Anne C BÃ¶hmer

List of Publications by Year
in descending order

25 papers	1,629 citations	567281 15 h-index	526287 27 g-index
27 all docs	27 docs citations	27 times ranked	3838 citing authors

#	Article	IF	Citations
1	Multitrait genetic association analysis identifies 50 new risk loci for gastro-oesophageal reflux, seven new loci for Barrettâ€™s oesophagus and provides insights into clinical heterogeneity in reflux diagnosis. Gut, 2022, 71, 1053-1061.	12.1	74
2	Germline variation in the insulin-like growth factor pathway and risk of Barrettâ€™s esophagus and esophageal adenocarcinoma. Carcinogenesis, 2021, 42, 369-377.	2.8	11
3	A comprehensive re-assessment of the association between vitamin D and cancer susceptibility using Mendelian randomization. Nature Communications, 2021, 12, 246.	12.8	39
4	Sex-Specific Genetic Associations for Barrettâ€™s Esophagus and Esophageal Adenocarcinoma. Gastroenterology, 2020, 159, 2065-2076.e1.	1.3	16
5	Shared Genetic Etiology of Obesity-Related Traits and Barrett's Esophagus/Adenocarcinoma: Insights from Genome-Wide Association Studies. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 427-433.	2.5	7
6	Gastroesophageal reflux GWAS identifies risk loci that also associate with subsequent severe esophageal diseases. Nature Communications, 2019, 10, 4219.	12.8	58
7	No Association Between Vitamin D Status and Risk of Barrett's Esophagus or Esophageal Adenocarcinoma: A Mendelian Randomization Study. Clinical Gastroenterology and Hepatology, 2019, 17, 2227-2235.e1.	4.4	16
8	Identification of loci of functional relevance to Barrettâ€™s esophagus and esophageal adenocarcinoma: Cross-referencing of expression quantitative trait loci data from disease-relevant tissues with genetic association data. PLoS ONE, 2019, 14, e0227072.	2.5	5
9	Nonsyndromic cleft palate: An association study at GWAS candidate loci in a multiethnic sample. Birth Defects Research, 2018, 110, 871-882.	1.5	11
10	Genetic estimators of DNA methylation provide insights into the molecular basis of polygenic traits. Translational Psychiatry, 2018, 8, 31.	4.8	12
11	Investigation of dominant and recessive inheritance models in genomeâ€wide association studies data of nonsyndromic cleft lip with or without cleft palate. Birth Defects Research, 2018, 110, 336-341.	1.5	8
12	Evidence for <scp>PTGER</scp>4,<scp>PSCA</scp>, and <scp>MBOAT</scp>7 as risk genes for gastric cancer on the genome and transcriptome level. Cancer Medicine, 2018, 7, 5057-5065.	2.8	22
13	Study of 300,486 individuals identifies 148 independent genetic loci influencing general cognitive function. Nature Communications, 2018, 9, 2098.	12.8	484
14	Imputation of Orofacial Clefting Data Identifies Novel Risk Loci and Sheds Light on the Genetic Background of Cleft Lip Â± Cleft Palate and Cleft Palate Only.. Human Molecular Genetics, 2017, 26, ddx012.	2.9	84
15	Meta-analysis Reveals Genome-Wide Significance at 15q13 for Nonsyndromic Clefting of Both the Lip and the Palate, and Functional Analyses Implicate GREM1 As a Plausible Causative Gene. PLoS Genetics, 2016, 12, e1005914.	3.5	66
16	The Barrettâ€associated variants at <scp>GDF</scp>7 and <scp>TBX</scp>5 also increase esophageal adenocarcinoma risk. Cancer Medicine, 2016, 5, 888-891.	2.8	21
17	Genome-wide association studies in oesophageal adenocarcinoma and Barrett's oesophagus: a large-scale meta-analysis. Lancet Oncology, The, 2016, 17, 1363-1373.	10.7	133
18	Replication analysis of 15 susceptibility loci for nonsyndromic cleft lip with or without cleft palate in an italian population. Birth Defects Research Part A: Clinical and Molecular Teratology, 2016, 106, 81-87.	1.6	10

#	Article	IF	Citations
19	Sequencing the GRHL3 Coding Region Reveals Rare Truncating Mutations and a Common Susceptibility Variant for Nonsyndromic Cleft Palate. American Journal of Human Genetics, 2016, 98, 755-762.	6.2	92
20	Genome-wide analysis of parent-of-origin effects in non-syndromic orofacial clefts. European Journal of Human Genetics, 2014, 22, 822-830.	2.8	12
21	Investigation of the involvement of MIR185 and its target genes in the development of schizophrenia. Journal of Psychiatry and Neuroscience, 2014, 39, 386-396.	2.4	23
22	Compound Heterozygosity of Low-Frequency Promoter Deletions and Rare Loss-of-Function Mutations in TXNL4A Causes Burn-McKeown Syndrome. American Journal of Human Genetics, 2014, 95, 698-707.	6.2	55
23	Nonsyndromic cleft lip with or without cleft palate in arab populations: Genetic analysis of 15 risk loci in a novel caseâ€“control sample recruited in Yemen. Birth Defects Research Part A: Clinical and Molecular Teratology, 2014, 100, 307-313.	1.6	26
24	Analysis of susceptibility loci for nonsyndromic orofacial clefting in a European trio sample. American Journal of Medical Genetics, Part A, 2013, 161, 2545-2549.	1.2	21
25	Genome-wide meta-analyses of nonsyndromic cleft lip with or without cleft palate identify six new risk loci. Nature Genetics, 2012, 44, 968-971.	21.4	311