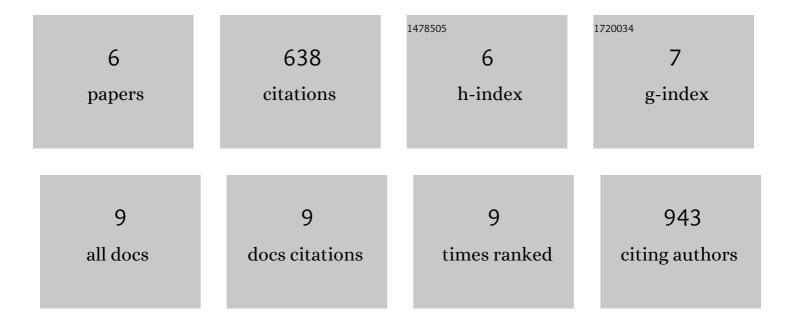
## Mitchell G Kluesner

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/396729/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	EditR: A Method to Quantify Base Editing from Sanger Sequencing. CRISPR Journal, 2018, 1, 239-250.	2.9	304
2	Highly efficient multiplex human T cell engineering without double-strand breaks using Cas9 base editors. Nature Communications, 2019, 10, 5222.	12.8	135
3	A Genetically Engineered Primary Human Natural Killer Cell Platform for Cancer Immunotherapy. Molecular Therapy, 2020, 28, 52-63.	8.2	120
4	CRISPR-Cas9 cytidine and adenosine base editing of splice-sites mediates highly-efficient disruption of proteins in primary and immortalized cells. Nature Communications, 2021, 12, 2437.	12.8	50
5	MultiEditR: The first tool for the detection and quantification of RNA editing from Sanger sequencing demonstrates comparable fidelity to RNA-seq. Molecular Therapy - Nucleic Acids, 2021, 25, 515-523.	5.1	11
6	C-to-U RNA Editing: From Computational Detection to Experimental Validation. Methods in Molecular Biology, 2021, 2181, 51-67.	0.9	4