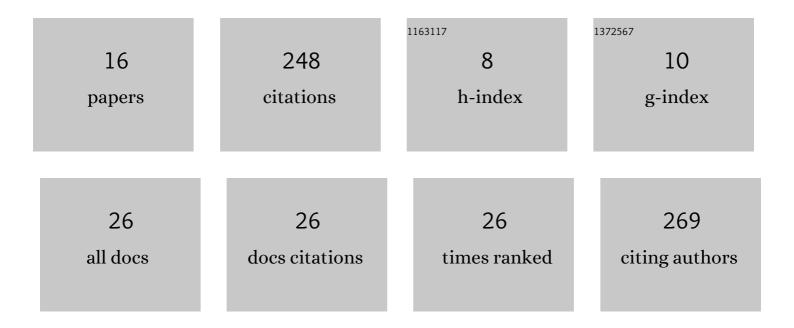
Danang Crysnanto

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9309788/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Structural variant-based pangenome construction has low sensitivity to variability of haplotype-resolved bovine assemblies. Nature Communications, 2022, 13, .	12.8	19
2	Characterization of a haplotype-reference panel for genotyping by low-pass sequencing in Swiss Large White pigs. BMC Genomics, 2021, 22, 290.	2.8	16
3	Novel functional sequences uncovered through a bovine multiassembly graph. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	42
4	Assessing genomic diversity and signatures of selection in Original Braunvieh cattle using whole-genome sequencing data. BMC Genomics, 2020, 21, 27.	2.8	47
5	Deletion of porcine <i>BOLL</i> is associated with defective acrosomes and subfertility in Yorkshire boars. Animal Genetics, 2020, 51, 945-949.	1.7	6
6	A 63â€bp insertion in exon 2 of the porcine <i>KIF21A</i> gene is associated with arthrogryposis multiplex congenita. Animal Genetics, 2020, 51, 820-823.	1.7	4
7	Bovine breed-specific augmented reference graphs facilitate accurate sequence read mapping and unbiased variant discovery. Genome Biology, 2020, 21, 184.	8.8	33
8	Activation of cryptic splicing in bovine WDR19 is associated with reduced semen quality and male fertility. PLoS Genetics, 2020, 16, e1008804.	3.5	26
9	Title is missing!. , 2020, 16, e1008804.		0
10	Title is missing!. , 2020, 16, e1008804.		0
11	Title is missing!. , 2020, 16, e1008804.		0
12	Title is missing!. , 2020, 16, e1008804.		0
13	Title is missing!. , 2020, 16, e1008804.		0
14	Title is missing!. , 2020, 16, e1008804.		0
15	Accurate sequence variant genotyping in cattle using variation-aware genome graphs. Genetics Selection Evolution, 2019, 51, 21.	3.0	26
16	Widespread gene duplication and adaptive evolution in the RNA interference pathways of the Drosophila obscura group. BMC Evolutionary Biology, 2019, 19, 99.	3.2	15