

Eng-Ti Leslie Low

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8815640/publications.pdf>

Version: 2024-02-01

10
papers

1,138
citations

1163117

8
h-index

1372567

10
g-index

10
all docs

10
docs citations

10
times ranked

1817
citing authors

#	ARTICLE	IF	CITATIONS
1	Oil palm genome sequence reveals divergence of interfertile species in Old and New worlds. <i>Nature</i> , 2013, 500, 335-339.	27.8	468
2	Loss of Karma transposon methylation underlies the mantled somaclonal variant of oil palm. <i>Nature</i> , 2015, 525, 533-537.	27.8	405
3	The oil palm SHELL gene controls oil yield and encodes a homologue of SEEDSTICK. <i>Nature</i> , 2013, 500, 340-344.	27.8	167
4	Seqping: gene prediction pipeline for plant genomes using self-training gene models and transcriptomic data. <i>BMC Bioinformatics</i> , 2017, 18, 1-7.	2.6	25
5	Evidence-based gene models for structural and functional annotations of the oil palm genome. <i>Biology Direct</i> , 2017, 12, 21.	4.6	24
6	PalmXplore: oil palm gene database. <i>Database: the Journal of Biological Databases and Curation</i> , 2018, 2018, .	3.0	19
7	Characterization of Oil Palm Acyl-CoA-Binding Proteins and Correlation of Their Gene Expression with Oil Synthesis. <i>Plant and Cell Physiology</i> , 2020, 61, 735-747.	3.1	14
8	Variation for heterodimerization and nuclear localization among known and novel oil palm SHELL alleles. <i>New Phytologist</i> , 2020, 226, 426-440.	7.3	11
9	DNA methylation changes in clonally propagated oil palm. <i>Plant Cell Reports</i> , 2020, 39, 1219-1233.	5.6	3
10	A genetic platform for predicting and reducing non-tenera contamination in oil palm (<i>Elaeis</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tj 50 382 T	1.6	2