Danang Crysnanto

List of Publications by Year
in descending order

16 papers	248 citations	1163117 8 h-index	1372567 10 g-index
26 all docs	26 docs citations	26 times ranked	269 citing authors

#	Article	IF	Citations
1	Assessing genomic diversity and signatures of selection in Original Braunvieh cattle using whole-genome sequencing data. BMC Genomics, 2020, 21, 27.	2.8	47
2	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
3	Bovine breed-specific augmented reference graphs facilitate accurate sequence read mapping and unbiased variant discovery. Genome Biology, 2020, 21, 184.	8.8	33
4	Accurate sequence variant genotyping in cattle using variation-aware genome graphs. Genetics Selection Evolution, 2019, 51, 21.	3.0	26
5	Activation of cryptic splicing in bovine WDR19 is associated with reduced semen quality and male fertility. PLoS Genetics, 2020, 16, e1008804.	3.5	26
6	Structural variant-based pangenome construction has low sensitivity to variability of haplotype-resolved bovine assemblies. Nature Communications, 2022, 13, .	12.8	19
7	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
8	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
9	Deletion of porcine *BOLL* is associated with defective acrosomes and subfertility in Yorkshire boars. Animal Genetics, 2020, 51, 945-949.	1.7	6
10	A 63‑bp insertion in exon 2 of the porcine *KIF21A* gene is associated with arthrogryposis multiplex congenita. Animal Genetics, 2020, 51, 820-823.	1.7	4
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	Title is missing!. , 2020, 16, e1008804.		0
16	Title is missing!. , 2020, 16, e1008804.		0