Bendik S Winsvold

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Association studies of up to 1.2 million individuals yield new insights into the genetic etiology of tobacco and alcohol use. Nature Genetics, 2019, 51, 237-244.	9.4	1,307
2	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	6.0	1,085
3	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. Nature Genetics, 2021, 53, 817-829.	9.4	629
4	Meta-analysis of 375,000 individuals identifies 38 susceptibility loci for migraine. Nature Genetics, 2016, 48, 856-866.	9.4	520
5	A genome-wide association study with 1,126,563 individuals identifies new risk loci for Alzheimer's disease. Nature Genetics, 2021, 53, 1276-1282.	9.4	430
6	Genome-wide meta-analysis identifies new susceptibility loci for migraine. Nature Genetics, 2013, 45, 912-917.	9.4	338
7	Genome-wide association analysis identifies susceptibility loci for migraine without aura. Nature Genetics, 2012, 44, 777-782.	9.4	294
8	Biological and clinical insights from genetics of insomnia symptoms. Nature Genetics, 2019, 51, 387-393.	9.4	250
9	A recurrent de novo mutation in KCNC1 causes progressive myoclonus epilepsy. Nature Genetics, 2015, 47, 39-46.	9.4	245
10	Deciphering osteoarthritis genetics across 826,690 individuals from 9 populations. Cell, 2021, 184, 4784-4818.e17.	13.5	188
11	Genome-wide association study of intracranial aneurysms identifies 17 risk loci and genetic overlap with clinical risk factors. Nature Genetics, 2020, 52, 1303-1313.	9.4	163
12	Genome-wide analysis of 102,084 migraine cases identifies 123 risk loci and subtype-specific risk alleles. Nature Genetics, 2022, 54, 152-160.	9.4	135
13	Genome-wide association analysis of self-reported daytime sleepiness identifies 42 loci that suggest biological subtypes. Nature Communications, 2019, 10, 3503.	5.8	117
14	Genetic Markers of Human Evolution Are Enriched in Schizophrenia. Biological Psychiatry, 2016, 80, 284-292.	0.7	92
15	Shared genetic basis for migraine and ischemic stroke. Neurology, 2015, 84, 2132-2145.	1.5	91
16	Headache, migraine and cardiovascular risk factors: The HUNT study. European Journal of Neurology, 2011, 18, 504-511.	1.7	82
17	Efficacy of antibiotic treatment in patients with chronic low back pain and Modic changes (the AIM) Tj ETQq1 1	0.784314	rgBT /Overlo

18Common Variant Burden Contributes to the Familial Aggregation of Migraine in 1,589 Families. Neuron,
2018, 98, 743-753.e4.3.863

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19	Genetic analysis for a shared biological basis between migraine and coronary artery disease. Neurology: Genetics, 2015, 1, e10.	0.9	61
20	Molecular genetic overlap between migraine and major depressive disorder. European Journal of Human Genetics, 2018, 26, 1202-1216.	1.4	56
21	Gene-based pleiotropy across migraine with aura and migraine without aura patient groups. Cephalalgia, 2016, 36, 648-657.	1.8	47
22	Gene co-expression analysis identifies brain regions and cell types involved in migraine pathophysiology: a GWAS-based study using the Allen Human Brain Atlas. Human Genetics, 2016, 135, 425-439.	1.8	47
23	Experiences of telemedicine in neurological outâ€patient clinics during the COVIDâ€19 pandemic. Annals of Clinical and Translational Neurology, 2021, 8, 440-447.	1.7	46
24	Shared genetic risk between migraine and coronary artery disease: A genome-wide analysis of common variants. PLoS ONE, 2017, 12, e0185663.	1.1	44
25	Blood pressure as a risk factor for headache and migraine: a prospective populationâ€based study. European Journal of Neurology, 2015, 22, 156.	1.7	42
26	Lifestyle factors and risk of migraine and tension-type headache. Follow-up data from the Nord-TrÃ,ndelag Health Surveys 1995–1997 and 2006–2008. Cephalalgia, 2018, 38, 1919-1926.	1.8	41
27	Migraine, obesity and body fat distribution – a population-based study. Journal of Headache and Pain, 2020, 21, 97.	2.5	36
28	Genetic variation in P2RX7 and pain tolerance. Pain, 2018, 159, 1064-1073.	2.0	34
29	Cross-trait analyses with migraine reveal widespread pleiotropy and suggest a vascular component to migraine headache. International Journal of Epidemiology, 2020, 49, 1022-1031.	0.9	34
30	Migraine, headache and development of metabolic syndrome: An 11-year follow-up in the Nord-Trï†ndelag Health Study (HUNT). Pain, 2013, 154, 1305-1311.	2.0	33
31	Concordance of genetic risk across migraine subgroups: Impact on current and future genetic association studies. Cephalalgia, 2015, 35, 489-499.	1.8	32
32	Genome-wide association study identifies <i>RNF123</i> locus as associated with chronic widespread musculoskeletal pain. Annals of the Rheumatic Diseases, 2021, 80, 1227-1235.	0.5	31
33	Genomeâ€Wide Association Study Identifies Risk Loci for Cluster Headache. Annals of Neurology, 2021, 90, 193-202.	2.8	31
34	Dissecting the shared genetic basis of migraine and mental disorders using novel statistical tools. Brain, 2022, 145, 142-153.	3.7	27
35	Epigenetic DNA methylation changes associated with headache chronification: A retrospective case-control study. Cephalalgia, 2018, 38, 312-322.	1.8	25
36	Genome-wide analysis identifies impaired axonogenesis in chronic overlapping pain conditions. Brain, 2022, 145, 1111-1123.	3.7	24

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37	Migraine genetics: from genome-wide association studies to translational insights. Genome Medicine, 2016, 8, 86.	3.6	22
38	The migraine–stroke connection: A genetic perspective. Cephalalgia, 2016, 36, 658-668.	1.8	22
39	Genetic Susceptibility Loci in Genomewide Association Study of Cluster Headache. Annals of Neurology, 2021, 90, 203-216.	2.8	22
40	Hospital-based headache care during the Covid-19 pandemic in Denmark and Norway. Journal of Headache and Pain, 2020, 21, 128.	2.5	21
41	Sex- and age-specific genetic analysis of chronic back pain. Pain, 2021, 162, 1176-1187.	2.0	21
42	Candidate-gene association study searching for genetic factors involved in migraine chronification. Cephalalgia, 2015, 35, 500-507.	1.8	20
43	Mitochondrial genome-wide association study of migraine – the HUNT Study. Cephalalgia, 2020, 40, 625-634.	1.8	19
44	Habitual sleep disturbances and migraine: a Mendelian randomization study. Annals of Clinical and Translational Neurology, 2020, 7, 2370-2380.	1.7	18
45	A re-evaluation of the phonological similarity effect in adults' short-term memory of words and nonwords. Memory, 2001, 9, 281-299.	0.9	16
46	Chronic musculoskeletal complaints as a predictor of mortality—The HUNT study. Pain, 2016, 157, 1443-1447.	2.0	16
47	The causal role of smoking on the risk of hip or knee replacement due to primary osteoarthritis: a Mendelian randomisation analysis of the HUNT study. Osteoarthritis and Cartilage, 2017, 25, 817-823.	0.6	16
48	Genome wide association study identifies variants in NBEA associated with migraine in bipolar disorder. Journal of Affective Disorders, 2015, 172, 453-461.	2.0	15
49	The causal role of smoking on the risk of headache. A Mendelian randomization analysis in the <scp>HUNT</scp> study. European Journal of Neurology, 2018, 25, 1148.	1.7	15
50	Neurology residents' knowledge of the management of headache. Cephalalgia, 2019, 39, 1396-1406.	1.8	13
51	Early menarche and chronic widespread musculoskeletal complaints―Results from the <scp>HUNT</scp> study. European Journal of Pain, 2016, 20, 458-464.	1.4	12
52	Migraine as a predictor of mortality: The HUNT study. Cephalalgia, 2016, 36, 351-357.	1.8	12
53	The impact of C-reactive protein levels on headache frequencyÂin the HUNT study 2006–2008. BMC Neurology, 2019, 19, 229.	0.8	12
54	Migraine and frequent tension-type headache are not associated with multiple sclerosis in a Norwegian case-control study. Multiple Sclerosis Journal - Experimental, Translational and Clinical, 2016, 2, 205521731668297.	0.5	10

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55	Parental migraine in relation to migraine in offspring: Family linkage analyses from the HUNT Study. Cephalalgia, 2019, 39, 854-862.	1.8	10
56	Prestige of neurological disorders among future neurologists in Norway. Acta Neurologica Scandinavica, 2019, 139, 555-558.	1.0	10
57	Clinical effect modifiers of antibiotic treatment in patients with chronic low back pain and Modic changes - secondary analyses of a randomised, placebo-controlled trial (the AIM study). BMC Musculoskeletal Disorders, 2020, 21, 458.	0.8	9
58	The effect of foetal growth restriction on the development of migraine and tension-type headache in adulthood. The HUNT Study. PLoS ONE, 2017, 12, e0175908.	1.1	9
59	Association of Modic change types and their short tau inversion recovery signals with clinical characteristics- a cross sectional study of chronic low back pain patients in the AIM-study. BMC Musculoskeletal Disorders, 2020, 21, 368.	0.8	8
60	Elucidating the relationship between migraine risk and brain structure using genetic data. Brain, 2022, 145, 3214-3224.	3.7	7
61	Urinary albumin excretion as a marker of endothelial dysfunction in migraine sufferers: the HUNT study, Norway. BMJ Open, 2013, 3, e003268.	0.8	6
62	Development and validation of a prediction model for incident hand osteoarthritis in the HUNT study. Osteoarthritis and Cartilage, 2020, 28, 932-940.	0.6	6
63	Acute stroke care during the first phase of COVIDâ€19 pandemic in Norway. Acta Neurologica Scandinavica, 2021, 143, 349-354.	1.0	6
64	The association between selected genetic variants and individual differences in experimental pain. Scandinavian Journal of Pain, 2021, 21, 163-173.	0.5	6
65	Remission of chronic headache: An 11-year follow-up study. Data from the Nord-TrÃ,ndelag Health Surveys 1995–1997 and 2006–2008. Cephalalgia, 2018, 38, 2026-2034.	1.8	5
66	Experiences, distress and burden among neurologists in Norway during the COVID-19 pandemic. PLoS ONE, 2021, 16, e0246567.	1.1	4
67	Genome-Wide Association Study to Identify Common Variants Associated with Brachial Circumference: A Meta-Analysis of 14 Cohorts. PLoS ONE, 2012, 7, e31369.	1.1	3
68	Low Back Pain With Persistent Radiculopathy; the Clinical Role of Genetic Variants in the Genes SOX5, CCDC26/GSDMC and DCC. Frontiers in Genetics, 2021, 12, 757632.	1.1	3
69	OS046. Genome-wide association scans identify novel maternalsusceptibility loci for preeclampsia. Pregnancy Hypertension, 2012, 2, 202.	0.6	2
70	Do incident musculoskeletal complaints influence mortality? The Nord-TrÃ,ndelag Health study. PLoS ONE, 2018, 13, e0203925.	1.1	2
71	The management and clinical knowledge of headache disorders among general practitioners in Norway: a questionnaire survey. Journal of Headache and Pain, 2021, 22, 136.	2.5	2
72	Genome-Wide Association Study of 2,093 Cases With Idiopathic Polyneuropathy and 445,256 Controls Identifies First Susceptibility Loci. Frontiers in Neurology, 2021, 12, 789093.	1.1	2

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73	The causal role of smoking on the risk of hip or knee replacement due to primary osteoarthritis. a mendelian randomization analysis of the nord-trÃ~ndelag health study. Osteoarthritis and Cartilage, 2017, 25, S181-S182.	0.6	1
74	The current state of cluster headache genetics. Headache, 2021, 61, 990-991.	1.8	1
75	Migraine without aura: genome-wide association analysis identifies several novel susceptibility. Journal of Headache and Pain, 2013, 14, .	2.5	0
76	Genetic variation in <i>P2RX7</i> and pain. Scandinavian Journal of Pain, 2016, 12, 127-127.	0.5	0
77	Response to Letter to the Editor: †Comments on the paper presenting prediction models for incident hand OA in the HUNT study'. Osteoarthritis and Cartilage, 2021, 29, 292-293.	0.6	0
78	Migraine and stroke. Tidsskrift for Den Norske Laegeforening, 2018, 138, .	0.2	0
79	Caesarean section and the association with migraine: a retrospective register-linked HUNT population cohort study. BMJ Open, 2020, 10, e040685.	0.8	0
80	Caesarean section and the association with migraine: a retrospective register-linked HUNT population cohort study. BMJ Open, 2020, 10, e040685.	0.8	0
81	A meta-analysis of genome-wide association studies identifies new genetic loci associated with all-cause and vascular dementia. Alzheimer's and Dementia. 2021, 17 Suppl 3, e056081	0.4	0