

Vanessa Vermeirssen

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

1,906
citations

586496

16
h-index

843174

20
g-index

23
all docs

23
docs citations

23
times ranked

2909
citing authors

#	ARTICLE	IF	CITATIONS
1	Nearby transposable elements impact plant stress gene regulatory networks: a meta-analysis in <i>A. thaliana</i> and <i>S. lycopersicum</i> . <i>BMC Genomics</i> , 2022, 23, 18.	1.2	19
2	RRM2 enhances MYCN-driven neuroblastoma formation and acts as a synergistic target with CHK1 inhibition. <i>Science Advances</i> , 2022, 8, .	4.7	15
3	From DNA Copy Number Gains and Tumor Dependencies to Novel Therapeutic Targets for High-Risk Neuroblastoma. <i>Journal of Personalized Medicine</i> , 2021, 11, 1286.	1.1	2
4	Function, dynamics and evolution of network motif modules in integrated gene regulatory networks of worm and plant. <i>Nucleic Acids Research</i> , 2018, 46, 6480-6503.	6.5	33
5	<i>Arabidopsis</i> Ensemble Reverse-Engineered Gene Regulatory Network Discloses Interconnected Transcription Factors in Oxidative Stress. <i>Plant Cell</i> , 2015, 26, 4656-4679.	3.1	79
6	Reciprocal Responses in the Interaction between <i>Arabidopsis</i> and the Cell-Content-Feeding Chelicerate Herbivore Spider Mite <i>Tetranychus</i> . <i>Plant Physiology</i> , 2014, 164, 384-399.	2.3	151
7	The Membrane-Bound NAC Transcription Factor ANAC013 Functions in Mitochondrial Retrograde Regulation of the Oxidative Stress Response in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2013, 25, 3472-3490.	3.1	293
8	Identification of cis-regulatory elements specific for different types of reactive oxygen species in <i>Arabidopsis thaliana</i> . <i>Gene</i> , 2012, 499, 52-60.	1.0	36
9	Transcription regulatory networks in <i>Caenorhabditis elegans</i> inferred through reverse-engineering of gene expression profiles constitute biological hypotheses for metazoan development. <i>Molecular BioSystems</i> , 2009, 5, 1817.	2.9	23
10	Transcription factor modularity in a gene-centered <i>C. elegans</i> core neuronal protein-DNA interaction network. <i>Genome Research</i> , 2007, 17, 1061-1071.	2.4	87
11	Matrix and Steiner-triple-system smart pooling assays for high-performance transcription regulatory network mapping. <i>Nature Methods</i> , 2007, 4, 659-664.	9.0	62
12	Gateway-Compatible Yeast One-Hybrid Screens. <i>Cold Spring Harbor Protocols</i> , 2006, 2006, pdb.prot4590-pdb.prot4590.	0.2	67
13	Fractionation of angiotensin I converting enzyme inhibitory activity from pea and whey protein in vitro gastrointestinal digests. <i>Journal of the Science of Food and Agriculture</i> , 2005, 85, 399-405.	1.7	61
14	In vitro intestinal transport and antihypertensive activity of ACE inhibitory pea and whey digests. <i>International Journal of Food Sciences and Nutrition</i> , 2005, 56, 415-430.	1.3	49
15	Bioavailability of angiotensin I converting enzyme inhibitory peptides. <i>British Journal of Nutrition</i> , 2004, 92, 357-366.	1.2	460
16	A quantitative in silico analysis calculates the angiotensin I converting enzyme (ACE) inhibitory activity in pea and whey protein digests. <i>Biochimie</i> , 2004, 86, 231-239.	1.3	107
17	Influence of the lactokinin Ala-Leu-Pro-Met-His-Ile-Arg (ALPMHIR) on the release of endothelin-1 by endothelial cells. <i>Regulatory Peptides</i> , 2004, 118, 105-109.	1.9	93
18	Release of Angiotensin I Converting Enzyme (ACE) Inhibitory Activity during in Vitro Gastrointestinal Digestion: From Batch Experiment to Semicontinuous Model. <i>Journal of Agricultural and Food Chemistry</i> , 2003, 51, 5680-5687.	2.4	64

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19	Optimisation and validation of an angiotensin-converting enzyme inhibition assay for the screening of bioactive peptides. <i>Journal of Proteomics</i> , 2002, 51, 75-87.	2.4	198
20	Development of a Six-Stage Culture System for Simulating the Gastrointestinal Microbiota of Weaned Infants. <i>Microbial Ecology in Health and Disease</i> , 2001, 13, .	3.8	2