

# Danang Crysanto

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9309788/publications.pdf>

Version: 2024-02-01

16  
papers

248  
citations

1163117  
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1372567  
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26  
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docs citations

26  
times ranked

269  
citing authors

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