

# Matthew P Harrigan

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2113299/publications.pdf>

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18  
papers

9,065  
citations

567281

15  
h-index

839539

18  
g-index

21  
all docs

21  
docs citations

21  
times ranked

9724  
citing authors

#	ARTICLE	IF	CITATIONS
1	Time-crystalline eigenstate order on a quantum processor. <i>Nature</i> , 2022, 601, 531-536.	27.8	138
2	Quantum approximate optimization of non-planar graph problems on a planar superconducting processor. <i>Nature Physics</i> , 2021, 17, 332-336.	16.7	262
3	Accurately computing the electronic properties of a quantum ring. <i>Nature</i> , 2021, 594, 508-512.	27.8	47
4	Exponential suppression of bit or phase errors with cyclic error correction. <i>Nature</i> , 2021, 595, 383-387.	27.8	172
5	Low-Depth Mechanisms for Quantum Optimization. <i>PRX Quantum</i> , 2021, 2, .	9.2	17
6	What the foundations of quantum computer science teach us about chemistry. <i>Journal of Chemical Physics</i> , 2021, 155, 150901.	3.0	9
7	Information scrambling in quantum circuits. <i>Science</i> , 2021, 374, 1479-1483.	12.6	127
8	Realizing topologically ordered states on a quantum processor. <i>Science</i> , 2021, 374, 1237-1241.	12.6	186
9	Hartree-Fock on a superconducting qubit quantum computer. <i>Science</i> , 2020, 369, 1084-1089.	12.6	453
10	Using models to improve optimizers for variational quantum algorithms. <i>Quantum Science and Technology</i> , 2020, 5, 044008.	5.8	46
11	Quantum supremacy using a programmable superconducting processor. <i>Nature</i> , 2019, 574, 505-510.	27.8	4,148
12	MSMBuilder: Statistical Models for Biomolecular Dynamics. <i>Biophysical Journal</i> , 2017, 112, 10-15.	0.5	228
13	Markov modeling reveals novel intracellular modulation of the human TREK-2 selectivity filter. <i>Scientific Reports</i> , 2017, 7, 632.	3.3	15
14	OpenMM 7: Rapid development of high performance algorithms for molecular dynamics. <i>PLoS Computational Biology</i> , 2017, 13, e1005659.	3.2	1,561
15	MSMExplorer: Data Visualizations for Biomolecular Dynamics. <i>Journal of Open Source Software</i> , 2017, 2, 188.	4.6	18
16	Osprey: Hyperparameter Optimization for Machine Learning. <i>Journal of Open Source Software</i> , 2016, 1, 34.	4.6	33
17	Conserve Water: A Method for the Analysis of Solvent in Molecular Dynamics. <i>Journal of Chemical Theory and Computation</i> , 2015, 11, 1094-1101.	5.3	14
18	MDTraj: A Modern Open Library for the Analysis of Molecular Dynamics Trajectories. <i>Biophysical Journal</i> , 2015, 109, 1528-1532.	0.5	1,576