

# Yang Li

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/412310/publications.pdf>

Version: 2024-02-01

16  
papers

1,060  
citations

687363

13  
h-index

996975

15  
g-index

18  
all docs

18  
docs citations

18  
times ranked

1795  
citing authors

#	ARTICLE	IF	CITATIONS
1	An oligonucleotide microarray for microRNA expression analysis based on labeling RNA with quantum dot and nanogold probe. <i>Nucleic Acids Research</i> , 2005, 33, e17-e17.	14.5	297
2	Study familial hypertrophic cardiomyopathy using patient-specific induced pluripotent stem cells. <i>Cardiovascular Research</i> , 2014, 104, 258-269.	3.8	167
3	Modeling and study of the mechanism of dilated cardiomyopathy using induced pluripotent stem cells derived from individuals with Duchenne muscular dystrophy. <i>DMM Disease Models and Mechanisms</i> , 2015, 8, 457-466.	2.4	111
4	Domestication of Transposable Elements into MicroRNA Genes in Plants. <i>PLoS ONE</i> , 2011, 6, e19212.	2.5	96
5	Overexpression of microRNA-1 promotes cardiomyocyte commitment from human cardiovascular progenitors via suppressing WNT and FGF signaling pathways. <i>Journal of Molecular and Cellular Cardiology</i> , 2013, 63, 146-154.	1.9	62
6	Evolution of MIR159/319 microRNA genes and their post-transcriptional regulatory link to siRNA pathways. <i>BMC Evolutionary Biology</i> , 2011, 11, 122.	3.2	61
7	Computational Identification of Novel Family Members of MicroRNA Genes in <i>Arabidopsis thaliana</i> and <i>Oryza sativa</i> . <i>Acta Biochimica Et Biophysica Sinica</i> , 2005, 37, 75-87.	2.0	58
8	The N-cadherin interactome in primary cardiomyocytes as defined by quantitative proximity proteomics. <i>Journal of Cell Science</i> , 2019, 132, .	2.0	53
9	HBL1 Is a Human Long Noncoding RNA that Modulates Cardiomyocyte Development from Pluripotent Stem Cells by Counteracting MIR1. <i>Developmental Cell</i> , 2017, 42, 333-348.e5.	7.0	48
10	Evolutionary rate covariation analysis of E-cadherin identifies Raskol as a regulator of cell adhesion and actin dynamics in <i>Drosophila</i> . <i>PLoS Genetics</i> , 2019, 15, e1007720.	3.5	30
11	Comparative Transcriptomic Analysis of Multiple Cardiovascular Fates from Embryonic Stem Cells Predicts Novel Regulators in Human Cardiogenesis. <i>Scientific Reports</i> , 2015, 5, 9758.	3.3	25
12	Computational identification of novel family members of microRNA genes in <i>Arabidopsis thaliana</i> and <i>Oryza sativa</i> . <i>Acta Biochimica Et Biophysica Sinica</i> , 2005, 37, 75-87.	2.0	22
13	Computational Identification of Novel Family Members of MicroRNA Genes in <i>Arabidopsis thaliana</i> and <i>Oryza sativa</i> . <i>Acta Biochimica Et Biophysica Sinica</i> , 2005, 37, 75-87.	2.0	14
14	Cloning of Novel Repeat-associated Small RNAs Derived from Hairpin Precursors in <i>Oryza sativa</i> . <i>Acta Biochimica Et Biophysica Sinica</i> , 2007, 39, 829-834.	2.0	13
15	Modeling and study of the mechanism of dilated cardiomyopathy using induced pluripotent stem cells derived from individuals with Duchenne muscular dystrophy. <i>Development (Cambridge)</i> , 2015, 142, e0905-e0905.	2.5	3
16	Reconciling computer models and stem cell models of human cardiac repolarization: reply. <i>Cardiovascular Research</i> , 2015, 106, 6-7.	3.8	0