

# Dongxu Li

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/993453/publications.pdf>

Version: 2024-02-01

12

papers

461

citations

10

h-index

12

g-index

12

all docs

12

docs citations

12

times ranked

541

citing authors

#	ARTICLE	IF	CITATIONS
1	Structure and regulation of ZCCHC4 in m6A-methylation of 28S rRNA. <i>Nature Communications</i> , 2019, 10, 5042.	12.8	72
2	A selective WDR5 degrader inhibits acute myeloid leukemia in patient-derived mouse models. <i>Science Translational Medicine</i> , 2021, 13, eabj1578.	12.4	67
3	Harnessing the E3 Ligase KEAP1 for Targeted Protein Degradation. <i>Journal of the American Chemical Society</i> , 2021, 143, 15073-15083.	13.7	66
4	ZFX Mediates Non-canonical Oncogenic Functions of the Androgen Receptor Splice Variant 7 in Castrate-Resistant Prostate Cancer. <i>Molecular Cell</i> , 2018, 72, 341-354.e6.	9.7	64
5	BAHCC1 binds H3K27me3 via a conserved BAH module to mediate gene silencing and oncogenesis. <i>Nature Genetics</i> , 2020, 52, 1384-1396.	21.4	57
6	Discovery and Characterization of a Cellular Potent Positive Allosteric Modulator of the Polycomb Repressive Complex 1 Chromodomain, CBX7. <i>Cell Chemical Biology</i> , 2019, 26, 1365-1379.e22.	5.2	38
7	A NSD3-targeted PROTAC suppresses NSD3 and cMyc oncogenic nodes in cancer cells. <i>Cell Chemical Biology</i> , 2022, 29, 386-397.e9.	5.2	30
8	Discovery of a dual WDR5 and Ikaros PROTAC degrader as an anti-cancer therapeutic. <i>Oncogene</i> , 2022, 41, 3328-3340.	5.9	18
9	Novel RNA-Affinity Proteogenomics Dissects Tumor Heterogeneity for Revealing Personalized Markers in Precision Prognosis of Cancer. <i>Cell Chemical Biology</i> , 2018, 25, 619-633.e5.	5.2	15
10	Reprogramming CBX8-PRC1 function with a positive allosteric modulator. <i>Cell Chemical Biology</i> , 2022, 29, 555-571.e11.	5.2	12
11	Epstein-Barr virus genomes in Hodgkin's disease and non-Hodgkin's lymphomas. <i>Pathology International</i> , 1995, 45, 735-741.	1.3	11
12	DOT1L activity in leukemia cells requires interaction with ubiquitylated H2B that promotes productive nucleosome binding. <i>Cell Reports</i> , 2022, 38, 110369.	6.4	11