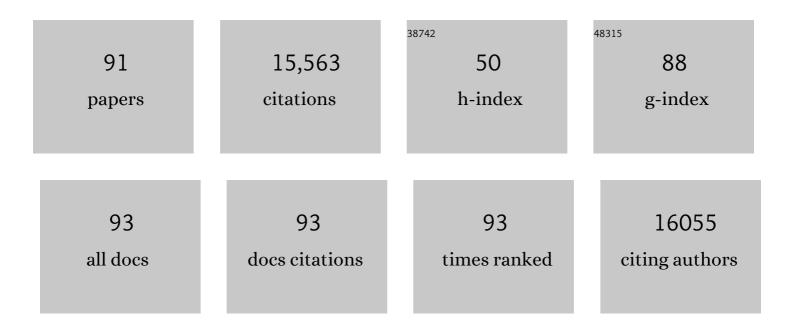
## Moritz F Sinner

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	2020 ESC Guidelines for the diagnosis and management of atrial fibrillation developed in collaboration with the European Association for Cardio-Thoracic Surgery (EACTS). European Heart Journal, 2021, 42, 373-498.	2.2	5,583
2	Simple Risk Model Predicts Incidence of Atrial Fibrillation in a Racially and Geographically Diverse Population: the CHARGEâ€AF Consortium. Journal of the American Heart Association, 2013, 2, e000102.	3.7	601
3	Multi-ethnic genome-wide association study for atrial fibrillation. Nature Genetics, 2018, 50, 1225-1233.	21.4	552
4	Meta-analysis identifies six new susceptibility loci for atrial fibrillation. Nature Genetics, 2012, 44, 670-675.	21.4	533
5	Incidence and Prevalence of Atrial Fibrillation and Associated Mortality Among Medicare Beneficiaries: 1993–2007. Circulation: Cardiovascular Quality and Outcomes, 2012, 5, 85-93.	2.2	476
6	Common variants in KCNN3 are associated with lone atrial fibrillation. Nature Genetics, 2010, 42, 240-244.	21.4	438
7	Genome-wide association study of PR interval. Nature Genetics, 2010, 42, 153-159.	21.4	400
8	Variants in ZFHX3 are associated with atrial fibrillation in individuals of European ancestry. Nature Genetics, 2009, 41, 879-881.	21.4	363
9	Common variants at ten loci modulate the QT interval duration in the QTSCD Study. Nature Genetics, 2009, 41, 407-414.	21.4	356
10	Common variants in 22 loci are associated with QRS duration and cardiac ventricular conduction. Nature Genetics, 2010, 42, 1068-1076.	21.4	308
11	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. Nature Genetics, 2014, 46, 826-836.	21.4	281
12	Large-scale analyses of common and rare variants identify 12 new loci associated with atrial fibrillation. Nature Genetics, 2017, 49, 946-952.	21.4	279
13	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
14	Atrial Fibrillation. Circulation, 2011, 124, 1982-1993.	1.6	225
15	MicroRNA29. Circulation, 2013, 127, 1466-1475.	1.6	222
16	Large scale replication and meta-analysis of variants on chromosome 4q25 associated with atrial fibrillation. European Heart Journal, 2008, 30, 813-819.	2.2	193
17	Searching for Atrial Fibrillation Poststroke. Circulation, 2019, 140, 1834-1850.	1.6	184
18	Integrating Genetic, Transcriptional, and Functional Analyses to Identify 5 Novel Genes for Atrial Fibrillation. Circulation, 2014, 130, 1225-1235.	1.6	183

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19	Symptoms and Functional Status of Patients With Atrial Fibrillation. Circulation, 2012, 125, 2933-2943.	1.6	175
20	P-wave duration and the risk of atrial fibrillation: Results from the Copenhagen ECG Study. Heart Rhythm, 2015, 12, 1887-1895.	0.7	152
21	A Large Candidate Gene Survey Identifies the <i>KCNE1</i> D85N Polymorphism as a Possible Modulator of Drug-Induced Torsades de Pointes. Circulation: Cardiovascular Genetics, 2012, 5, 91-99.	5.1	150
22	Clinical course of atrial fibrillation in older adults: the importance of cardiovascular events beyond stroke. European Heart Journal, 2014, 35, 250-256.	2.2	148
23	B-type natriuretic peptide and C-reactive protein in the prediction of atrial fibrillation risk: the CHARGE-AF Consortium of community-based cohort studies. Europace, 2014, 16, 1426-1433.	1.7	144
24	Outcomes of Medicare Beneficiaries Undergoing Catheter Ablation for Atrial Fibrillation. Circulation, 2012, 126, 2200-2207.	1.6	138
25	Independent Susceptibility Markers for Atrial Fibrillation on Chromosome 4q25. Circulation, 2010, 122, 976-984.	1.6	137
26	A genome-wide association study identifies 6p21 as novel risk locus for dilated cardiomyopathy. European Heart Journal, 2014, 35, 1069-1077.	2.2	137
27	Variants in the 3′ untranslated region of the KCNQ1-encoded Kv7.1 potassium channel modify disease severity in patients with type 1 long QT syndrome in an allele-specific manner. European Heart Journal, 2012, 33, 714-723.	2.2	130
28	Novel Genetic Markers Associate With Atrial Fibrillation Risk in Europeans and Japanese. Journal of the American College of Cardiology, 2014, 63, 1200-1210.	2.8	127
29	Defining the major health modifiers causing atrial fibrillation: a roadmap to underpin personalized prevention and treatment. Nature Reviews Cardiology, 2016, 13, 230-237.	13.7	122
30	A roadmap to improve the quality of atrial fibrillation management: proceedings from the fifth Atrial Fibrillation Network/European Heart Rhythm Association consensus conference. Europace, 2016, 18, 37-50.	1.7	121
31	52 Genetic Loci Influencing MyocardialÂMass. Journal of the American College of Cardiology, 2016, 68, 1435-1448.	2.8	113
32	Incidence of complications related to catheter ablation of atrial fibrillation and atrial flutter: a nationwide in-hospital analysis of administrative data for Germany in 2014. European Heart Journal, 2018, 39, 4020-4029.	2.2	108
33	Preventive or Deferred Ablation of Ventricular Tachycardia in Patients With Ischemic Cardiomyopathy and Implantable Defibrillator (BERLIN VT). Circulation, 2020, 141, 1057-1067.	1.6	104
34	The non-synonymous coding IKr-channel variant KCNH2-K897T is associated with atrial fibrillation: results from a systematic candidate gene-based analysis of KCNH2 (HERG). European Heart Journal, 2008, 29, 907-914.	2.2	103
35	Determination and Interpretation of the QT Interval. Circulation, 2018, 138, 2345-2358.	1.6	100
36	Common Genetic Variants and Response to Atrial Fibrillation Ablation. Circulation: Arrhythmia and Electrophysiology, 2015, 8, 296-302.	4.8	98

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37	Usefulness of Short-Term Variability of QT Intervals as a Predictor for Electrical Remodeling and Proarrhythmia in Patients With Nonischemic Heart Failure. American Journal of Cardiology, 2010, 106, 216-220.	1.6	96
38	Genetic loci associated with heart rate variability and their effects on cardiac disease risk. Nature Communications, 2017, 8, 15805.	12.8	95
39	Integrating new approaches to atrial fibrillation management: the 6th AFNET/EHRA Consensus Conference. Europace, 2018, 20, 395-407.	1.7	95
40	Relation of Circulating Liver Transaminase Concentrations to Risk of New-Onset Atrial Fibrillation. American Journal of Cardiology, 2013, 111, 219-224.	1.6	85
41	Mutations in the mitochondrial thioredoxin reductase gene TXNRD2 cause dilated cardiomyopathy. European Heart Journal, 2011, 32, 1121-1133.	2.2	84
42	Clinical effectiveness of primary prevention implantable cardioverter-defibrillators: results of the EU-CERT-ICD controlled multicentre cohort study. European Heart Journal, 2020, 41, 3437-3447.	2.2	78
43	Large-Scale Candidate Gene Analysis in Whites and African Americans Identifies <i>IL6R</i> Polymorphism in Relation to Atrial Fibrillation. Circulation: Cardiovascular Genetics, 2011, 4, 557-564.	5.1	74
44	Genome-wide association studies of atrial fibrillation: past, present, and future. Cardiovascular Research, 2011, 89, 701-709.	3.8	66
45	White Blood Cell Count and Risk of Incident Atrial Fibrillation (From the Framingham Heart Study). American Journal of Cardiology, 2012, 109, 533-537.	1.6	66
46	A novel trafficking-defective HCN4 mutation is associated with early-onset atrial fibrillation. Heart Rhythm, 2014, 11, 1055-1062.	0.7	64
47	Alcohol consumption, sinus tachycardia, and cardiac arrhythmias at the Munich Octoberfest: results from the Munich Beer Related Electrocardiogram Workup Study (MunichBREW). European Heart Journal, 2017, 38, 2100-2106.	2.2	61
48	A meta-analysis of genome-wide association studies of the electrocardiographic early repolarization pattern. Heart Rhythm, 2012, 9, 1627-1634.	0.7	58
49	Genome Wide Analysis of Drug-Induced Torsades de Pointes: Lack of Common Variants with Large Effect Sizes. PLoS ONE, 2013, 8, e78511.	2.5	57
50	Incidence of sudden cardiac death in Germany: results from an emergency medical service registry in Lower Saxony. Europace, 2014, 16, 1752-1758.	1.7	54
51	Common Variants in <i>CASQ2</i> , <i>GPD1L</i> , and <i>NOS1AP</i> Are Significantly Associated With Risk of Sudden Death in Patients With Coronary Artery Disease. Circulation: Cardiovascular Genetics, 2011, 4, 397-402.	5.1	53
52	Spontaneous Brugada electrocardiogram patterns are rare in the German general population: results from the KORA study. Europace, 2009, 11, 1338-1344.	1.7	52
53	Analysis for Genetic Modifiers of Disease Severity in Patients With Long-QT Syndrome Type 2. Circulation: Cardiovascular Genetics, 2015, 8, 447-456.	5.1	51
54	Prediction of mortality benefit based on periodic repolarisation dynamics in patients undergoing prophylactic implantation of a defibrillator: a prospective, controlled, multicentre cohort study. Lancet, The, 2019, 394, 1344-1351.	13.7	49

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55	Exome-chip meta-analysis identifies novel loci associated with cardiac conduction, including ADAMTS6. Genome Biology, 2018, 19, 87.	8.8	47
56	Early repolarization pattern is the strongest predictor of arrhythmia recurrence in patients with idiopathic ventricular fibrillation: results from a single centre long-term follow-up over 20 years. Europace, 2016, 18, 718-725.	1.7	44
57	Reduced left atrial cardiomyocyte PITX2 and elevated circulating BMP10 predict atrial fibrillation after ablation. JCI Insight, 2020, 5, .	5.0	44
58	Dynamic risk assessment to improve quality of care in patients with atrial fibrillation: the 7th AFNET/EHRA Consensus Conference. Europace, 2021, 23, 329-344.	1.7	38
59	Deciphering the Plasma Proteome of Type 2 Diabetes. Diabetes, 2020, 69, 2766-2778.	0.6	34
60	The Role of MicroRNAs in Antiarrhythmic Therapy for Atrial Fibrillation. Arrhythmia and Electrophysiology Review, 2015, 4, 146.	2.4	30
61	Genetic Susceptibility for Atrial Fibrillation in Patients Undergoing Atrial Fibrillation Ablation. Circulation: Arrhythmia and Electrophysiology, 2020, 13, e007676.	4.8	30
62	Discovery of novel heart rate-associated loci using the Exome Chip. Human Molecular Genetics, 2017, 26, 2346-2363.	2.9	29
63	Genome-wide association meta-analysis of 30,000 samples identifies seven novel loci for quantitative ECG traits. European Journal of Human Genetics, 2019, 27, 952-962.	2.8	29
64	The common non-synonymous variant G38S of the KCNE1-(minK)-gene is not associated to QT interval in Central European Caucasians: results from the KORA study. European Heart Journal, 2007, 28, 305-309.	2.2	27
65	Targeted sequencing in candidate genes for atrial fibrillation: The Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) Targeted Sequencing Study. Heart Rhythm, 2014, 11, 452-457.	0.7	24
66	Chronically elevated branched chain amino acid levels are pro-arrhythmic. Cardiovascular Research, 2022, 118, 1742-1757.	3.8	24
67	Lack of replication in polymorphisms reported to be associated with atrial fibrillation. Heart Rhythm, 2011, 8, 403-409.	0.7	22
68	One-year clinical outcome after ablation with a novel multipolar irrigated ablation catheter for treatment of atrial fibrillation: potential implications for clinical use. Europace, 2016, 18, 1170-1178.	1.7	17
69	Repolarization Heterogeneity Measured With T-Wave Area Dispersion in Standard 12-Lead ECG Predicts Sudden Cardiac Death in General Population. Circulation: Arrhythmia and Electrophysiology, 2018, 11, e005762.	4.8	17
70	Manual Compression versus Vascular Closing Device for Closing Access Puncture Site in Femoral Left-Heart Catheterization and Percutaneous Coronary Interventions: A Retrospective Cross-Sectional Comparison of Costs and Effects in Inpatient Care. Value in Health, 2017, 20, 769-776.	0.3	12
71	Impact of acute ethanol intake on cardiac autonomic regulation. Scientific Reports, 2021, 11, 13255.	3.3	12
72	Geographic variation in the use of catheter ablation for atrial fibrillation among Medicare beneficiaries. American Heart Journal, 2015, 169, 775-782.e2.	2.7	11

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73	Initial single centre experience with the novel Rhythmia© high density mapping system in an all comer collective of 400 electrophysiological patients. International Journal of Cardiology, 2018, 272, 168-174.	1.7	11
74	Development and external validation of predictive models for prevalent and recurrent atrial fibrillation: a protocol for the analysis of the CATCH ME combined dataset. BMC Cardiovascular Disorders, 2019, 19, 120.	1.7	10
75	A practical guide to setting up pig models for cardiovascular catheterization, electrophysiological assessment and heart disease research. Lab Animal, 2022, 51, 46-67.	0.4	10
76	Outcomes of ablation in Wolff-Parkinson-White-syndrome: Data from the German Ablation Registry. International Journal of Cardiology, 2021, 323, 106-112.	1.7	9
77	A genetic variant alters the secondary structure of the lncRNA H19 and is associated with dilated cardiomyopathy. RNA Biology, 2021, 18, 409-415.	3.1	9
78	Early decision-analytic modeling – a case study on vascular closure devices. BMC Health Services Research, 2015, 15, 486.	2.2	8
79	Cardiac Risk Factors for Stroke: A Comprehensive Mendelian Randomization Study. Stroke, 2022, 53, STROKEAHA121036306.	2.0	8
80	Completion of Guidelineâ€Recommended Initial Evaluation of Atrial Fibrillation. Clinical Cardiology, 2012, 35, 585-593.	1.8	7
81	Catch-up-ESUS - follow-up in embolic stroke of undetermined source (ESUS) in a prospective, open-label, observational study: study protocol and initial baseline data. BMJ Open, 2019, 9, e031716.	1.9	5
82	Singleâ€center experience of ultraâ€highâ€density mapping guided catheter ablation of focal atrial tachycardia. Clinical Cardiology, 2022, , .	1.8	5
83	Effects of acute alcohol consumption on cardiac excitation, conduction, and repolarization: results from the Munich Beer Related Electrocardiogram Workup Study (MunichBREW). Clinical Research in Cardiology, 2021, 110, 916-918.	3.3	4
84	Benefit of Contact Force Sensing Catheter Technology for Successful Left Atrial Anterior Line Formation: A Prospective Randomized Trial. BioMed Research International, 2018, 2018, 1-8.	1.9	3
85	Arrhythmias at the Munich Octoberfest: ECG under the influence?. European Heart Journal, 2017, 38, 2641-2643.	2.2	2
86	Keep it simple: the ECG and sudden cardiac death risk. Heart, 2020, 106, 403-404.	2.9	2
87	Genome-Wide Association Studies Revealing the Heritability of Common Atrial Fibrillation. Circulation: Cardiovascular Genetics, 2017, 10, .	5.1	1
88	Atrioventricular block grade III in the context of acute alcohol intake. European Journal of Emergency Medicine, 2021, 28, 75-76.	1.1	1
89	Common electrocardiogram measures are not associated with telomere length. Aging, 0, , .	3.1	1
90	Recurrent Stroke in a Young Patient with Embolic Stroke of Undetermined Source and Patent Foramen Ovale: Quo Vadis?. Case Reports in Neurology, 2020, 12, 45-49.	0.7	0

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	Atrial fibrillation in Iran: Familiar findings in familial AF. International Journal of Cardiology, 2020, 314, 75-76.	1.7	0