Stefan Böhringer

List of Publications by Year in descending order

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37111 94269 10,624 110 37 96 citations g-index h-index papers 112 112 112 21236 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Genetic studies of body mass index yield new insights for obesity biology. Nature, 2015, 518, 197-206.	13.7	3,823
2	New genetic loci link adipose and insulin biology to body fat distribution. Nature, 2015, 518, 187-196.	13.7	1,328
3	Identification of seven loci affecting mean telomere length and their association with disease. Nature Genetics, 2013, 45, 422-427.	9.4	808
4	Novel Loci for Adiponectin Levels and Their Influence on Type 2 Diabetes and Metabolic Traits: A Multi-Ethnic Meta-Analysis of 45,891 Individuals. PLoS Genetics, 2012, 8, e1002607.	1.5	419
5	A Genome-Wide Association Study Identifies Five Loci Influencing Facial Morphology in Europeans. PLoS Genetics, 2012, 8, e1002932.	1.5	274
6	Genome-wide association meta-analysis of human longevity identifies a novel locus conferring survival beyond 90 years of age. Human Molecular Genetics, 2014, 23, 4420-4432.	1.4	227
7	A Meta-Analysis of Thyroid-Related Traits Reveals Novel Loci and Gender-Specific Differences in the Regulation of Thyroid Function. PLoS Genetics, 2013, 9, e1003266.	1.5	194
8	A comprehensive molecular study on Coffin–Siris and Nicolaides–Baraitser syndromes identifies a broad molecular and clinical spectrum converging on altered chromatin remodeling. Human Molecular Genetics, 2013, 22, 5121-5135.	1.4	190
9	Oculo-auriculo-vertebral spectrum (OAVS): clinical evaluation and severity scoring of 53 patients and proposal for a new classification. European Journal of Medical Genetics, 2005, 48, 397-411.	0.7	184
10	Genome-wide association study identifies novel genetic variants contributing to variation in blood metabolite levels. Nature Communications, 2015, 6, 7208.	5.8	178
11	Genome-wide meta-analysis uncovers novel loci influencing circulating leptin levels. Nature Communications, 2016, 7, 10494.	5.8	153
12	Genotyping in 46 patients with tentative diagnosis of Treacher Collins syndrome revealed unexpected phenotypic variation. European Journal of Human Genetics, 2004, 12, 879-890.	1.4	149
13	A novel real-time PCR assay for quantitative analysis of methylated alleles (QAMA): analysis of the retinoblastoma locus. Nucleic Acids Research, 2004, 32, e125-e125.	6.5	120
14	Gene set analysis of GWAS data for human longevity highlights the relevance of the insulin/IGF-1 signaling and telomere maintenance pathways. Age, 2013, 35, 235-249.	3.0	105
15	Genetic determination of human facial morphology: links between cleft-lips and normal variation. European Journal of Human Genetics, 2011, 19, 1192-1197.	1.4	89
16	Quaking promotes monocyte differentiation into pro-atherogenic macrophages by controlling pre-mRNA splicing and gene expression. Nature Communications, 2016, 7, 10846.	5.8	87
17	Validation of genetic modifiers for Duchenne muscular dystrophy: a multicentre study assessing <i>SPP1</i> and <i>LTBP4</i> variants. Journal of Neurology, Neurosurgery and Psychiatry, 2015, 86, 1060-1065.	0.9	86
18	A family with autosomal dominant oculo-auriculo-vertebral spectrum. Clinical Dysmorphology, 2007, 16, 1-7.	0.1	81

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19	Triple Positivity for Anti–Citrullinated Protein Autoantibodies, Rheumatoid Factor, and Anti–Carbamylated Protein Antibodies Conferring High Specificity for Rheumatoid Arthritis. Arthritis and Rheumatology, 2018, 70, 1721-1731.	2.9	81
20	Ensemble landmarking of 3D facial surface scans. Scientific Reports, 2018, 8, 12.	1.6	78
21	Results of a 6-week treatment with 10 mg prednisolone in patients with hand osteoarthritis (HOPE): a double-blind, randomised, placebo-controlled trial. Lancet, The, 2019, 394, 1993-2001.	6.3	78
22	Syndrome identification based on 2D analysis software. European Journal of Human Genetics, 2006, 14, 1082-1089.	1.4	77
23	The Prognostic Value of AJCC Staging in Uveal Melanoma Is Enhanced by Adding Chromosome 3 and 8q Status. , 2017, 58, 833.		77
24	Somatic mosaicism in patients with Angelman syndrome and an imprinting defect. Human Molecular Genetics, 2004, 13, 2547-2555.	1.4	74
25	A metabolomic profile is associated with the risk of incident coronary heart disease. American Heart Journal, 2014, 168, 45-52.e7.	1.2	74
26	Genomewide Linkage Screen for Waldenstr \tilde{A} ¶m Macroglobulinemia Susceptibility Loci in High-Risk Families. American Journal of Human Genetics, 2006, 79, 695-701.	2.6	72
27	Tamoxifen Pharmacogenetics and Metabolism: Results From the Prospective CYPTAM Study. Journal of Clinical Oncology, 2019, 37, 636-646.	0.8	72
28	Five-Year Graft Survival and Clinical Outcomes of 500 Consecutive Cases After Descemet Membrane Endothelial Keratoplasty. Cornea, 2020, 39, 290-297.	0.9	69
29	Histone acetylation dependent allelic expression imbalance of BAPX1 in patients with the oculo-auriculo-vertebral spectrum. Human Molecular Genetics, 2006, 15, 581-587.	1.4	68
30	The ACPA recognition profile and subgrouping of ACPA-positive RA patients. Annals of the Rheumatic Diseases, 2012, 71, 268-274.	0.5	61
31	Genes expressed in blood link osteoarthritis with apoptotic pathways. Annals of the Rheumatic Diseases, 2014, 73, 1844-1853.	0.5	61
32	Novel genetic loci affecting facial shape variation in humans. ELife, 2019, 8, .	2.8	58
33	Treat to target (drug-free) inactive disease in DMARD-naive juvenile idiopathic arthritis: 24-month clinical outcomes of a three-armed randomised trial. Annals of the Rheumatic Diseases, 2019, 78, 51-59.	0.5	56
34	Long-Term Graft Survival in Penetrating Keratoplasty: The Biexponential Model of Chronic Endothelial Cell Loss Revisited. Cornea, 2010, 29, 1113-1117.	0.9	55
35	The effect of a face mask for respiratory support on breathing in preterm infants at birth. Resuscitation, 2019, 144, 178-184.	1.3	48
36	Reproduction abnormalities and twin pregnancies in parents of sporadic patients with oculo-auriculo-vertebral spectrum/Goldenhar syndrome. Human Genetics, 2007, 121, 369-376.	1.8	46

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37	Automated segmentation of the corneal endothelium in a large set of â€real-world' specular microscopy images using the U-Net architecture. Scientific Reports, 2019, 9, 4752.	1.6	41
38	Association Between <i>CNDP1</i> Genotype and Diabetic Nephropathy Is Sex Specific. Diabetes, 2010, 59, 1555-1559.	0.3	39
39	Distribution of GNAQ and GNA11 Mutation Signatures in Uveal Melanoma Points to a Light Dependent Mutation Mechanism. PLoS ONE, 2015, 10, e0138002.	1.1	39
40	Automated syndrome detection in a set of clinical facial photographs. American Journal of Medical Genetics, Part A, 2011, 155, 2161-2169.	0.7	38
41	TCTEX1D1 is a genetic modifier of disease progression in Duchenne muscular dystrophy. European Journal of Human Genetics, 2020, 28, 815-825.	1.4	36
42	Epistasis between two HLA antigens defines a subset of individuals at a very high risk for ankylosing spondylitis. Annals of the Rheumatic Diseases, 2013, 72, 974-978.	0.5	35
43	An Automatic 3D Facial Landmarking Algorithm Using 2D Gabor Wavelets. IEEE Transactions on Image Processing, 2016, 25, 580-588.	6.0	31
44	Effect of Surgical Indication and Preoperative Lens Status on Descemet Membrane Endothelial Keratoplasty Outcomes. American Journal of Ophthalmology, 2020, 212, 79-87.	1.7	31
45	Insulin-like growth factor 1 receptor expression and IGF1R 3129G > T polymorphism are associated with response to neoadjuvant chemotherapy in breast cancer patients: results from the NEOZOTAC trial (BOOG 2010-01). Breast Cancer Research, 2016, 18, 3.	2.2	30
46	Impact of geometry and viewing angle on classification accuracy of 2D based analysis of dysmorphic faces. European Journal of Medical Genetics, 2008, 51, 44-53.	0.7	29
47	MicroRNA Classifier and Nomogram for Metastasis Prediction in Colon Cancer. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 187-197.	1.1	28
48	Novel quantitative pigmentation phenotyping enhances genetic association, epistasis, and prediction of human eye colour. Scientific Reports, 2017, 7, 43359.	1.6	27
49	A genome-wide association study identifies genetic loci associated with specific lobar brain volumes. Communications Biology, 2019, 2, 285.	2.0	27
50	How to deal with the early GWAS data when imputing and combining different arrays is necessary. European Journal of Human Genetics, 2012, 20, 572-576.	1.4	26
51	Generating evidence for precision medicine: considerations made by the Ubiquitous Pharmacogenomics Consortium when designing and operationalizing the PREPARE study. Pharmacogenetics and Genomics, 2020, 30, 131-144.	0.7	26
52	Genetic associations and regulation of expression indicate an independent role for 14q32 snoRNAs in human cardiovascular disease. Cardiovascular Research, 2019, 115, 1519-1532.	1.8	25
53	Suppression of compensatory erythropoiesis in hemolytic disease of the fetus and newborn due to intrauterine transfusions. American Journal of Obstetrics and Gynecology, 2020, 223, 119.e1-119.e10.	0.7	24
54	HLA class I/II matching and chronic endothelial cell loss in penetrating normal risk keratoplasty. Acta Ophthalmologica, 2004, 82, 13-18.	0.4	23

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55	Complement receptor 1 gene polymorphisms are associated with cardiovascular risk. Atherosclerosis, 2017, 257, 16-21.	0.4	22
56	Exploratory analysis of 1936 SNPs in ADME genes for association with busulfan clearance in adult hematopoietic stem cell recipients. Pharmacogenetics and Genomics, 2013, 23, 675-683.	0.7	17
57	Personalised drug repositioning for Clear Cell Renal Cell Carcinoma using gene expression. Scientific Reports, 2018, 8, 5250.	1.6	14
58	Genome Wide Association Study for Predictors of Progression Free Survival in Patients on Capecitabine, Oxaliplatin, Bevacizumab and Cetuximab in First-Line Therapy of Metastatic Colorectal Cancer. PLoS ONE, 2015, 10, e0131091.	1.1	13
59	Age-dependent clinical prognostic value of histone modifications in colorectal cancer. Translational Research, 2015, 165, 578-588.	2.2	12
60	Influence of batch effect correction methods on drug induced differential gene expression profiles. BMC Bioinformatics, 2019, 20, 437.	1.2	12
61	Prediction of aspiration in dysphagia using logistic regression: oral intake and self-evaluation. European Archives of Oto-Rhino-Laryngology, 2020, 277, 197-205.	0.8	12
62	Acquired Melanocytic Nevi in Childhood and Familial Melanoma. JAMA Dermatology, 2014, 150, 35.	2.0	11
63	Automated human skull landmarking with 2D Gabor wavelets. Physics in Medicine and Biology, 2018, 63, 105011.	1.6	11
64	Quantification of Facial Traits. Frontiers in Genetics, 2019, 10, 397.	1.1	11
65	Quality of life of children with hearing loss in special and mainstream education: A longitudinal study. International Journal of Pediatric Otorhinolaryngology, 2020, 128, 109701.	0.4	11
66	Frequent mutated <i>B2M</i> , <i>EZH2</i> , <i>IRF8</i> , and <i>TNFRSF14</i> in primary bone diffuse large B-cell lymphoma reflect a GCB phenotype. Blood Advances, 2021, 5, 3760-3775.	2.5	11
67	Correcting for multiple testing in genetic association studies: the legend lives on. Human Genetics, 2001, 109, 566-567.	1.8	10
68	Classification and Visualization Based on Derived Image Features: Application to Genetic Syndromes. PLoS ONE, 2014, 9, e109033.	1.1	9
69	Voice outcome after unilateral ELS type III or bilateral type II resections for T1â€₹2 glottic carcinoma: Results after 1 year. Head and Neck, 2019, 41, 1638-1647.	0.9	9
70	Improving the aseptic transfer procedures in hospital pharmacies. Part B: evaluation of disinfection methods for materials with a non-sterile surface. European Journal of Hospital Pharmacy, 2021, 28, 271-275.	0.5	9
71	Transcriptome Signature Reversion as a Method to Reposition Drugs Against Cancer for Precision Oncology. Cancer Journal (Sudbury, Mass), 2019, 25, 116-120.	1.0	9
72	Distinctive facial features in idiopathic Moyamoya disease in Caucasians: a first systematic analysis. Peerl, 2018, 6, e4740.	0.9	8

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73	Multilocus statistics to uncover epistasis and heterogeneity in complex diseases: revisiting a set of multiple sclerosis data. European Journal of Human Genetics, 2003, 11, 573-584.	1.4	7
74	A Genetic Polymorphism in <i>CTLA-4</i> Is Associated with Overall Survival in Sunitinib-Treated Patients with Clear Cell Metastatic Renal Cell Carcinoma. Clinical Cancer Research, 2018, 24, 2350-2356.	3.2	7
75	Genome wide association study to identify predictors for severe skin toxicity in colorectal cancer patients treated with cetuximab. PLoS ONE, 2018, 13, e0208080.	1.1	7
76	Powerful testing via hierarchical linkage disequilibrium in haplotype association studies. Biometrical Journal, 2019, 61, 747-768.	0.6	7
77	Channel discrimination along all contacts of the cochlear implant electrode array and its relation to speech perception. International Journal of Audiology, 2019, 58, 262-268.	0.9	7
78	The impact of estimated tumour purity on gene expression-based drug repositioning of Clear Cell Renal Cell Carcinoma samples. Scientific Reports, 2019, 9, 2495.	1.6	7
79	Comparing the effect of two different interfaces on breathing of preterm infants at birth: A matched-pairs analysis. Resuscitation, 2020, 157, 60-66.	1.3	7
80	MiRNAs Correlate with HLA Expression in Uveal Melanoma: Both Up- and Downregulation Are Related to Monosomy 3. Cancers, 2021, 13, 4020.	1.7	7
81	Pathway analysis to identify genetic variants associated with efficacy of adalimumab in rheumatoid arthritis. Pharmacogenomics, 2017, 18, 945-953.	0.6	6
82	Noise Exposure and Hearing Loss among Brewery Workers in Lagos, Nigeria. International Journal of Environmental Research and Public Health, 2020, 17, 2880.	1.2	6
83	Prognostic Factors for the Outcome of Translabyrinthine Surgery for Vestibular Schwannomas. Otology and Neurotology, 2021, 42, 475-482.	0.7	6
84	Time to Functional Recovery After Laser Tonsillotomy Performed Under Local Anesthesia vs Conventional Tonsillectomy With General Anesthesia Among Adults. JAMA Network Open, 2022, 5, e2148655.	2.8	6
85	Reconstruction of images from Gabor graphs with applications in facial image processing. International Journal of Wavelets, Multiresolution and Information Processing, 2015, 13, 1550019.	0.9	5
86	Ouabain Does Not Induce Selective Spiral Ganglion Cell Degeneration in Guinea Pigs. BioMed Research International, 2018, 2018, 1-15.	0.9	5
87	Effectiveness of Phantom Stimulation in Shifting the Pitch Percept in Cochlear Implant Users. Ear and Hearing, 2020, 41, 1258-1269.	1.0	5
88	The variant T allele of Pvull in ESR1 gene is a prognostic marker in early breast cancer survival. Scientific Reports, 2021, 11, 3249.	1.6	5
89	Genetic variants in Cell Adhesion Molecule 1 (CADM1): A validation study of a novel endothelial cell venous thrombosis risk factor. Thrombosis Research, 2014, 134, 1186-1192.	0.8	4
90	Improving the aseptic transfer procedures in hospital pharmacies part C: evaluation and redesign of the transfer process. European Journal of Hospital Pharmacy, 2022, 29, 12-17.	0.5	4

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91	Identification of pharmacogenetic biomarkers for efficacy of cytoreductive surgery plus hyperthermic intraperitoneal mitomycin C in patients with colorectal peritoneal metastases. European Journal of Surgical Oncology, 2020, 46, 1925-1931.	0.5	4
92	Blood pressure, antihypertensive medication and neuropsychiatric symptoms in older people with dementia: The COSMOS study. International Journal of Geriatric Psychiatry, 2021, 36, 46-53.	1.3	4
93	Changes in healthcare utilisation for paediatric tonsillectomy and adenoidectomy in the Netherlands: a populationâ€based study. Clinical Otolaryngology, 2021, 46, 347-356.	0.6	4
94	Choosing and changing the analysis scale in non-inferiority trials with a binary outcome. Clinical Trials, 2021, , 174077452110537.	0.7	4
95	Nationwide inventory on retinopathy of prematurity screening in the Netherlands. British Journal of Ophthalmology, 2023, 107, 712-716.	2.1	4
96	Improving the aseptic transfer procedures in hospital pharmacies part A: methods for the determination of the surface bioburden on ampoules and vials. European Journal of Hospital Pharmacy, 2021, 28, 38-41.	0.5	3
97	Modelling growth curves of the normal infant's mandible: 3D measurements using computed tomography. Clinical Oral Investigations, 2021, 25, 6365-6375.	1.4	3
98	Feasibility and Effect of Physiological-Based CPAP in Preterm Infants at Birth. Frontiers in Pediatrics, 2021, 9, 777614.	0.9	3
99	A prediction model for recurrence after translabyrinthine surgery for vestibular schwannoma: toward personalized postoperative surveillance. European Archives of Oto-Rhino-Laryngology, 2022, , 1.	0.8	3
100	A Model for Fine Mapping in Family Based Association Studies. Human Heredity, 2009, 67, 226-236.	0.4	2
101	Combining Family and Twin Data in Association Studies to Estimate the Noninherited Maternal Antigens Effect. Genetic Epidemiology, 2012, 36, 811-819.	0.6	2
102	Combining information from linkage and association mapping for next-generation sequencing longitudinal family data. BMC Proceedings, 2014, 8, S34.	1.8	2
103	<i>SLC04A1</i> , <i>SLC22A2</i> and <i>SLC28A2</i> variants not related to methotrexate efficacy or toxicity in rheumatoid arthritis patients. Pharmacogenomics, 2018, 19, 613-619.	0.6	2
104	Reply to C.L. Braal et al, H. Brauch et al, and M.P. Goetz et al. Journal of Clinical Oncology, 2019, 37, 1984-1985.	0.8	2
105	Effectiveness of endolymphatic duct blockage versus endolymphatic sac decompression in patients with intractable MéniÃ⁻re's disease: study protocol for a double-blinded, randomised controlled trial. BMJ Open, 2021, 11, e054514.	0.8	2
106	Comments on: Hierarchical inference for genome-wide association studies by Jelle J. Goeman and Stefan Böhringer. Computational Statistics, 2020, 35, 41-45.	0.8	1
107	Genetic variants determining survival and fertility in an adverse African environment: a population-based large-scale candidate gene association study. Aging, 2016, 8, 1364-1383.	1.4	1
108	Response to a letter to the editor "A prediction model for recurrence after translabyrinthine surgery for vestibular schwannoma: towards personalized postoperative surveillance― European Archives of Oto-Rhino-Laryngology, 2022, , 1.	0.8	1

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109	OP0180â€LOW-DOSE PREDNISOLONE IN PATIENTS WITH HAND OSTEOARTHRITIS (HOPE): RESULTS FROM A RANDOMISED DOUBLE-BLIND PLACEBO-CONTROLLED TRIAL., 2019, , .		0
110	Novel Genetic Loci Affecting Facial Shape Variation in Humans. SSRN Electronic Journal, 0, , .	0.4	0