

Battles and hijacks: noncoding transcription in plants

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Citation Report

#	ARTICLE	IF	CITATIONS
1	RNA-Seq Analysis of Rice Roots Reveals the Involvement of Post-Transcriptional Regulation in Response to Cadmium Stress. <i>Frontiers in Plant Science</i> , 2015, 6, 1136.	1.7	90
2	lncRNAs in Stress Response. <i>Current Topics in Microbiology and Immunology</i> , 2015, 394, 203-236.	0.7	24
3	Long Non-coding RNAs and Their Biological Roles in Plants. <i>Genomics, Proteomics and Bioinformatics</i> , 2015, 13, 137-147.	3.0	231
4	miRNA-encoded peptides (miPEPs): A new tool to analyze the roles of miRNAs in plant biology. <i>RNA Biology</i> , 2015, 12, 1178-1180.	1.5	48
5	Long non-coding RNAs are major contributors to transcriptome changes in sunflower meiocytes with different recombination rates. <i>BMC Genomics</i> , 2016, 17, 490.	1.2	28
6	Understanding the Functions of Long Non-Coding RNAs through Their Higher-Order Structures. <i>International Journal of Molecular Sciences</i> , 2016, 17, 702.	1.8	78
7	Sm-Like Protein-Mediated RNA Metabolism Is Required for Heat Stress Tolerance in Arabidopsis. <i>Frontiers in Plant Science</i> , 2016, 7, 1079.	1.7	26
8	Plant micro<scp>RNA</scp>s: key regulators of root architecture and biotic interactions. <i>New Phytologist</i> , 2016, 212, 22-35.	3.5	53
9	Long Non-coding RNAs in Human Disease. <i>Current Topics in Microbiology and Immunology</i> , 2016, , .	0.7	4
10	Put your 3D glasses on: plant chromatin is on show. <i>Journal of Experimental Botany</i> , 2016, 67, 3205-3221.	2.4	59
11	Noncoding RNAs, Emerging Regulators in Root Endosymbioses. <i>Molecular Plant-Microbe Interactions</i> , 2016, 29, 170-180.	1.4	44
12	The role of microRNA in abiotic stress response in plants. <i>Molecular Biology</i> , 2016, 50, 337-343.	0.4	26
13	Genome-wide identification and functional prediction of novel and fungi-responsive lincRNAs in <i>Triticum aestivum</i> . <i>BMC Genomics</i> , 2016, 17, 238.	1.2	74
14	Viroids, infectious long non-coding RNAs with autonomous replication. <i>Virus Research</i> , 2016, 212, 12-24.	1.1	42
15	Functions of plants long non-coding RNAs. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2016, 1859, 155-162.	0.9	72
16	Transposable elements (<scp>TE</scp>s) contribute to stress-related long intergenic noncoding <scp>RNA</scp>s in plants. <i>Plant Journal</i> , 2017, 90, 133-146.	2.8	116
17	Plant Epigenetics: Non-coding RNAs as Emerging Regulators. <i>RNA Technologies</i> , 2017, , 129-147.	0.2	0
18	Global analysis of ribosome-associated noncoding RNAs unveils new modes of translational regulation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E10018-E10027.	3.3	168

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19	Antisense movement on the clock. <i>New Phytologist</i> , 2017, 216, 626-628.	3.5	4
20	A Nucleus-Localized Long Non-Coding RNA Enhances Drought and Salt Stress Tolerance. <i>Plant Physiology</i> , 2017, 175, 1321-1336.	2.3	251
21	Long noncoding RNAs in the model species <i>Brachypodium distachyon</i> . <i>Scientific Reports</i> , 2017, 7, 11252.	1.6	51
22	The antiphase regulatory module comprising <i>CDF5</i> and its antisense <i>scp</i> RNA links the circadian clock to photoperiodic flowering. <i>New Phytologist</i> , 2017, 216, 854-867.	3.5	112
23	History, Discovery, and Classification of lncRNAs. <i>Advances in Experimental Medicine and Biology</i> , 2017, 1008, 1-46.	0.8	659
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25	Diversity, expansion, and evolutionary novelty of plant DNA-binding transcription factor families. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2017, 1860, 3-20.	0.9	75
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27	Survey of High Throughput RNA-Seq Data Reveals Potential Roles for lncRNAs during Development and Stress Response in Bread Wheat. <i>Frontiers in Plant Science</i> , 2017, 8, 1019.	1.7	111
28	Transcription factors network in root endosymbiosis establishment and development. <i>World Journal of Microbiology and Biotechnology</i> , 2018, 34, 37.	1.7	34
29	Splicing regulation by long noncoding RNAs. <i>Nucleic Acids Research</i> , 2018, 46, 2169-2184.	6.5	226
30	<i>Arabidopsis</i> HEAT SHOCK TRANSCRIPTION FACTOR1b regulates multiple developmental genes under benign and stress conditions. <i>Journal of Experimental Botany</i> , 2018, 69, 2847-2862.	2.4	56
31	Emerging roles of long non-coding RNAs in plant response to biotic and abiotic stresses. <i>Critical Reviews in Biotechnology</i> , 2018, 38, 93-105.	5.1	100
32	Transcriptional and Posttranscriptional Regulation of Drought Stress Treatments in <i>Brachypodium</i> Leaves. <i>Methods in Molecular Biology</i> , 2018, 1667, 21-29.	0.4	2
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39	Ascophyllum nodosum extract mitigates salinity stress in Arabidopsis thaliana by modulating the expression of miRNA involved in stress tolerance and nutrient acquisition. <i>PLoS ONE</i> , 2018, 13, e0206221.	1.1	54
40	Functions of long non-coding RNAs in plants: a riddle to explore. <i>Nucleus (India)</i> , 2018, 61, 261-272.	0.9	5
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123	Transcriptional Association between mRNAs and Their Paired Natural Antisense Transcripts Following <i>Fusarium oxysporum</i> Inoculation in <i>Brassica rapa</i> L. <i>Horticulturae</i> , 2022, 8, 17.	1.2	8
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131	Regulation mechanism of long non-coding RNA in plant secondary metabolite biosynthesis. <i>Notulae Botanicae Horti Agrobotanici Cluj-Napoca</i> , 2022, 50, 12604.	0.5	1
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142	Whole-transcriptome profiles of <i>Chrysanthemum seticuspe</i> improve genome annotation and shed new light on mRNA-miRNA-lncRNA networks in ray florets and disc florets. <i>BMC Plant Biology</i> , 2022, 22, .	1.6	3
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144	Single-Molecule Real-Time Sequencing of Full-Length Transcriptome and Identification of Genes Related to Male Development in <i>Cannabis sativa</i> . <i>Plants</i> , 2022, 11, 3559.	1.6	0
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169	Non-Coding RNA and Its Prospective Utilization in Plant Breeding. , 0, , .		0

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