## Marc Coram

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6643400/publications.pdf

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17 papers	6,128 citations	14 h-index	993246 17 g-index
18	18	18	11006 citing authors
all docs	docs citations	times ranked	

#	Article	IF	CITATIONS
1	Integrating deep learning and unbiased automated high-content screening to identify complex disease signatures in human fibroblasts. Nature Communications, 2022, 13, 1590.	5.8	29
2	Quantum optimization with a novel Gibbs objective function and ansatz architecture search. Physical Review Research, 2020, 2, .	1.3	48
3	Learning causal networks using inducible transcription factors and transcriptomeâ€wide time series. Molecular Systems Biology, 2020, 16, e9174.	3.2	51
4	Performance of a Deep-Learning Algorithm vs Manual Grading for Detecting Diabetic Retinopathy in India. JAMA Ophthalmology, 2019, 137, 987.	1.4	171
5	A Poisson Log-Normal Model for Constructing Gene Covariation Network Using RNA-seq Data. Journal of Computational Biology, 2017, 24, 721-731.	0.8	4
6	Development and Validation of a Deep Learning Algorithm for Detection of Diabetic Retinopathy in Retinal Fundus Photographs. JAMA - Journal of the American Medical Association, 2016, 316, 2402.	3.8	4,738
7	A minimum variance method for genome-wide data-driven normalization of quantitative real-time polymerase chain reaction expression data. Analytical Biochemistry, 2014, 458, 11-13.	1.1	5
8	Antibodies specifically target AML antigen NuSAP1 after allogeneic bone marrow transplantation. Blood, 2010, 115, 2077-2087.	0.6	29
9	Single-Cell Profiling Identifies Aberrant STAT5 Activation in Myeloid Malignancies with Specific Clinical and Biologic Correlates. Cancer Cell, 2008, 14, 335-343.	7.7	219
10	The Significance of Monoamine Oxidase-A Expression in High Grade Prostate Cancer. Journal of Urology, 2008, 180, 2206-2211.	0.2	53
11	H-Y Antibody Development Associates With Acute Rejection in Female Patients With Male Kidney Transplants. Transplantation, 2008, 86, 75-81.	0.5	84
12	A statistical method for chromatographic alignment of LC-MS data. Biostatistics, 2007, 8, 357-367.	0.9	60
13	Reconstructing Genetic Ancestry Blocks in Admixed Individuals. American Journal of Human Genetics, 2006, 79, 1-12.	2.6	240
14	Quantitative Analysis of Acrylamide Labeled Serum Proteins by LCâ^'MS/MS. Journal of Proteome Research, 2006, 5, 2009-2018.	1.8	111
15	Consistency of Bayes estimators of a binary regression function. Annals of Statistics, 2006, 34, 1233.	1.4	5
16	A suite of algorithms for the comprehensive analysis of complex protein mixtures using high-resolution LC-MS. Bioinformatics, 2006, 22, 1902-1909.	1.8	250
17	New tests of the correspondence between unitary eigenvalues and the zeros of RiemannÂs zeta function. Journal of Physics A, 2003, 36, 2883-2906.	1.6	26