Sven O Twardziok

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2550871/publications.pdf

Version: 2024-02-01

38 papers 8,342 citations

24 h-index

257450

345221 36 g-index

41 all docs

41 docs citations

41 times ranked

11401 citing authors

#	Article	IF	CITATIONS
1	Complex landscape of alternative splicing in myeloid neoplasms. Leukemia, 2021, 35, 1108-1120.	7.2	39
2	Hypertension delays viral clearance and exacerbates airway hyperinflammation in patients with COVID-19. Nature Biotechnology, 2021, 39, 705-716.	17. 5	129
3	Implementing FAIR data management within the German Network for Bioinformatics Infrastructure (de.NBI) exemplified by selected use cases. Briefings in Bioinformatics, 2021, 22, .	6.5	18
4	Homozygous BCMA gene deletion in response to anti-BCMA CAR T cells in a patient with multiple myeloma. Nature Medicine, 2021, 27, 616-619.	30.7	140
5	The combination of WGS and RNA-Seq is superior to conventional diagnostic tests in multiple myeloma: Ready for prime time?. Cancer Genetics, 2020, 242, 15-24.	0.4	32
6	Molecular landscape and clonal architecture of adult myelodysplastic/myeloproliferative neoplasms. Blood, 2020, 136, 1851-1862.	1.4	112
7	COVID-19 severity correlates with airway epithelium–immune cell interactions identified by single-cell analysis. Nature Biotechnology, 2020, 38, 970-979.	17.5	887
8	"Somatic―and "pathogenic― is the classification strategy applicable in times of large-scale sequencing?. Haematologica, 2019, 104, 1515-1520.	3.5	9
9	Dark-matter matters: Discriminating subtle blood cancers using the darkest DNA. PLoS Computational Biology, 2019, 15, e1007332.	3.2	7
10	Durum wheat genome highlights past domestication signatures and future improvement targets. Nature Genetics, 2019, 51, 885-895.	21.4	576
11	Combining RNA-seq data and homology-based gene prediction for plants, animals and fungi. BMC Bioinformatics, 2018, 19, 189.	2.6	192
12	Gene Prediction in the Barley Genome. Compendium of Plant Genomes, 2018, , 73-88.	0.5	0
13	Shifting the limits in wheat research and breeding using a fully annotated reference genome. Science, 2018, 361, .	12.6	2,424
14	Chromosome-scale comparative sequence analysis unravels molecular mechanisms of genome dynamics between two wheat cultivars. Genome Biology, 2018, 19, 104.	8.8	54
15	The pseudogenes of barley. Plant Journal, 2018, 93, 502-514.	5.7	14
16	A chromosome conformation capture ordered sequence of the barley genome. Nature, 2017, 544, 427-433.	27.8	1,365
17	Light and Plastid Signals Regulate Different Sets of Genes in the Albino Mutant Pap7-1. Plant Physiology, 2017, 175, 1203-1219.	4.8	29
18	Wild emmer genome architecture and diversity elucidate wheat evolution and domestication. Science, 2017, 357, 93-97.	12.6	781

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19	Genome sequence of the progenitor of the wheat D genome Aegilops tauschii. Nature, 2017, 551, 498-502.	27.8	563
20	Towards a wholeâ€genome sequence for rye (<i>Secale cereale</i> L.). Plant Journal, 2017, 89, 853-869.	5.7	238
21	The repetitive landscape of the 5100 Mbp barley genome. Mobile DNA, 2017, 8, 22.	3.6	49
22	Probiotic Treatment Decreases the Number of CD14-Expressing Cells in Porcine Milk Which Correlates with Several Intestinal Immune Parameters in the Piglets. Frontiers in Immunology, 2015, 6, 108.	4.8	25
23	Effects of age and zinc supplementation on transport properties in the jejunum of piglets. Journal of Animal Physiology and Animal Nutrition, 2015, 99, 542-552.	2.2	10
24	The General Composition of the Faecal Virome of Pigs Depends on Age, but Not on Feeding with a Probiotic Bacterium. PLoS ONE, 2014, 9, e88888.	2.5	34
25	Crossâ€ŧalk Between Host, Microbiome and Probiotics: A Systems Biology Approach for Analyzing the Effects of Probiotic <i>Enterococcus faecium</i> NCIMB 10415 in Piglets. Molecular Informatics, 2014, 33, 171-182.	2.5	4
26	Elevated dietary zinc oxide levels do not have a substantial effect on porcine reproductive and respiratory syndrome virus (PPRSV) vaccination and infection. Virology Journal, 2014, 11, 140.	3.4	3
27	High-dose dietary zinc oxide mitigates infection with transmissible gastroenteritis virus in piglets. BMC Veterinary Research, 2014, 10, 75.	1.9	31
28	Enterococcus faecium NCIMB 10415 supplementation affects intestinal immune-associated gene expression in post-weaning piglets. Veterinary Immunology and Immunopathology, 2014, 157, 65-77.	1.2	35
29	Dietary Enterococcus faecium NCIMB 10415 and Zinc Oxide Stimulate Immune Reactions to Trivalent Influenza Vaccination in Pigs but Do Not Affect Virological Response upon Challenge Infection. PLoS ONE, 2014, 9, e87007.	2.5	14
30	Identification of an avian group A rotavirus containing a novel VP4 gene with a close relationship to those of mammalian rotaviruses. Journal of General Virology, 2013, 94, 136-142.	2.9	245
31	Antiviral effects of a probiotic Enterococcus faecium strain against transmissible gastroenteritis coronavirus. Archives of Virology, 2013, 158, 799-807.	2.1	66
32	Porcine intestinal mast cells. Evaluation of different fixatives for histochemical staining techniques considering tissue shrinkage. European Journal of Histochemistry, 2013, 57, 21.	1.5	27
33	Characterization of the effects of Enterococcus faecium on intestinal epithelial transport properties in piglets1. Journal of Animal Science, 2013, 91, 1707-1718.	0.5	29
34	Inhibitory Influence of Enterococcus faecium on the Propagation of Swine Influenza A Virus In Vitro. PLoS ONE, 2013, 8, e53043.	2.5	54
35	Allelic variations in coding regions of the vitamin D receptor gene in dairy cows and potential susceptibility to periparturient hypocalcaemia. Journal of Dairy Research, 2012, 79, 423-428.	1.4	8
36	Simultaneous Identification of DNA and RNA Viruses Present in Pig Faeces Using Process-Controlled Deep Sequencing. PLoS ONE, 2012, 7, e34631.	2.5	77

#	Article	IF	CITATIONS
37	Evidence for Regulated Interleukin-4 Expression in Chondrocyte-Scaffolds under In Vitro Inflammatory Conditions. PLoS ONE, 2011, 6, e25749.	2.5	18
38	Stochasticity in reactions. , 2010, , .		3