

# Moussa Benhamed

## List of Publications by Year in descending order

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Version: 2024-02-01

62  
papers

7,737  
citations

117625

34  
h-index

128289

60  
g-index

70  
all docs

70  
docs citations

70  
times ranked

9274  
citing authors

#	ARTICLE	IF	CITATIONS
1	Shifting the limits in wheat research and breeding using a fully annotated reference genome. <i>Science</i> , 2018, 361, .	12.6	2,424
2	The transcriptional landscape of polyploid wheat. <i>Science</i> , 2018, 361, .	12.6	768
3	The <i>Rosa</i> genome provides new insights into the domestication of modern roses. <i>Nature Genetics</i> , 2018, 50, 772-777.	21.4	344
4	Noncoding Transcription by Alternative RNA Polymerases Dynamically Regulates an Auxin-Driven Chromatin Loop. <i>Molecular Cell</i> , 2014, 55, 383-396.	9.7	330
5	<i>Arabidopsis</i> GCN5, HD1, and TAF1/HAF2 Interact to Regulate Histone Acetylation Required for Light-Responsive Gene Expression. <i>Plant Cell</i> , 2006, 18, 2893-2903.	6.6	302
6	Splicing regulation by long noncoding RNAs. <i>Nucleic Acids Research</i> , 2018, 46, 2169-2184.	14.5	226
7	ANGUSTIFOLIA3 Binds to SWI/SNF Chromatin Remodeling Complexes to Regulate Transcription during <i>Arabidopsis</i> Leaf Development. <i>Plant Cell</i> , 2014, 26, 210-229.	6.6	219
8	Plant Immunity: From Signaling to Epigenetic Control of Defense. <i>Trends in Plant Science</i> , 2018, 23, 833-844.	8.8	198
9	Whole-genome landscape of <i>Medicago truncatula</i> symbiotic genes. <i>Nature Plants</i> , 2018, 4, 1017-1025.	9.3	192
10	Battles and hijacks: noncoding transcription in plants. <i>Trends in Plant Science</i> , 2015, 20, 362-371.	8.8	176
11	R-Loop Mediated trans Action of the APOLO Long Noncoding RNA. <i>Molecular Cell</i> , 2020, 77, 1055-1065.e4.	9.7	164
12	To die or not to die? Lessons from lesion mimic mutants. <i>Frontiers in Plant Science</i> , 2015, 6, 24.	3.6	157
13	The Plant DNA Damage Response: Signaling Pathways Leading to Growth Inhibition and Putative Role in Response to Stress Conditions. <i>Frontiers in Plant Science</i> , 2019, 10, 653.	3.6	137
14	Genome-scale <i>Arabidopsis</i> promoter array identifies targets of the histone acetyltransferase GCN5. <i>Plant Journal</i> , 2008, 56, 493-504.	5.7	120
15	Thermopriming triggers splicing memory in <i>Arabidopsis</i> . <i>Journal of Experimental Botany</i> , 2018, 69, 2659-2675.	4.8	119
16	<i>Arabidopsis</i> HAF2 Gene Encoding TATA-binding Protein (TBP)-associated Factor TAF1, Is Required to Integrate Light Signals to Regulate Gene Expression and Growth. <i>Journal of Biological Chemistry</i> , 2005, 280, 1465-1473.	3.4	117
17	Modify the Histone to Win the Battle: Chromatin Dynamics in Plant-Pathogen Interactions. <i>Frontiers in Plant Science</i> , 2018, 9, 355.	3.6	106
18	Wheat chromatin architecture is organized in genome territories and transcription factories. <i>Genome Biology</i> , 2020, 21, 104.	8.8	99

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19	LHP1 Regulates H3K27me3 Spreading and Shapes the Three-Dimensional Conformation of the Arabidopsis Genome. <i>PLoS ONE</i> , 2016, 11, e0158936.	2.5	97
20	Ethylene induced plant stress tolerance by <i>Enterobacter</i> sp. SA187 is mediated by 2-oxo-4-methylthiobutyric acid production. <i>PLoS Genetics</i> , 2018, 14, e1007273.	3.5	95
21	The MYST histone acetyltransferases are essential for gametophyte development in Arabidopsis. <i>BMC Plant Biology</i> , 2008, 8, 121.	3.6	90
22	The BAF60 Subunit of the SWI/SNF Chromatin-Remodeling Complex Directly Controls the Formation of a Gene Loop at <i>FLOWERING LOCUS C</i> in Arabidopsis. <i>Plant Cell</i> , 2014, 26, 538-551.	6.6	82
23	MAPK-triggered chromatin reprogramming by histone deacetylase in plant innate immunity. <i>Genome Biology</i> , 2017, 18, 131.	8.8	73
24	The lncRNA APOLO interacts with the transcription factor WRKY42 to trigger root hair cell expansion in response to cold. <i>Molecular Plant</i> , 2021, 14, 937-948.	8.3	72
25	Plant-Specific Histone Deacetylases HDT1/2 Regulate <i>GIBBERELLIN 2-OXIDASE2</i> Expression to Control Arabidopsis Root Meristem Cell Number. <i>Plant Cell</i> , 2017, 29, 2183-2196.	6.6	69
26	Put your 3D glasses on: plant chromatin is on show. <i>Journal of Experimental Botany</i> , 2016, 67, 3205-3221.	4.8	59
27	The <i>Arabidopsis</i> lncRNA <i>ASCO</i> modulates the transcriptome through interaction with splicing factors. <i>EMBO Reports</i> , 2020, 21, e48977.	4.5	57
28	The Polyadenylation Factor Subunit CLEAVAGE AND POLYADENYLATION SPECIFICITY FACTOR30: A Key Factor of Programmed Cell Death and a Regulator of Immunity in Arabidopsis. <i>Plant Physiology</i> , 2014, 165, 732-746.	4.8	54
29	The Arabidopsis SWI/SNF protein BAF60 mediates seedling growth control by modulating DNA accessibility. <i>Genome Biology</i> , 2017, 18, 114.	8.8	53
30	The Polycomb protein <i>LHP1</i> regulates <i>Arabidopsis thaliana</i> stress responses through the repression of the <i>MYC</i> -dependent branch of immunity. <i>Plant Journal</i> , 2019, 100, 1118-1131.	5.7	52
31	Ploidy-dependent changes in the epigenome of symbiotic cells correlate with specific patterns of gene expression. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 4543-4548.	7.1	50
32	GCN5 modulates salicylic acid homeostasis by regulating H3K14ac levels at the 5' and 3' ends of its target genes. <i>Nucleic Acids Research</i> , 2020, 48, 5953-5966.	14.5	44
33	Involvement of Arabidopsis Hexokinase1 in Cell Death Mediated by <i>Myo</i> -Inositol Accumulation. <i>Plant Cell</i> , 2015, 27, 1801-1814.	6.6	42
34	Multiple Functions of Kip-Related Protein5 Connect Endoreduplication and Cell Elongation. <i>Plant Physiology</i> , 2013, 161, 1694-1705.	4.8	41
35	Chloroplast Dysfunction Causes Multiple Defects in Cell Cycle Progression in the Arabidopsis <i>crumpled leaf</i> Mutant. <i>Plant Physiology</i> , 2014, 166, 152-167.	4.8	37
36	Polycomb-dependent differential chromatin compartmentalization determines gene coregulation in <i>Arabidopsis</i> . <i>Genome Research</i> , 2021, 31, 1230-1244.	5.5	36

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37	Chromatin meets the cell cycle. <i>Journal of Experimental Botany</i> , 2014, 65, 2677-2689.	4.8	35
38	<i>Arabidopsis</i> ATRX Modulates H3.3 Occupancy and Fine-Tunes Gene Expression. <i>Plant Cell</i> , 2017, 29, 1773-1793.	6.6	35
39	A new role for histone demethylases in the maintenance of plant genome integrity. <i>ELife</i> , 2020, 9, .	6.0	33
40	Herboxidiene triggers splicing repression and abiotic stress responses in plants. <i>BMC Genomics</i> , 2017, 18, 260.	2.8	31
41	Chloroplast Activity and 3â€²phosphadenosine 5â€²phosphate Signaling Regulate Programmed Cell Death in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2016, 170, 1745-1756.	4.8	30
42	The quest for epigenetic regulation underlying unisexual flower development in <i>Cucumis melo</i> . <i>Epigenetics and Chromatin</i> , 2017, 10, 22.	3.9	27
43	Function of the Plant DNA Polymerase Epsilon in Replicative Stress Sensing, a Genetic Analysis. <i>Plant Physiology</i> , 2017, 173, 1735-1749.	4.8	26
44	A SWI/SNF Chromatin Remodelling Protein Controls Cytokinin Production through the Regulation of Chromatin Architecture. <i>PLoS ONE</i> , 2015, 10, e0138276.	2.5	25
45	The lncRNA MARS modulates the epigenetic reprogramming of the marneral cluster in response to ABA. <i>Molecular Plant</i> , 2022, 15, 840-856.	8.3	25
46	Evidence for a Role of <i>Arabidopsis</i> CDT1 Proteins in Gametophyte Development and Maintenance of Genome Integrity. <i>Plant Cell</i> , 2012, 24, 2779-2791.	6.6	24
47	A hierarchical transcriptional network activates specific CDK inhibitors that regulate G2 to control cell size and number in <i>Arabidopsis</i> . <i>Nature Communications</i> , 2022, 13, 1660.	12.8	22
48	The plant DNA polymerase theta is essential for the repair of replication-associated DNA damage. <i>Plant Journal</i> , 2021, 106, 1197-1207.	5.7	19
49	Integrative genome-wide analysis reveals the role of WIP proteins in inhibition of growth and development. <i>Communications Biology</i> , 2020, 3, 239.	4.4	16
50	DNA polymerase epsilon is required for heterochromatin maintenance in <i>Arabidopsis</i> . <i>Genome Biology</i> , 2020, 21, 283.	8.8	14
51	Cantaloupe melon genome reveals 3D chromatin features and structural relationship with the ancestral cucurbitaceae karyotype. <i>IScience</i> , 2022, 25, 103696.	4.1	12
52	The Mitochondrial DNA (mtDNA)-Associated Protein SWIB5 Influences mtDNA Architecture and Homologous Recombination. <i>Plant Cell</i> , 2017, 29, tpc.00899.2016.	6.6	11
53	The matrix revolutions: towards the decoding of the plant chromatin three-dimensional reality. <i>Journal of Experimental Botany</i> , 2020, 71, 5129-5147.	4.8	11
54	Immunity onset alters plant chromatin and utilizes EDA16 to regulate oxidative homeostasis. <i>PLoS Pathogens</i> , 2021, 17, e1009572.	4.7	10

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55	CmLHP1 proteins play a key role in plant development and sex determination in melon (<i>Cucumis) Tj ETQq1 1 0.784314 rgBT /Over	5.7	6
56	Sex Determination in Cucumis. Plant Genetics and Genomics: Crops and Models, 2017, , 307-319.	0.3	5
57	New partners for old friends: Plant SWI/SNF complexes. Molecular Plant, 2021, 14, 870-872.	8.3	4
58	Histone modification CHIP-seq on Arabidopsis thaliana plantlets. Bio-protocol, 2021, 11, e4211.	0.4	4
59	Chromatin architecture: A new dimension in the dynamic control of gene expression. Plant Signaling and Behavior, 2016, 11, e1232224.	2.4	1
60	Profiling Developmentally and Environmentally Controlled Chromatin Reprogramming. Methods in Molecular Biology, 2018, 1675, 3-30.	0.9	1
61	Plant Epigenetics: Non-coding RNAs as Emerging Regulators. RNA Technologies, 2017, , 129-147.	0.3	0
62	Three bona fide plant-specific SAGA subunits and their regulatory function. Molecular Plant, 2021, 14, 1033-1035.	8.3	0