## Dirk Roeland Boer

List of Publications by Year in descending order

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31 papers

1,690 citations

471509 17 h-index 32 g-index

34 all docs

34 docs citations

34 times ranked 2634 citing authors

#	Article	IF	CITATIONS
1	Structural Basis for DNA Binding Specificity by the Auxin-Dependent ARF Transcription Factors. Cell, 2014, 156, 577-589.	28.9	348
2	Molecular Recognition of a Three-Way DNA Junction by a Metallosupramolecular Helicate. Angewandte Chemie - International Edition, 2006, 45, 1227-1231.	13.8	278
3	DNA-binding drugs caught in action: the latest 3D pictures of drug-DNA complexes. Dalton Transactions, 2009, , 399-414.	3.3	153
4	The crystallography stations at the Alba synchrotron. European Physical Journal Plus, 2015, 130, 1.	2.6	96
5	Design principles of a minimal auxin response system. Nature Plants, 2020, 6, 473-482.	9.3	71
6	Unveiling the Molecular Mechanism of a Conjugative Relaxase: The Structure of TrwC Complexed with a 27-mer DNA Comprising the Recognition Hairpin and the Cleavage Site. Journal of Molecular Biology, 2006, 358, 857-869.	4.2	68
7	Molecular mechanism of light-driven sodium pumping. Nature Communications, 2020, 11, 2137.	12.8	67
8	Selfâ€Assembly of Functionalizable Twoâ€Component 3D DNA Arrays through the Induced Formation of DNA Threeâ€Wayâ€Junction Branch Points by Supramolecular Cylinders. Angewandte Chemie - International Edition, 2010, 49, 2336-2339.	13.8	65
9	Architecture of DNA elements mediating ARF transcription factor binding and auxin-responsive gene expression in <i>Arabidopsis</i> . Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 24557-24566.	7.1	53
10	Discovery of a new family of relaxases in Firmicutes bacteria. PLoS Genetics, 2017, 13, e1006586.	3.5	49
11	Plasmid replication initiator RepB forms a hexamer reminiscent of ring helicases and has mobile nuclease domains. EMBO Journal, 2009, 28, 1666-1678.	7.8	45
12	Thread Insertion of a Bis(dipyridophenazine) Diruthenium Complex into the DNA Double Helix by the Extrusion of AT Base Pairs and Crossâ€Linking of DNA Duplexes. Angewandte Chemie - International Edition, 2014, 53, 1949-1952.	13.8	42
13	Superstar: comparison of CSD and PDB-based interaction fields as a basis for the prediction of protein-ligand interactions 1 1Edited by R. Huber. Journal of Molecular Biology, 2001, 312, 275-287.	4.2	40
14	X-ray Crystal Structure and EPR Spectra of "Arsenite-Inhibitedâ€DesulfovibriogigasAldehyde Dehydrogenase:Â A Member of the Xanthine Oxidase Family. Journal of the American Chemical Society, 2004, 126, 8614-8615.	13.7	35
15	<i>MXCuBE2</i> : the dawn of <i>MXCuBE</i> Collaboration. Journal of Synchrotron Radiation, 2019, 26, 393-405.	2.4	30
16	Structural basis of a histidine-DNA nicking/joining mechanism for gene transfer and promiscuous spread of antibiotic resistance. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E6526-E6535.	7.1	27
17	A C2HC zinc finger is essential for the RING-E2 interaction of the ubiquitin ligase RNF125. Scientific Reports, 2016, 6, 29232.	3.3	20
18	Correlating EPR and X-ray structural analysis of arsenite-inhibited forms of aldehyde oxidoreductase. Journal of Biological Inorganic Chemistry, 2007, 12, 353-366.	2.6	15

#	Article	IF	CITATIONS
19	Multiple Layered Control of the Conjugation Process of the Bacillus subtilis Plasmid pLS20. Frontiers in Molecular Biosciences, 2021, 8, 648468.	3.5	15
20	Nicking activity of the pMV158 MobM relaxase on cognate and heterologous origins of transfer. Plasmid, 2013, 70, 120-130.	1.4	12
21	Conformational plasticity of RepB, the replication initiator protein of promiscuous streptococcal plasmid pMV158. Scientific Reports, 2016, 6, 20915.	3.3	11
22	Differential Inhibition of Human and Trypanosome Ubiquitin E1S by TAK-243 Offers Possibilities for Parasite Selective Inhibitors. Scientific Reports, 2019, 9, 16195.	3.3	9
23	Novel regulatory mechanism of establishment genes of conjugative plasmids. Nucleic Acids Research, 2018, 46, 11910-11926.	14.5	8
24	Functional Properties and Structural Requirements of the Plasmid pMV158-Encoded MobM Relaxase Domain. Journal of Bacteriology, 2013, 195, 3000-3008.	2.2	7
25	Relation between the molecular electrostatic potential and activity of some FF-MAS related sterol compounds. Bioorganic and Medicinal Chemistry, 2001, 9, 2653-2659.	3.0	6
26	On the purification and preliminary crystallographic analysis of isoquinoline 1-oxidoreductase fromBrevundimonas diminuta7. Acta Crystallographica Section F: Structural Biology Communications, 2005, 61, 137-140.	0.7	6
27	Calculated heats of formation of sterol diene isomers compared with synthetic yields of isomerisation reactions of $\hat{l}$ ,7 sterols. Perkin Transactions II RSC, 2000, , 1701-1704.	1.1	5
28	Cobalt-, zinc- and iron-bound forms of adenylate kinase (AK) from the sulfate-reducing bacteriumDesulfovibrio gigas: purification, crystallization and preliminary X-ray diffraction analysis. Acta Crystallographica Section F: Structural Biology Communications, 2009, 65, 926-929.	0.7	4
29	Inactivation of the dimeric RappLS20 anti-repressor of the conjugation operon is mediated by peptide-induced tetramerization. Nucleic Acids Research, 2020, 48, 8113-8127.	14.5	4
30	Molecular Recognition of a Three-Way DNA Junction by a Metallosupramolecular Helicate. Angewandte Chemie - International Edition, 2006, 45, 1834-1834.	13.8	2
31	Structural and biochemical characterization of the relaxosome auxiliary proteins encoded on the Bacillus subtilis plasmid pLS20. Computational and Structural Biotechnology Journal, 2022, 20, 757-765	4.1	O