

# Deepak Reyon

## List of Publications by Year in descending order

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Version: 2024-02-01

21  
papers

5,403  
citations

535685

17  
h-index

799663

21  
g-index

23  
all docs

23  
docs citations

23  
times ranked

9651  
citing authors

#	ARTICLE	IF	CITATIONS
1	Zebrafish <i>dscaml1</i> Deficiency Impairs Retinal Patterning and Oculomotor Function. <i>Journal of Neuroscience</i> , 2020, 40, 143-158.	1.7	15
2	Disruption of the kringle 1 domain of prothrombin leads to late onset mortality in zebrafish. <i>Scientific Reports</i> , 2020, 10, 4049.	1.6	10
3	Disruption of <i>asxl1</i> results in myeloproliferative neoplasms in zebrafish. <i>DMM Disease Models and Mechanisms</i> , 2019, 12, .	1.2	18
4	Efficient CRISPR/Cas9-mediated editing of trinucleotide repeat expansion in myotonic dystrophy patient-derived iPS and myogenic cells. <i>Nucleic Acids Research</i> , 2018, 46, 8275-8298.	6.5	78
5	Genome editing of factor X in zebrafish reveals unexpected tolerance of severe defects in the common pathway. <i>Blood</i> , 2017, 130, 666-676.	0.6	22
6	Uromodulin p.Cys147Trp mutation drives kidney disease by activating ER stress and apoptosis. <i>Journal of Clinical Investigation</i> , 2017, 127, 3954-3969.	3.9	49
7	Nodal patterning without Lefty inhibitory feedback is functional but fragile. <i>ELife</i> , 2017, 6, .	2.8	52
8	Context influences on TALE DNA binding revealed by quantitative profiling. <i>Nature Communications</i> , 2015, 6, 7440.	5.8	30
9	Targeted disruption of DNMT1, DNMT3A and DNMT3B in human embryonic stem cells. <i>Nature Genetics</i> , 2015, 47, 469-478.	9.4	409
10	Factor X Mutant Zebrafish Tolerate a Severe Hemostatic Defect in Early Development Yet Develop Lethal Hemorrhage in Adulthood. <i>Blood</i> , 2015, 126, 426-426.	0.6	1
11	Targeted Genome Editing in Human Cells Using CRISPR/Cas Nucleases and Truncated Guide RNAs. <i>Methods in Enzymology</i> , 2014, 546, 21-45.	0.4	43
12	Systematic screening reveals a role for BRCA1 in the response to transcription-associated DNA damage. <i>Genes and Development</i> , 2014, 28, 1957-1975.	2.7	86
13	Broad specificity profiling of TALENs results in engineered nucleases with improved DNA-cleavage specificity. <i>Nature Methods</i> , 2014, 11, 429-435.	9.0	182
14	Dimeric CRISPR RNA-guided FokI nucleases for highly specific genome editing. <i>Nature Biotechnology</i> , 2014, 32, 569-576.	9.4	852
15	Improving CRISPR-Cas nuclease specificity using truncated guide RNAs. <i>Nature Biotechnology</i> , 2014, 32, 279-284.	9.4	1,706
16	I $\kappa$ B Kinase $\hat{2}$ (IKK $\hat{2}$ ) Mutations in Lymphomas That Constitutively Activate Canonical Nuclear Factor $\hat{B}$ (NF $\hat{B}$ ) Signaling. <i>Journal of Biological Chemistry</i> , 2014, 289, 26960-26972.	1.6	20
17	Interactome Maps of Mouse Gene Regulatory Domains Reveal Basic Principles of Transcriptional Regulation. <i>Cell</i> , 2013, 155, 1507-1520.	13.5	299
18	Engineering Customized TALE Nucleases (TALENs) and TALE Transcription Factors by Fast Ligation-Based Automatable Solid-Phase High-Throughput (FLASH) Assembly. <i>Current Protocols in Molecular Biology</i> , 2013, 103, Unit 12.16.	2.9	28

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19	Engineering Designer Transcription Activatorâ€œLike Effector Nucleases (TALENs) by REAL or REALâ€œFast Assembly. <i>Current Protocols in Molecular Biology</i> , 2012, 100, Unit 12.15.	2.9	68
20	FLASH assembly of TALENs for high-throughput genome editing. <i>Nature Biotechnology</i> , 2012, 30, 460-465.	9.4	1,070
21	ZiFit (Zinc Finger Targeter): an updated zinc finger engineering tool. <i>Nucleic Acids Research</i> , 2010, 38, W462-W468.	6.5	365