Xia Li

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5300130/publications.pdf

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31	1,707	16	30
papers	citations	h-index	g-index
34	34	34	3237
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Editorial: Chemo-Resistance in Gastrointestinal Cancers. Frontiers in Oncology, 2022, 12, 821212.	1.3	1
2	Reconstruction of functional uterine tissues through recellularizing the decellularized rat uterine scaffolds by MSCs in vivo and in vitro. Biomedical Materials (Bristol), 2021, 16, 035023.	1.7	17
3	Nipped-B-like Protein Sensitizes Esophageal Squamous Cell Carcinoma Cells to Cisplatin via Upregulation of PUMA. Technology in Cancer Research and Treatment, 2020, 19, 153303382096072.	0.8	O
4	Arsenic trioxide (ATO) induced degradation of Cyclin D1 sensitized PD-1/PD-L1 checkpoint inhibitor in oral and esophageal squamous cell carcinoma. Journal of Cancer, 2020, 11, 6516-6529.	1.2	9
5	Characterisation of naturally occurring isothiocyanates as glutathione reductase inhibitors. Journal of Enzyme Inhibition and Medicinal Chemistry, 2020, 35, 1773-1780.	2.5	9
6	Ultrastable Anode Interface Achieved by Fluorinating Electrolytes for All-Solid-State Li Metal Batteries. ACS Energy Letters, 2020, 5, 1035-1043.	8.8	176
7	Borneol promotes apoptosis of Human Glioma Cells through regulating HIF-1a expression via mTORC1/eIF4E pathway. Journal of Cancer, 2020, 11, 4810-4822.	1.2	9
8	Arsenic nano complex induced degradation of YAP sensitized ESCC cancer cells to radiation and chemotherapy. Cell and Bioscience, 2020, 10, 146.	2.1	12
9	Risk stratification of cutaneous melanoma reveals carcinogen metabolism enrichment and immune inhibition in high-risk patients. Aging, 2020, 12, 16457-16475.	1.4	2
10	Better prognostic determination and feature characterization of cutaneous melanoma through integrative genomic analysis. Aging, 2019, 11, 5081-5107.	1.4	12
11	Methylation-Based Classification of Cervical Squamous Cell Carcinoma into Two New Subclasses Differing in Immune-Related Gene Expression. International Journal of Molecular Sciences, 2018, 19, 3607.	1.8	4
12	Histone variants H2A.Z and H3.3 coordinately regulate PRC2-dependent H3K27me3 deposition and gene expression regulation in mES cells. BMC Biology, 2018, 16, 107.	1.7	54
13	Wholeâ€exome sequencing predicted cancer epitope trees of 23 early cervical cancers in Chinese women. Cancer Medicine, 2017, 6, 207-219.	1.3	12
14	Emerging role of mutations in epigenetic regulators including MLL2 derived from The Cancer Genome Atlas for cervical cancer. BMC Cancer, 2017, 17, 252.	1.1	13
15	Dynamic changes of driver genes' mutations across clinical stages in nine cancer types. Cancer Medicine, 2016, 5, 1556-1565.	1.3	12
16	Soybean miR172c Targets the Repressive AP2 Transcription Factor NNC1 to Activate <i>ENOD40</i> Expression and Regulate Nodule Initiation Â. Plant Cell, 2015, 26, 4782-4801.	3.1	188
17	MicroRNA167-Directed Regulation of the Auxin Response Factors <i>GmARF8a</i> and <i>GmARF8b</i> Is Required for Soybean Nodulation and Lateral Root Development. Plant Physiology, 2015, 168, 984-999.	2.3	183
18	ABA signalling is fine-tuned by antagonistic HAB1 variants. Nature Communications, 2015, 6, 8138.	5.8	95

#	Article	IF	CITATIONS
19	GmmiR156b overexpression delays flowering time in soybean. Plant Molecular Biology, 2015, 89, 353-363.	2.0	49
20	Identification of Cold-Responsive miRNAs and Their Target Genes in Nitrogen-Fixing Nodules of Soybean. International Journal of Molecular Sciences, 2014, 15, 13596-13614.	1.8	54
21	PEG-mediated osmotic stress induces premature differentiation of the root apical meristem and outgrowth of lateral roots in wheat. Journal of Experimental Botany, 2014, 65, 4863-4872.	2.4	79
22	Identification of functional cooperative mutations of SETD2 in human acute leukemia. Nature Genetics, 2014, 46, 287-293.	9.4	213
23	H3.3 actively marks enhancers and primes gene transcription via opening higher-ordered chromatin. Genes and Development, 2013, 27, 2109-2124.	2.7	185
24	Differential principal component analysis of ChIP-seq. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 6789-6794.	3.3	48
25	iASeq: integrating multiple chip-seq datasets for detecting allele-specific binding. BMC Bioinformatics, 2012, 13, .	1.2	1
26	iASeq: integrative analysis of allele-specificity of protein-DNA interactions in multiple ChIP-seq datasets. BMC Genomics, 2012, 13, 681.	1.2	22
27	SOS3 mediates lateral root development under low salt stress through regulation of auxin redistribution and maxima in Arabidopsis. New Phytologist, 2011, 189, 1122-1134.	3.5	113
28	Highly efficient in vitro adventitious shoot regeneration of peppermint (Mentha x piperita L.) using internodal explants. In Vitro Cellular and Developmental Biology - Plant, 2009, 45, 435-440.	0.9	10
29	Genetic analysis of involvement of ETR1 in plant response to salt and osmotic stress. Plant Growth Regulation, 2008, 54, 261-269.	1.8	41
30	Salt-avoidance tropism in Arabidopsis thaliana. Plant Signaling and Behavior, 2008, 3, 351-353.	1.2	30
31	GA signaling and CO/FT regulatory module mediate salt-induced late flowering in Arabidopsis thaliana. Plant Growth Regulation, 2007, 53, 195-206.	1.8	54