

# Rob Ww Hooft

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

37  
papers

8,515  
citations

20  
h-index

39  
g-index

39  
ext. papers

11,111  
ext. citations

7.4  
avg, IF

5.19  
L-index

#	Paper	IF	Citations
37	Application Profile for Machine-Actionable Data Management Plans. <i>Data Science Journal</i> , <b>2021</b> , 20, 32	2	
36	FAIR Principles: Interpretations and Implementation Considerations. <i>Data Intelligence</i> , <b>2020</b> , 2, 10-29	3	66
35	Facilities that make the PDB data collection more powerful. <i>Protein Science</i> , <b>2020</b> , 29, 330-344	6.3	4
34	Data Management Planning: How Requirements and Solutions are Beginning to Converge. <i>Data Intelligence</i> , <b>2020</b> , 2, 208-219	3	12
33	Data Stewardship Wizard – A Tool Bringing Together Researchers, Data Stewards, and Data Experts around Data Management Planning. <i>Data Science Journal</i> , <b>2019</b> , 18,	2	8
32	Bioinformatics in the Netherlands: the value of a nationwide community. <i>Briefings in Bioinformatics</i> , <b>2019</b> , 20, 540-550	13.4	2
31	Four simple recommendations to encourage best practices in research software. <i>F1000Research</i> , <b>2017</b> , 6,	3.6	58
30	Top 10 metrics for life science software good practices. <i>F1000Research</i> , <b>2016</b> , 5,	3.6	12
29	The FAIR Guiding Principles for scientific data management and stewardship. <i>Scientific Data</i> , <b>2016</b> , 3, 160018	8.2	4154
28	The Dutch Techcentre for Life Sciences: Enabling data-intensive life science research in the Netherlands. <i>F1000Research</i> , <b>2015</b> , 4, 33	3.6	8
27	The Dutch Techcentre for Life Sciences: Enabling data-intensive life science research in the Netherlands. <i>F1000Research</i> , <b>2015</b> , 4, 33	3.6	7
26	A series of PDB related databases for everyday needs. <i>Nucleic Acids Research</i> , <b>2011</b> , 39, D411-9	20.1	487
25	A high-throughput processing service for retention time alignment of complex proteomics and metabolomics LC-MS data. <i>Bioinformatics</i> , <b>2011</b> , 27, 1176-8	7.2	5
24	Using the t-distribution to improve the absolute structure assignment with likelihood calculations. <i>Journal of Applied Crystallography</i> , <b>2010</b> , 43, 665-668	3.8	81
23	Probability plots based on Student's t-distribution. <i>Acta Crystallographica Section A: Foundations and Advances</i> , <b>2009</b> , 65, 319-21		23
22	Determination of absolute structure using Bayesian statistics on Bijvoet differences. <i>Journal of Applied Crystallography</i> , <b>2008</b> , 41, 96-103	3.8	579
21	Accurate cells from area-detector images. <i>Journal of Applied Crystallography</i> , <b>2000</b> , 33, 893-898	3.8	169

20	Who checks the checkers? Four validation tools applied to eight atomic resolution structures. EU 3-D Validation Network. <i>Journal of Molecular Biology</i> , <b>1998</b> , 276, 417-36	6.5	106
19	Objectively judging the quality of a protein structure from a Ramachandran plot. <i>Bioinformatics</i> , <b>1997</b> , 13, 425-30	7.2	122
18	Crystal Engineering of Melamine-Imide Complexes; Tuning the Stoichiometry by Steric Hindrance of the Imide Carbonyl Groups. <i>Angewandte Chemie International Edition in English</i> , <b>1997</b> , 36, 969-971		55
17	Verification of Protein Structures: Side-Chain Planarity. <i>Journal of Applied Crystallography</i> , <b>1996</b> , 29, 714-716	3.6	59
16	Positioning hydrogen atoms by optimizing hydrogen-bond networks in protein structures. <i>Proteins: Structure, Function and Bioinformatics</i> , <b>1996</b> , 26, 363-76	4.2	256
15	Improved coordinate reconstruction from stereo diagrams. <i>Journal of Molecular Graphics</i> , <b>1996</b> , 14, 168-72		4
14	Errors in protein structures. <i>Nature</i> , <b>1996</b> , 381, 272	50.4	1730
13	Divergent evolution of a beta/alpha-barrel subclass: detection of numerous phosphate-binding sites by motif search. <i>Protein Science</i> , <b>1995</b> , 4, 268-74	6.3	42
12	The use of position-specific rotamers in model building by homology. <i>Proteins: Structure, Function and Bioinformatics</i> , <b>1995</b> , 23, 415-21	4.2	126
11	Structural studies on sweet taste inhibitors: lactisole, dl-2(4-methoxyphenoxy)-propanoic acid. <i>Journal of Molecular Structure</i> , <b>1994</b> , 326, 25-34	3.4	7
10	Reconstruction of symmetry-related molecules from protein data bank (PDB) files. <i>Journal of Applied Crystallography</i> , <b>1994</b> , 27, 1006-1009	3.8	8
9	Molecular dynamics study of conformational and anomeric equilibria in aqueous D-glucose. <i>The Journal of Physical Chemistry</i> , <b>1993</b> , 97, 12093-12099		51
8	The structure of 1,6-diazacyclododecane-7,12-dione, a cyclic monomeric model of polyamide 46. <i>Journal of Crystallographic and Spectroscopic Research</i> , <b>1993</b> , 23, 917-920		5
7	Structural studies on sweet taste inhibitors: methyl-4,6-dichloro-4,6-dideoxygalactopyranoside. <i>Journal of Molecular Structure</i> , <b>1993</b> , 291, 173-182	3.4	6
6	Use of molecular dynamics methods in conformational analysis. Glycol. A model study. <i>Journal of Chemical Physics</i> , <b>1992</b> , 97, 3639-3646	3.9	34
5	An adaptive umbrella sampling procedure in conformational analysis using molecular dynamics and its application to glycol. <i>Journal of Chemical Physics</i> , <b>1992</b> , 97, 6690-6694	3.9	114
4	Structural studies on artificial sweeteners: itN-(4-(1-propyloxy)-phenyl)-urea. <i>Journal of Molecular Structure</i> , <b>1991</b> , 263, 267-279	3.4	2
3	Structure of racemic 4-isopropenyl-1-cyclohexene-1-carbaldehyde oxime (perillartine). <i>Acta Crystallographica Section C: Crystal Structure Communications</i> , <b>1990</b> , 46, 1133-1135		0

2	Low-temperature, high-resolution crystal structure study of tricyclo[3.3.1.1 <sup>3,7</sup> ]decan-2-ol (adamantanol-2). <i>Journal of Crystallographic and Spectroscopic Research</i> , <b>1990</b> , 20, 123-131	3
1	Software vs. data in the context of citation	10