1-2.	1.2	18
38	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. <i>American Journal of Human Genetics</i> , 2021, 108, 564-582.	6.2	18
39	Increased Prevalence of Inflammatory Bowel Disease in Patients with Mutations in Genes Encoding the Receptor Subunits for TGF $\beta$ 2. <i>Inflammatory Bowel Diseases</i> , 2016, 22, 2058-2062.	1.9	15
40	Late-Onset Crohn's Disease Is A Subgroup Distinct in Genetic and Behavioral Risk Factors With UC-Like Characteristics. <i>Inflammatory Bowel Diseases</i> , 2018, 24, 2413-2422.	1.9	14
41	Ileal Gene Expression Data from Crohn's Disease Small Bowel Resections Indicate Distinct Clinical Subgroups. <i>Journal of Crohn's and Colitis</i> , 2019, 13, 1055-1066.	1.3	14
42	Association of NOD2 and IL23R with Inflammatory Bowel Disease in Puerto Rico. <i>PLoS ONE</i> , 2014, 9, e108204.	2.5	14
43	The TNF family member TL1A induces IL-22 secretion in committed human T <sub>H</sub> 17 cells via IL-9 induction. <i>Journal of Leukocyte Biology</i> , 2017, 101, 727-737.	3.3	13
44	An Intergenic Variant rs9268877 Between HLA-DRA and HLA-DRB Contributes to the Clinical Course and Long-term Outcome of Ulcerative Colitis. <i>Journal of Crohn's and Colitis</i> , 2018, 12, 1113-1121.	1.3	12
45	Prevalence and Effect of Genetic Risk of Thromboembolic Disease in Inflammatory Bowel Disease. <i>Gastroenterology</i> , 2021, 160, 771-780.e4.	1.3	11
46	Crohn's disease-associated ATG16L1 T300A genotype is associated with improved survival in gastric cancer. <i>EBioMedicine</i> , 2021, 67, 103347.	6.1	10
47	Serological, genetic and clinical associations with increased health care resource utilization in inflammatory bowel disease. <i>Journal of Digestive Diseases</i> , 2018, 19, 15-23.	1.5	7
48	Genetic associations with adverse events from anti-tumor necrosis factor therapy in inflammatory bowel disease patients. <i>World Journal of Gastroenterology</i> , 2017, 23, 7265-7273.	3.3	7
49	Stratification of risk of progression to colectomy in ulcerative colitis via measured and predicted gene expression. <i>American Journal of Human Genetics</i> , 2021, 108, 1765-1779.	6.2	6
50	Genes in IBD-Associated Risk Loci Demonstrate Genotype-, Tissue-, and Inflammation-Specific Patterns of Expression in Terminal Ileum and Colon Mucosal Tissue. <i>Inflammatory Bowel Diseases</i> , 2016, 22, S1.	1.9	4
51	Improved Performance of Dynamic Measures of Insulin Response Over Surrogate Indices to Identify Genetic Contributors of Type 2 Diabetes: The GUARDIAN Consortium. <i>Diabetes</i> , 2016, 65, 2072-2080.	0.6	4
52	Reply. <i>Gastroenterology</i> , 2017, 152, 2083-2084.	1.3	0
53	Novel Deuterated Gemini-Vitamin D3 Analogs Have Potent Antitumor Activity.. <i>Blood</i> , 2007, 110, 4210-4210.	1.4	0