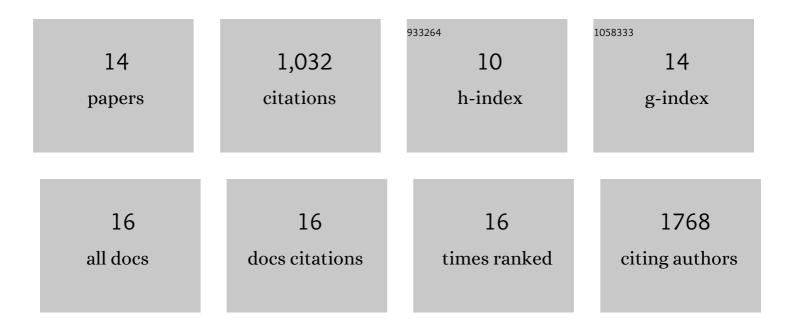
## David Sturgill

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6177186/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Oncogenic IncRNAs alter epigenetic memory at a fragile chromosomal site in human cancer cells. Science Advances, 2022, 8, eabl5621.	4.7	18
2	The X Chromosome from Telomere to Telomere: Key Achievements and Future Opportunities. Faculty Reviews, 2021, 10, 63.	1.7	1
3	Activation of Crtc2/Creb1 in skeletal muscle enhances weight loss during intermittent fasting. FASEB Journal, 2021, 35, e21999.	0.2	3
4	Membraneless nuclear organelles and the search for phases within phases. Wiley Interdisciplinary Reviews RNA, 2019, 10, e1514.	3.2	111
5	Immunoprecipitation and Sequencing of Acetylated RNA. Bio-protocol, 2019, 9, e3278.	0.2	11
6	Replication Stress Shapes a Protective Chromatin Environment across Fragile Genomic Regions. Molecular Cell, 2018, 69, 36-47.e7.	4.5	75
7	Acetylation of Cytidine in mRNA Promotes Translation Efficiency. Cell, 2018, 175, 1872-1886.e24.	13.5	409
8	HJURP antagonizes CENP-A mislocalization driven by the H3.3 chaperones HIRA and DAXX. PLoS ONE, 2018, 13, e0205948.	1.1	37
9	Independence between pre-mRNA splicing and DNA methylation in an isogenic minigene resource. Nucleic Acids Research, 2017, 45, 12780-12797.	6.5	4
10	<scp>TET</scp> â€catalyzed oxidation of intragenic 5â€methylcytosine regulates <scp>CTCF</scp> â€dependent alternative splicing. EMBO Journal, 2016, 35, 335-355.	3.5	111
11	Cajal body function in genome organization and transcriptome diversity. BioEssays, 2016, 38, 1197-1208.	1.2	56
12	Cajal bodies are linked to genome conformation. Nature Communications, 2016, 7, 10966.	5.8	127
13	Controlled DNA double-strand break induction in mice reveals post-damage transcriptome stability. Nucleic Acids Research, 2016, 44, e64-e64.	6.5	44
14	<i>Sxl</i> -Dependent, <i>tra/tra2</i> -Independent Alternative Splicing of the <i>Drosophila melanogaster</i> X-Linked Gene <i>found in neurons</i> . G3: Genes, Genomes, Genetics, 2015, 5, 2865-2874.	0.8	17