## Stefano Colella

List of Publications by Year in descending order

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Version: 2024-02-01

44 papers

5,633 citations

172207 29 h-index 253896 43 g-index

48 all docs 48 docs citations

48 times ranked

9778 citing authors

#	Article	IF	CITATIONS
1	Genome Sequence of the Pea Aphid Acyrthosiphon pisum. PLoS Biology, 2010, 8, e1000313.	2.6	913
2	Rapid transcriptional plasticity of duplicated gene clusters enables a clonally reproducing aphid to colonise diverse plant species. Genome Biology, 2017, 18, 27.	3.8	624
3	hsa-miR-210 Is Induced by Hypoxia and Is an Independent Prognostic Factor in Breast Cancer. Clinical Cancer Research, 2008, 14, 1340-1348.	3.2	617
4	QuantiSNP: an Objective Bayes Hidden-Markov Model to detect and accurately map copy number variation using SNP genotyping data. Nucleic Acids Research, 2007, 35, 2013-2025.	6.5	525
5	Sensitive and quantitative universal Pyrosequencingâ,,¢ methylation analysis of CpG sites. BioTechniques, 2003, 35, 146-150.	0.8	457
6	LRRTM1 on chromosome 2p12 is a maternally suppressed gene that is associated paternally with handedness and schizophrenia. Molecular Psychiatry, 2007, 12, 1129-1139.	4.1	300
7	Genomic insight into the amino acid relations of the pea aphid, <i>Acyrthosiphon pisum</i> , with its symbiotic bacterium <i>Buchnera aphidicola</i> . Insect Molecular Biology, 2010, 19, 249-258.	1.0	219
8	Molecular Analysis of Mutations in the CSB(ERCC6) Gene in Patients with Cockayne Syndrome. American Journal of Human Genetics, 1998, 62, 77-85.	2.6	145
9	Loss of Heterozygosity on Chromosome 10 Is More Extensive in Primary (De Novo) Than in Secondary Glioblastomas. Laboratory Investigation, 2000, 80, 65-72.	1.7	145
10	MicroRNA-10b and breast cancer metastasis. Nature, 2008, 455, E8-E9.	13.7	134
11	A statistical approach for detecting genomic aberrations in heterogeneous tumor samples from single nucleotide polymorphism genotyping data. Genome Biology, 2010, 11, R92.	3.8	125
12	Genome-Wide Hypomethylation in Head and Neck Cancer Is More Pronounced in HPV-Negative Tumors and Is Associated with Genomic Instability. PLoS ONE, 2009, 4, e4941.	1.1	114
13	Molecular evolutionary trends and feeding ecology diversification in the Hemiptera, anchored by the milkweed bug genome. Genome Biology, 2019, 20, 64.	3.8	114
14	Matrix-Assisted Laser Desorption/Ionisation, Time-of-Flight Mass Spectrometry in Genomics Research. PLoS Genetics, 2006, 2, e100.	1.5	103
15	alpha-Haemoglobin stabilising protein is a quantitative trait gene that modifies the phenotype of beta-thalassaemia. British Journal of Haematology, 2006, 133, 675-682.	1.2	79
16	New insight into the RNA interference response against cathepsin-L gene in the pea aphid, Acyrthosiphon pisum: Molting or gut phenotypes specifically induced by injection or feeding treatments. Insect Biochemistry and Molecular Biology, 2014, 51, 20-32.	1.2	75
17	Direct flow cytometry measurements reveal a fine-tuning of symbiotic cell dynamics according to the host developmental needs in aphid symbiosis. Scientific Reports, 2016, 6, 19967.	1.6	71
18	GenoSNP: a variational Bayes within-sample SNP genotyping algorithm that does not require a reference population. Bioinformatics, 2008, 24, 2209-2214.	1.8	65

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19	Reduced expression of the A? subunit of protein phosphatase 2A in human gliomas in the absence of mutations in the A? and A? subunit genes. International Journal of Cancer, 2001, 93, 798-804.	2.3	60
20	Identical mutations in the CSB gene associated with either Cockayne syndrome or the DeSanctis-Cacchione variant of xeroderma pigmentosum. Human Molecular Genetics, 2000, 9, 1171-1175.	1.4	53
21	Tyrosine pathway regulation is host-mediated in the pea aphid symbiosis during late embryonic and early larval development. BMC Genomics, 2013, 14, 235.	1.2	51
22	Alterations in the CSB Gene in Three Italian Patients with the Severe Form of Cockayne Syndrome (CS) But Without Clinical Photosensitivity. Human Molecular Genetics, 1999, 8, 935-941.	1.4	49
23	Systemic analysis of the symbiotic function of Buchnera aphidicola, the primary endosymbiont of the pea aphid Acyrthosiphon pisum. Comptes Rendus - Biologies, 2009, 332, 1034-1049.	0.1	49
24	A Genomic Reappraisal of Symbiotic Function in the Aphid/Buchnera Symbiosis: Reduced Transporter Sets and Variable Membrane Organisations. PLoS ONE, 2011, 6, e29096.	1.1	44
25	The transposable element-rich genome of the cereal pest Sitophilus oryzae. BMC Biology, 2021, 19, 241.	1.7	40
26	N-(4-Hydroxyphenyl)retinamide and nitric oxide pro-drugs exhibit apoptotic and anti-invasive effects against bone metastatic breast cancer cells. Carcinogenesis, 2006, 27, 568-577.	1.3	37
27	Multimodal dynamic response of the <i>Buchnera aphidicola</i> pLeu plasmid to variations in leucine demand of its host, the pea aphid <i>Acyrthosiphon pisum</i> Molecular Microbiology, 2011, 81, 1271-1285.	1.2	35
28	Cloning a new human gene from chromosome 21q22.3 encoding a glutamic acid-rich protein expressed in heart and skeletal muscle. Human Genetics, 1997, 99, 387-392.	1.8	34
29	Disruption of phenylalanine hydroxylase reduces adult lifespan and fecundity, and impairs embryonic development in parthenogenetic pea aphids. Scientific Reports, 2016, 6, 34321.	1.6	34
30	Hostâ€specific competitiveness to form nodules in <i>Rhizobium leguminosarum</i> symbiovar <i>viciae</i> . New Phytologist, 2020, 226, 555-568.	3.5	33
31	Mutation analysis of hBUB1, hBUBR1 and hBUB3 genes in glioblastomas. Acta Neuropathologica, 2001, 101, 297-304.	3.9	29
32	Genome-wide loss of heterozygosity analysis of WT1-wild-type and WT1-mutant Wilms tumors. Genes Chromosomes and Cancer, 2005, 43, 172-180.	1.5	29
33	DiCoExpress: a tool to process multifactorial RNAseq experiments from quality controls to co-expression analysis through differential analysis based on contrasts inside GLM models. Plant Methods, 2020, 16, 68.	1.9	29
34	Identification and characterization of Vietnamese coffee bacterial endophytes displaying in vitro antifungal and nematicidal activities. Microbiological Research, 2021, 242, 126613.	2.5	28
35	Molecular signatures of metastasis in head and neck cancer. Head and Neck, 2008, 30, 1273-1283.	0.9	27
36	Altered Intra-Nuclear Organisation of Heterochromatin and Genes in ICF Syndrome. PLoS ONE, 2010, 5, e11364.	1.1	25

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37	ArthropodaCyc: a CycADS powered collection of BioCyc databases to analyse and compare metabolism of arthropods. Database: the Journal of Biological Databases and Curation, 2016, 2016, baw081.	1.4	22
38	Responses of mature symbiotic nodules to the whole-plant systemic nitrogen signaling. Journal of Experimental Botany, 2020, 71, 5039-5052.	2.4	22
39	The anatomy of an aphid genome: From sequence to biology. Comptes Rendus - Biologies, 2010, 333, 464-473.	0.1	20
40	CycADS: an annotation database system to ease the development and update of BioCyc databases. Database: the Journal of Biological Databases and Curation, 2011, 2011, bar008-bar008.	1.4	16
41	Bacteriocyte Reprogramming to Cope With Nutritional Stress in a Phloem Sap Feeding Hemipteran, the Pea Aphid Acyrthosiphon pisum. Frontiers in Physiology, 2018, 9, 1498.	1.3	15
42	Systemic control of nodule formation by plant nitrogen demand requires autoregulation-dependent and independent mechanisms. Journal of Experimental Botany, 2021, 72, 7942-7956.	2.4	7
43	Prenatal diagnosis of 30 fetuses at risk for fragile X syndrome. , 1996, 64, 187-190.		2
44	Analysis of DNA Methylation at the Human Alpha Globin Cluster during Hematopoiesis Blood, 2008, 112, 1861-1861.	0.6	0