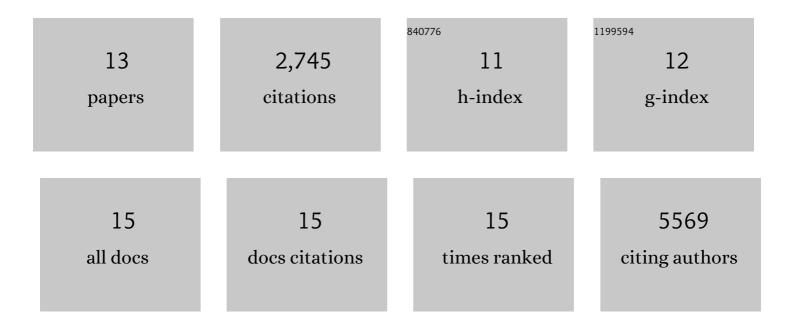
Summer B Thyme

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8901191/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Macromolecular modeling and design in Rosetta: recent methods and frameworks. Nature Methods, 2020, 17, 665-680.	19.0	513
2	Phenotypic Landscape of Schizophrenia-Associated Genes Defines Candidates and Their Shared Functions. Cell, 2019, 177, 478-491.e20.	28.9	159
3	Kctd13 deletion reduces synaptic transmission via increased RhoA. Nature, 2017, 551, 227-231.	27.8	125
4	Polq-Mediated End Joining Is Essential for Surviving DNA Double-Strand Breaks during Early Zebrafish Development. Cell Reports, 2016, 15, 707-714.	6.4	56
5	Computational Design of DNA-Binding Proteins. Methods in Molecular Biology, 2016, 1414, 265-283.	0.9	0
6	Internal guide RNA interactions interfere with Cas9-mediated cleavage. Nature Communications, 2016, 7, 11750.	12.8	133
7	CHOPCHOP v2: a web tool for the next generation of CRISPR genome engineering. Nucleic Acids Research, 2016, 44, W272-W276.	14.5	801
8	Reprogramming homing endonuclease specificity through computational design and directed evolution. Nucleic Acids Research, 2014, 42, 2564-2576.	14.5	31
9	Progressive engineering of a homing endonuclease genome editing reagent for the murine X-linked immunodeficiency locus. Nucleic Acids Research, 2014, 42, 6463-6475.	14.5	8
10	Efficient Mutagenesis by Cas9 Protein-Mediated Oligonucleotide Insertion and Large-Scale Assessment of Single-Guide RNAs. PLoS ONE, 2014, 9, e98186.	2.5	794
11	Improved Modeling of Side-Chain–Base Interactions and Plasticity in Protein–DNA Interface Design. Journal of Molecular Biology, 2012, 419, 255-274.	4.2	20
12	Mining Endonuclease Cleavage Determinants in Genomic Sequence Data. Journal of Biological Chemistry, 2011, 286, 32617-32627.	3.4	15
13	Exploitation of binding energy for catalysis and design. Nature, 2009, 461, 1300-1304.	27.8	86