## Kieron D Edwards

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5842501/publications.pdf

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20 papers

1,531 citations

623734 14 h-index 713466 21 g-index

24 all docs

24 docs citations

times ranked

24

1861 citing authors

#	Article	IF	Citations
1	Evidence for anti-inflammatory effects and modulation of neurotransmitter metabolism by Salvia officinalis L BMC Complementary Medicine and Therapies, 2022, 22, 131.	2.7	5
2	Effects of Salvia officinalis L. and Chamaemelum nobile (L.) extracts on inflammatory responses in two models of human cells: Primary subcutaneous adipocytes and neuroblastoma cell line (SK-N-SH). Journal of Ethnopharmacology, 2021, 268, 113614.	4.1	7
3	Targeting Acr3 from <i>Ensifer medicae</i> to the plasma membrane or to the tonoplast of tobacco hairy roots allows arsenic extrusion or improved accumulation. Effect of <i>acr3</i> expression on the root transcriptome. Metallomics, 2019, 11, 1864-1886.	2.4	9
4	Accelerated flowering time reduces lifetime water use without penalizing reproductive performance in Arabidopsis. Plant, Cell and Environment, 2019, 42, 1847-1867.	5.7	10
5	Circadian clock components control daily growth activities by modulating cytokinin levels and cell divisionâ€associated gene expression in ⟨i⟩Populus⟨ i⟩ trees. Plant, Cell and Environment, 2018, 41, 1468-1482.	5 <b>.</b> 7	22
6	The clock gene circuit in $\langle i \rangle$ Arabidopsis $\langle  i \rangle$ includes a repressilator with additional feedback loops. Molecular Systems Biology, 2012, 8, 574.	7.2	386
7	Deciphering the complex leaf transcriptome of the allotetraploid species Nicotiana tabacum: a phylogenomic perspective. BMC Genomics, 2012, 13, 406.	2.8	39
8	Light inputs shape the Arabidopsis circadian system. Plant Journal, 2011, 66, 480-491.	5.7	78
9	Quantitative analysis of regulatory flexibility under changing environmental conditions. Molecular Systems Biology, 2010, 6, 424.	7.2	99
10	TobEA: an atlas of tobacco gene expression from seed to senescence. BMC Genomics, 2010, 11, 142.	2.8	55
11	Data assimilation constrains new connections and components in a complex, eukaryotic circadian clock model. Molecular Systems Biology, 2010, 6, 416.	7.2	145
12	Efficient utility-based clustering over high dimensional partition spaces. Bayesian Analysis, 2009, 4, .	3.0	5
13	BAG-1 is up-regulated in colorectal tumour progression and promotes colorectal tumour cell survival through increased NF-κB activity. Carcinogenesis, 2008, 29, 849-857.	2.8	49
14	Modelling non-stationary gene regulatory processes with a non-homogeneous Bayesian network and the allocation sampler. Bioinformatics, 2008, 24, 2071-2078.	4.1	55
15	Reconstruction of transcriptional dynamics from gene reporter data using differential equations. Bioinformatics, 2008, 24, 2901-2907.	4.1	58
16	Analysis of Circadian Leaf Movement Rhythms in Arabidopsis thaliana. Methods in Molecular Biology, 2007, 362, 103-113.	0.9	23
17	Analysis of Phase of LUCIFERASE Expression Reveals Novel Circadian Quantitative Trait Loci in Arabidopsis. Plant Physiology, 2006, 140, 1464-1474.	4.8	36
18	FLOWERING LOCUS C Mediates Natural Variation in the High-Temperature Response of the Arabidopsis Circadian Clock. Plant Cell, 2006, 18, 639-650.	6.6	276

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#	Article	lF	CITATIONS
19	Natural Allelic Variation in the Temperature-Compensation Mechanisms of the Arabidopsis thaliana Circadian ClockSequence data from this article have been deposited with the EMBL/GenBank Data Libraries under accession nos. AY685131 and AY685132 Genetics, 2005, 170, 387-400.	2.9	153
20	QTL for timing: a natural diversity of clock genes. Trends in Genetics, 2002, 18, 115-118.	6.7	6