

# Tony Kuo

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11315920/publications.pdf>

Version: 2024-02-01

8  
papers

616  
citations

1478505

6  
h-index

1588992

8  
g-index

8  
all docs

8  
docs citations

8  
times ranked

990  
citing authors

#	ARTICLE	IF	CITATIONS
1	Multiple wheat genomes reveal global variation in modern breeding. <i>Nature</i> , 2020, 588, 277-283.	27.8	513
2	Trithorax Group Proteins Act Together with a Polycomb Group Protein to Maintain Chromatin Integrity for Epigenetic Silencing during Seed Germination in <i>Arabidopsis</i> . <i>Molecular Plant</i> , 2018, 11, 659-677.	8.3	47
3	EAGLE: Explicit Alternative Genome Likelihood Evaluator. <i>BMC Medical Genomics</i> , 2018, 11, 28.	1.5	16
4	De Novo Genome Assembly of the Japanese Wheat Cultivar Norin 61 Highlights Functional Variation in Flowering Time and Fusarium-Resistant Genes in East Asian Genotypes. <i>Plant and Cell Physiology</i> , 2021, 62, 8-27.	3.1	16
5	Identification of Binding Targets of a Pyrrole-Imidazole Polyamide KR12 in the LS180 Colorectal Cancer Genome. <i>PLoS ONE</i> , 2016, 11, e0165581.	2.5	11
6	Long non-coding RNA 2310069B03Rik functions as a suppressor of Ucp1 expression under prolonged cold exposure in murine beige adipocytes. <i>Bioscience, Biotechnology and Biochemistry</i> , 2020, 84, 305-313.	1.3	9
7	Involvement of Rev1 in alkylating agent-induced loss of heterozygosity in <i>Oryzias latipes</i> . <i>Genes To Cells</i> , 2020, 25, 124-138.	1.2	3
8	CRED: a rapid peak caller for Chem-seq data. <i>Journal of Open Source Software</i> , 2019, 4, 1423.	4.6	1