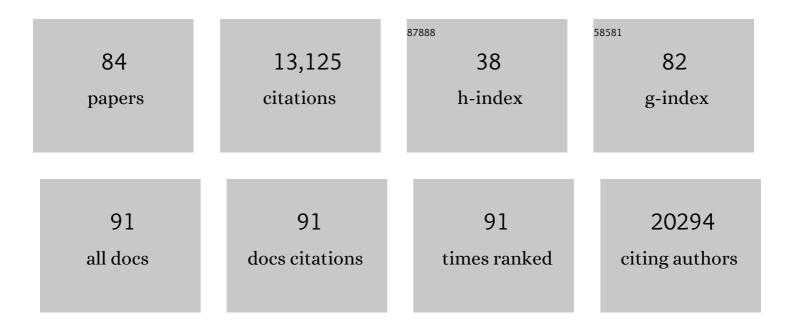
Klaus J Stark

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Large-scale association analysis identifies 13 new susceptibility loci for coronary artery disease. Nature Genetics, 2011, 43, 333-338.	21.4	1,685
2	Large-scale association analysis identifies new risk loci for coronary artery disease. Nature Genetics, 2013, 45, 25-33.	21.4	1,439
3	A large genome-wide association study of age-related macular degeneration highlights contributions of rare and common variants. Nature Genetics, 2016, 48, 134-143.	21.4	1,167
4	Genome-wide association of early-onset myocardial infarction with single nucleotide polymorphisms and copy number variants. Nature Genetics, 2009, 41, 334-341.	21.4	990
5	Meta-analysis identifies 13 new loci associated with waist-hip ratio and reveals sexual dimorphism in the genetic basis of fat distribution. Nature Genetics, 2010, 42, 949-960.	21.4	836
6	Genome-wide meta-analysis identifies 11 new loci for anthropometric traits and provides insights into genetic architecture. Nature Genetics, 2013, 45, 501-512.	21.4	578
7	A genome-wide meta-analysis identifies 22 loci associated with eight hematological parameters in the HaemGen consortium. Nature Genetics, 2009, 41, 1182-1190.	21.4	481
8	New susceptibility locus for coronary artery disease on chromosome 3q22.3. Nature Genetics, 2009, 41, 280-282.	21.4	440
9	Genome-wide haplotype association study identifies the SLC22A3-LPAL2-LPA gene cluster as a risk locus for coronary artery disease. Nature Genetics, 2009, 41, 283-285.	21.4	427
10	FTO genotype is associated with phenotypic variability of body mass index. Nature, 2012, 490, 267-272.	27.8	383
11	Repeated Replication and a Prospective Meta-Analysis of the Association Between Chromosome 9p21.3 and Coronary Artery Disease. Circulation, 2008, 117, 1675-1684.	1.6	356
12	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	27.8	353
13	A genome-wide association study identifies two loci associated with heart failure due to dilated cardiomyopathy. European Heart Journal, 2011, 32, 1065-1076.	2.2	292
14	Association of Early Repolarization Pattern on ECG with Risk of Cardiac and All-Cause Mortality: A Population-Based Prospective Cohort Study (MONICA/KORA). PLoS Medicine, 2010, 7, e1000314.	8.4	246
15	Dysfunctional nitric oxide signalling increases risk of myocardial infarction. Nature, 2013, 504, 432-436.	27.8	230
16	Large-Scale Gene-Centric Analysis Identifies Novel Variants for Coronary Artery Disease. PLoS Genetics, 2011, 7, e1002260.	3.5	203
17	A Genome-Wide Association Study Reveals Variants in ARL15 that Influence Adiponectin Levels. PLoS Genetics, 2009, 5, e1000768.	3.5	148
18	Genetic Regulation of Serum Phytosterol Levels and Risk of Coronary Artery Disease. Circulation: Cardiovascular Genetics, 2010, 3, 331-339.	5.1	141

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19	Genetic variation at chromosome 1p13.3 affects sortilin mRNA expression, cellular LDL-uptake and serum LDL levels which translates to the risk of coronary artery disease. Atherosclerosis, 2010, 208, 183-189.	0.8	141
20	Large Scale Association Analysis of Novel Genetic Loci for Coronary Artery Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2009, 29, 774-780.	2.4	140
21	Lifelong Reduction of LDL-Cholesterol Related to a Common Variant in the LDL-Receptor Gene Decreases the Risk of Coronary Artery Disease—A Mendelian Randomisation Study. PLoS ONE, 2008, 3, e2986.	2.5	137
22	A Genome-Wide Association Study Identifies <i>LIPA</i> as a Susceptibility Gene for Coronary Artery Disease. Circulation: Cardiovascular Genetics, 2011, 4, 403-412.	5.1	130
23	Discovery and prioritization of variants and genes for kidney function in >1.2 million individuals. Nature Communications, 2021, 12, 4350.	12.8	125
24	Genome-wide association study identifies a new locus for coronary artery disease on chromosome 10p11.23. European Heart Journal, 2011, 32, 158-168.	2.2	124
25	Genetic Association Study Identifies HSPB7 as a Risk Gene for Idiopathic Dilated Cardiomyopathy. PLoS Genetics, 2010, 6, e1001167.	3.5	110
26	Exome-wide association study reveals novel susceptibility genes to sporadic dilated cardiomyopathy. PLoS ONE, 2017, 12, e0172995.	2.5	92
27	Common Polymorphisms Influencing Serum Uric Acid Levels Contribute to Susceptibility to Gout, but Not to Coronary Artery Disease. PLoS ONE, 2009, 4, e7729.	2.5	90
28	Heritability of Early Repolarization. Circulation: Cardiovascular Genetics, 2011, 4, 134-138.	5.1	89
29	Lack of Association Between the Trp719Arg Polymorphism in Kinesin-Like Protein-6 and Coronary Artery Disease in 19 Case-Control Studies. Journal of the American College of Cardiology, 2010, 56, 1552-1563.	2.8	84
30	The novel genetic variant predisposing to coronary artery disease in the region of the PSRC1 and CELSR2 genes on chromosome 1 associates with serum cholesterol. Journal of Molecular Medicine, 2008, 86, 1233-1241.	3.9	80
31	Association of Common Polymorphisms in GLUT9 Gene with Gout but Not with Coronary Artery Disease in a Large Case-Control Study. PLoS ONE, 2008, 3, e1948.	2.5	75
32	Longâ€term pattern of brain natriuretic peptide and Nâ€terminal pro brain natriuretic peptide and its determinants in the general population: contribution of age, gender, and cardiac and extraâ€cardiac factors. European Journal of Heart Failure, 2013, 15, 859-867.	7.1	70
33	Meta-analysis of Gene-Level Associations for Rare Variants Based on Single-Variant Statistics. American Journal of Human Genetics, 2013, 93, 236-248.	6.2	60
34	Volvox germline-specific genes that are putative targets of RegA repression encode chloroplast proteins. Current Genetics, 1999, 36, 363-370.	1.7	59
35	Genome-wide association meta-analysis for early age-related macular degeneration highlights novel loci and insights for advanced disease. BMC Medical Genomics, 2020, 13, 120.	1.5	56
36	Genetic pleiotropy between age-related macular degeneration and 16 complex diseases and traits. Genome Medicine, 2017, 9, 29.	8.2	52

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37	Genome-wide association analysis in dilated cardiomyopathy reveals two new players in systolic heart failure on chromosomes 3p25.1 and 22q11.23. European Heart Journal, 2021, 42, 2000-2011.	2.2	49
38	Generation of Highly Purified Human Cardiomyocytes from Peripheral Blood Mononuclear Cell-Derived Induced Pluripotent Stem Cells. PLoS ONE, 2015, 10, e0126596.	2.5	46
39	The Bacterial Paromomycin Resistance Gene, , as a Dominant Selectable Marker in. Protist, 2004, 155, 381-393.	1.5	42
40	Association of common polymorphisms in known susceptibility genes with rheumatoid arthritis in a Slovak population using osteoarthritis patients as controls. Arthritis Research and Therapy, 2009, 11, R70.	3.5	42
41	The lipoprotein subfraction profile: heritability and identification of quantitative trait loci. Journal of Lipid Research, 2008, 49, 715-723.	4.2	41
42	RANTES/CCL5 and Risk for Coronary Events: Results from the MONICA/KORA Augsburg Case-Cohort, Athero-Express and CARDIoGRAM Studies. PLoS ONE, 2011, 6, e25734.	2.5	40
43	Epistatic interaction between haplotypes of the ghrelin ligand and receptor genes influence susceptibility to myocardial infarction and coronary artery disease. Human Molecular Genetics, 2007, 16, 887-899.	2.9	35
44	Retinal Layer Thicknesses in Early Age-Related Macular Degeneration: Results From the German AugUR Study. , 2019, 60, 1581.		34
45	Genetic variation in the arachidonate 5-lipoxygenase-activating protein (<i>ALOX5AP</i>) is associated with myocardial infarction in the German population. Clinical Science, 2008, 115, 309-315.	4.3	32
46	Large-Scale Candidate Gene Analysis of HDL Particle Features. PLoS ONE, 2011, 6, e14529.	2.5	32
47	The German AugUR study: study protocol of a prospective study to investigate chronic diseases in the elderly. BMC Geriatrics, 2015, 15, 130.	2.7	31
48	On the impact of different approaches to classify age-related macular degeneration: Results from the German AugUR study. Scientific Reports, 2018, 8, 8675.	3.3	31
49	Common Genetic Variants in <i>ANK2</i> Modulate QT Interval. Circulation: Cardiovascular Genetics, 2008, 1, 93-99.	5.1	29
50	Association of Smoking, Alcohol Consumption, Blood Pressure, Body Mass Index, and Glycemic Risk Factors With Age-Related Macular Degeneration. JAMA Ophthalmology, 2021, 139, 1299.	2.5	29
51	Genetic associations with lipoprotein subfractions provide information on their biological nature. Human Molecular Genetics, 2012, 21, 1433-1443.	2.9	28
52	Features of Age-Related Macular Degeneration in the General Adults and Their Dependency on Age, Sex, and Smoking: Results from the German KORA Study. PLoS ONE, 2016, 11, e0167181.	2.5	27
53	Lymphotoxin-α and galectin-2 SNPs are not associated with myocardial infarction in two different German populations. Journal of Molecular Medicine, 2007, 85, 997-1004.	3.9	25
54	NT-proBNP Predicts Cardiovascular Death in the General Population Independent of Left Ventricular Mass and Function: Insights from a Large Population-Based Study with Long-Term Follow-Up. PLoS ONE, 2016, 11, e0164060.	2.5	25

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55	The common Y402H variant in complement factor H gene is not associated with susceptibility to myocardial infarction and its related risk factors. Clinical Science, 2007, 113, 213-218.	4.3	24
56	CYB5A polymorphism increases androgens and reduces risk of rheumatoid arthritis in women. Arthritis Research and Therapy, 2015, 17, 56.	3.5	24
57	Identification and MS-assisted interpretation of genetically influenced NMR signals in human plasma. Genome Medicine, 2013, 5, 13.	8.2	23
58	FGF21 signalling pathway and metabolic traits – genetic association analysis. European Journal of Human Genetics, 2010, 18, 1344-1348.	2.8	22
59	Estimates and Determinants of SARS-Cov-2 Seroprevalence and Infection Fatality Ratio Using Latent Class Analysis: The Population-Based Tirschenreuth Study in the Hardest-Hit German County in Spring 2020. Viruses, 2021, 13, 1118.	3.3	22
60	Expression pattern in human macrophages dependent on 9p21.3 coronary artery disease risk locus. Atherosclerosis, 2013, 227, 244-249.	0.8	21
61	Slit3 inhibits Robo3-induced invasion of synovial fibroblasts in rheumatoid arthritis. Arthritis Research and Therapy, 2010, 12, R45.	3.5	19
62	Proteomic Profiling Implies Mitochondrial Dysfunction in Tachycardia-Induced Heart Failure. Journal of Cardiac Failure, 2012, 18, 660-673.	1.7	19
63	Investigating the modulation of genetic effects on late AMD by age and sex: Lessons learned and two additional loci. PLoS ONE, 2018, 13, e0194321.	2.5	19
64	Adiponectin Multimeric Forms but not Total Adiponectin Levels are Associated with Myocardial Infarction in Non-Diabetic Men. Journal of Atherosclerosis and Thrombosis, 2011, 18, 616-627.	2.0	19
65	Association between <i>PPAR</i> α gene polymorphisms and myocardial infarction. Clinical Science, 2008, 115, 301-308.	4.3	17
66	N adherin promoter polymorphisms and risk of osteoarthritis. FASEB Journal, 2014, 28, 683-691.	0.5	15
67	KCND3 potassium channel gene variant confers susceptibility to electrocardiographic early repolarization pattern. JCI Insight, 2019, 4, .	5.0	15
68	Physical Activity, Incidence, and Progression of Age-Related Macular Degeneration: A Multicohort Study. American Journal of Ophthalmology, 2022, 236, 99-106.	3.3	13
69	Lack of Association Between a Common Polymorphism Near the <i>INSIG2</i> Gene and BMI, Myocardial Infarction, and Cardiovascular Risk Factors. Obesity, 2009, 17, 1390-1395.	3.0	12
70	IDGenerator: unique identifier generator for epidemiologic or clinical studies. BMC Medical Research Methodology, 2016, 16, 120.	3.1	12
71	Chances and challenges of machine learningâ€based disease classification in genetic association studies illustrated on ageâ€related macular degeneration. Genetic Epidemiology, 2020, 44, 759-777.	1.3	12
72	Genetic Control of Germ-Soma Differentiation in Volvox carteri. Protist, 2002, 153, 99-107.	1.5	10

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73	Genetics in neuroendocrine immunology: implications for rheumatoid arthritis and osteoarthritis. Annals of the New York Academy of Sciences, 2010, 1193, 10-14.	3.8	8
74	Poor risk factor control in outpatients with diabetes mellitus type 2 in Germany: The DIAbetes COhoRtE (DIACORE) study. PLoS ONE, 2019, 14, e0213157.	2.5	8
75	1,25-dihydroxyvitamin-D3 but not the clinically applied marker 25-hydroxyvitamin-D3 predicts survival after stem cell transplantation. Bone Marrow Transplantation, 2021, 56, 419-433.	2.4	8
76	Incidence, progression and risk factors of age-related macular degeneration in 35–95-year-old individuals from three jointly designed German cohort studies. BMJ Open Ophthalmology, 2022, 7, e000912.	1.6	7
77	Changes in healthcare seeking and lifestyle in old aged individuals during COVID-19 lockdown in Germany: the population-based AugUR study. BMC Geriatrics, 2022, 22, 34.	2.7	7
78	Lack of association of genetic variants in the LRP8 gene with familial and sporadic myocardial infarction. Journal of Molecular Medicine, 2008, 86, 1163-1170.	3.9	6
79	Distribution and specificity of high-sensitivity cardiac troponin T in older adults without acute cardiac conditions: cross-sectional results from the population-based AugUR study. BMJ Open, 2021, 11, e052004.	1.9	6
80	Relative Telomere Length Is Associated With Age-Related Macular Degeneration in Women. , 2022, 63, 30.		6
81	Anti-Thymocyte Globulin Treatment Augments 1,25-Dihydroxyvitamin D3 Serum Levels in Patients Undergoing Hematopoietic Stem Cell Transplantation. Frontiers in Immunology, 2021, 12, 803726.	4.8	3
82	Harmonization of Study and Reference Data by PhaseLift: Saving Time When Imputing Study Data. Genetic Epidemiology, 2014, 38, 381-388.	1.3	1
83	Frequency of hand eczema in the elderly: Crossâ€ s ectional findings from the <scp>German AugUR</scp> study. Contact Dermatitis, 2021, 85, 489-493.	1.4	1
84	Reply to the Letter by Hayashi et al. Circulation: Cardiovascular Genetics, 2011, 4, .	5.1	0