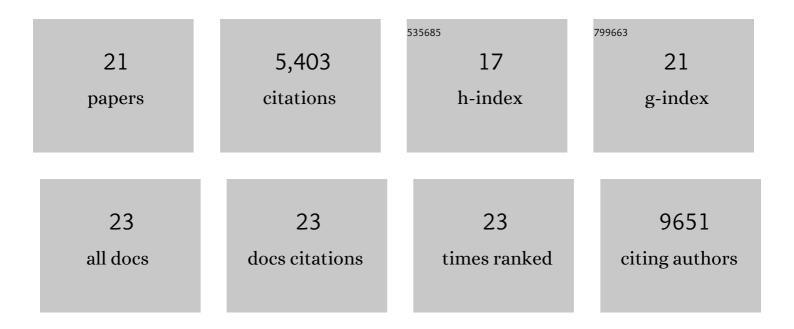
Deepak Reyon

List of Publications by Year in descending order

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DEEDAK REVON

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

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