

# Tiziana Lischetti

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8358321/publications.pdf>

Version: 2024-02-01

11  
papers

664  
citations

1040056

9  
h-index

1372567

10  
g-index

11  
all docs

11  
docs citations

11  
times ranked

797  
citing authors

#	ARTICLE	IF	CITATIONS
1	Direct binding between BubR1 and B56â€“PP2A phosphatase complexes regulate mitotic progression. <i>Journal of Cell Science</i> , 2013, 126, 1086-1092.	2.0	187
2	Distinct domains in Bub1 localize RZZ and BubR1 to kinetochores to regulate the checkpoint. <i>Nature Communications</i> , 2015, 6, 7162.	12.8	99
3	A minimal number of MELT repeats supports all functions of KNL1 in chromosome segregation. <i>Journal of Cell Science</i> , 2014, 127, 871-84.	2.0	94
4	Structure of a Blinkin-BUBR1 Complex Reveals an Interaction Crucial for Kinetochores-Mitotic Checkpoint Regulation via an Unanticipated Binding Site. <i>Structure</i> , 2011, 19, 1691-1700.	3.3	68
5	The internal Cdc20 binding site in BubR1 facilitates both spindle assembly checkpoint signalling and silencing. <i>Nature Communications</i> , 2014, 5, 5563.	12.8	55
6	Epigenetic inactivation of suppressors of cytokine signalling in Philadelphia-negative chronic myeloproliferative disorders. <i>British Journal of Haematology</i> , 2008, 141, 504-511.	2.5	54
7	Regulation of mitotic progression by the spindle assembly checkpoint. <i>Molecular and Cellular Oncology</i> , 2015, 2, e970484.	0.7	45
8	The coding and long noncoding single-cell atlas of the developing human fetal striatum. <i>Science</i> , 2021, 372, .	12.6	40
9	The evolutionary history of the polyQ tract in huntingtin sheds light on its functional pro-neural activities. <i>Cell Death and Differentiation</i> , 2022, 29, 293-305.	11.2	12
10	Conformation-specific anti-Mad2 monoclonal antibodies for the dissection of checkpoint signaling. <i>MAbs</i> , 2016, 8, 689-697.	5.2	10
11	Structure of a Blinkin-BUBR1 Complex Reveals an Interaction Crucial for Kinetochores-Mitotic Checkpoint Regulation via an Unanticipated Binding Site. <i>Structure</i> , 2011, 19, 1895.	3.3	0