Giancarlo Russo

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7519373/publications.pdf

Version: 2024-02-01

22 papers 883 citations

15 h-index 19 g-index

27 all docs

27 docs citations

27 times ranked

1694 citing authors

#	Article	IF	CITATIONS
1	Re-evaluation of single nucleotide variants and identification of structural variants in a cohort of 45 sudden unexplained death cases. International Journal of Legal Medicine, 2021, 135, 1341-1349.	1.2	8
2	Assessing time dependent changes in microbial composition of biological crime scene traces using microbial RNA markers. Forensic Science International: Genetics, 2021, 53, 102537.	1.6	17
3	Degradation of human mRNA transcripts over time as an indicator of the time since deposition (TsD) in biological crime scene traces. Forensic Science International: Genetics, 2021, 53, 102524.	1.6	23
4	Genome-wide transcriptomics identifies an early preclinical signature of prion infection. PLoS Pathogens, 2020, 16, e1008653.	2.1	40
5	Germâ€free and microbiotaâ€associated mice yield small intestinal epithelial organoids with equivalent and robust transcriptome/proteome expression phenotypes. Cellular Microbiology, 2020, 22, e13191.	1.1	26
6	Genome-wide transcriptomics identifies an early preclinical signature of prion infection. , 2020, 16, e1008653.		0
7	Genome-wide transcriptomics identifies an early preclinical signature of prion infection. , 2020, 16, e1008653.		O
8	Periodontal bacterial supernatants modify differentiation, migration and inflammatory cytokine expression in human periodontal ligament stem cells. PLoS ONE, 2019, 14, e0219181.	1.1	17
9	De novo genome sequencing and comparative stage-specific transcriptomic analysis of Dirofilaria repens. International Journal for Parasitology, 2019, 49, 911-919.	1.3	3
10	Transcription and microbial profiling of body fluids using a massively parallel sequencing approach. Forensic Science International: Genetics, 2019, 43, 102149.	1.6	23
11	Lack of the pH-sensing Receptor TDAG8 [GPR65] in Macrophages Plays a Detrimental Role in Murine Models of Inflammatory Bowel Disease. Journal of Crohn's and Colitis, 2019, 13, 245-258.	0.6	39
12	Exome analysis in 34 sudden unexplained death (SUD) victims mainly identified variants in channelopathy-associated genes. International Journal of Legal Medicine, 2018, 132, 1057-1065.	1.2	38
13	Post-mortem whole-exome analysis in a large sudden infant death syndrome cohort with a focus on cardiovascular and metabolic genetic diseases. European Journal of Human Genetics, 2017, 25, 404-409.	1.4	98
14	Stable core virome despite variable microbiome after fecal transfer. Gut Microbes, 2017, 8, 214-220.	4.3	60
15	RNase H As Gene Modifier, Driver of Evolution and Antiviral Defense. Frontiers in Microbiology, 2017, 8, 1745.	1.5	49
16	Long-term changes of bacterial and viral compositions in the intestine of a recovered <i>Clostridium difficile<ii> patient after fecal microbiota transplantation. Journal of Physical Education and Sports Management, 2016, 2, a000448.</i>	0.5	50
17	Strictly co-isogenic C57BL/6J- $\langle i \rangle$ Prnp $\langle i \rangle$ â $^{\circ}$ /â $^{\circ}$ mice: A rigorous resource for prion science. Journal of Experimental Medicine, 2016, 213, 313-327.	4.2	98
18	Cell Cycle Constraints and Environmental Control of Local DNA Hypomethylation in α-Proteobacteria. PLoS Genetics, 2016, 12, e1006499.	1.5	25

#	Article	IF	CITATION
19	Strictly co-isogenic C57BL/6J-Prnpâ^/lâ^mice: A rigorous resource for prion science. Journal of Cell Biology, 2016, 212, 2126OIA42.	2.3	O
20	Highly sensitive, non-invasive detection of colorectal cancer mutations using single molecule, third generation sequencing. Applied & Translational Genomics, 2015, 7, 32-39.	2.1	16
21	Targeted Combinatorial Alternative Splicing Generates Brain Region-Specific Repertoires of Neurexins. Neuron, 2014, 84, 386-398.	3 . 8	165
22	SIRPα polymorphisms, but not the prion protein, control phagocytosis of apoptotic cells. Journal of Experimental Medicine, 2013, 210, 2539-2552.	4.2	67