Fulvia Ferrazzi

List of Publications by Year
in descending order

53 papers	1,647 citations	394421 19 h-index	315739 38 g-index
59 all docs	59 docs citations	59 times ranked	3176 citing authors

#	Article	IF	Citations
1	Light Chain Restriction in Proximal Tubulesâ€"Implications for Light Chain Proximal Tubulopathy. Frontiers in Medicine, 2022, 9, 723758.	2.6	2
2	Can Gene Expression Analysis in Zero-Time Biopsies Predict Kidney Transplant Rejection?. Frontiers in Medicine, 2022, 9, 793744.	2.6	1
3	Recurrent novel HMGA2-NCOR2 fusions characterize a subset of keratin-positive giant cell-rich soft tissue tumors. Modern Pathology, 2021, 34, 1507-1520.	5.5	22
4	Validation of the â€˜Inflammatory Bowel Diseaseâ€"Distribution, Chronicity, Activity [IBD-DCA] Scoreâ€™ for Ulcerative Colitis and CrohnÂ´s Disease. Journal of Crohn's and Colitis, 2021, 15, 1621-1630.	1.3	21
5	Significance of Glomerular Immune Reactivity in Time Zero Biopsies for Allograft Survival Beyond IgA. Frontiers in Medicine, 2021, 8, 656840.	2.6	2
6	Functional genomics meta-analysis to identify gene set enrichment networks in cardiac hypertrophy. Biological Chemistry, 2021, 402, 953-972.	2.5	3
7	IQGAP3, a YAP Target, Is Required for Proper Cell-Cycle Progression and Genome Stability. Molecular Cancer Research, 2021, 19, 1712-1726.	3.4	11
8	Complement in Renal Disease as a Potential Contributor to Arterial Hypertension. Kidney and Blood Pressure Research, 2021, 46, 362-376.	2.0	4
9	Lipomatous Solitary Fibrous Tumors Harbor Rare NAB2-STAT6 Fusion Variants and Show Up-Regulation of the Gene PPARG, Encoding for a Regulator of Adipocyte Differentiation. American Journal of Pathology, 2021, 191, 1314-1324.	3.8	5
10	Reproducibility of mRNA-Based Testing of ESR1, PGR, ERBB2, and MKI67 Expression in Invasive Breast Cancerâ€"A Europe-Wide External Quality Assessment. Cancers, 2021, 13, 4718.	3.7	6
11	Myogenin controls via AKAP6 non-centrosomal microtubule-organizing center formation at the nuclear envelope. ELife, 2021, 10, .	6.0	6
12	Relevance of glomerular C4d deposition in pediatric patients with Henoch-SchÃ¶nlein Purpura compared to IgA nephritis. Journal of Nephropathology, 2021, 10, e16-e16.	0.2	1
13	28P Europe-side external quality assessment (EQA) of RNA based testing of ER, PR, HER2 and Ki67 in invasive breast cancer. Annals of Oncology, 2020, 31, S25.	1.2	0
14	High Stroma T-Cell Infiltration is Associated with Better Survival in Stage pT1 Bladder Cancer. International Journal of Molecular Sciences, 2020, 21, 8407.	4.1	14
15	Loss of PHF6 leads to aberrant development of human neuron-like cells. Scientific Reports, 2020, 10, 19030.	3.3	3
16	Sox11 is an Activity-Regulated Gene with Dentate-Gyrus-Specific Expression Upon General Neural Activation. Cerebral Cortex, 2020, 30, 3731-3743.	2.9	7
17	Genetic interaction screen for severe neurodevelopmental disorders reveals a functional link between Ube3a and Mef2 in Drosophila melanogaster. Scientific Reports, 2020, 10, 1204.	3.3	8
18	Genomeâ€wide cooperation of <scp>EMT</scp> transcription factor <scp>ZEB</scp> 1 with <scp>YAP</scp> and <scp>AP</scp> â€1 in breast cancer. EMBO Journal, 2020, 39, e103209.	7.8	104

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19	Locally renewing resident synovial macrophages provide a protective barrier for the joint. Nature, 2019, 572, 670-675.	27.8	345
20	Evolutionary conserved networks of human height identify multiple Mendelian causes of short stature. European Journal of Human Genetics, 2019, 27, 1061-1071.	2.8	11
21	Polyol Pathway Links Glucose Metabolism to the Aggressiveness of Cancer Cells. Cancer Research, 2018, 78, 1604-1618.	0.9	83
22	Clinical relevance of systematic phenotyping and exome sequencing in patients with short stature. Genetics in Medicine, 2018, 20, 630-638.	2.4	101
23	Integrative bioinformatics analysis characterizing the role of EDC3 in mRNA decay and its association to intellectual disability. BMC Medical Genomics, 2018, 11, 41.	1.5	5
24	Effects of Anti-Integrin Treatment With Vedolizumab on Immune Pathways and Cytokines in Inflammatory Bowel Diseases. Frontiers in Immunology, 2018, 9, 1700.	4.8	38
25	Hypomorphic Pathogenic Variants in TAF13 Are Associated with Autosomal-Recessive Intellectual Disability and Microcephaly. American Journal of Human Genetics, 2017, 100, 555-561.	6.2	26
26	Pseudoexfoliation syndrome-associated genetic variants affect transcription factor binding and alternative splicing of LOXL1. Nature Communications, 2017, 8, 15466.	12.8	57
27	FAM13A is associated with non-small cell lung cancer (NSCLC) progression and controls tumor cell proliferation and survival. OncoImmunology, 2017, 6, e1256526.	4.6	44
28	Cardiac injury of the newborn mammalian heart accelerates cardiomyocyte terminal differentiation. Scientific Reports, 2017, 7, 8362.	3.3	32
29	Identification of Genetic Signatures and Immune Mechanisms That Define Therapeutic Response and Failure to Anti-Integrin Therapy with Vedolizumab in Patients with IBD. Gastroenterology, 2017, 152, S386.	1.3	0
30	PEDF Is Associated with the Termination of Chondrocyte Phenotype and Catabolism of Cartilage Tissue. BioMed Research International, 2017, 2017, 1-13.	1.9	7
31	From basic mechanisms to clinical applications in heart protection, new players in cardiovascular diseases and cardiac theranostics: meeting report from the third international symposium on "New frontiers in cardiovascular research" Basic Research in Cardiology, 2016, 111, 69.	5.9	41
32	CD163+ M2c-like macrophages predominate in renal biopsies from patients with lupus nephritis. Arthritis Research and Therapy, 2016, 18, 90.	3.5	92
33	Rhinovirus inhibits IL-17A and the downstream immune responses in allergic asthma. Mucosal Immunology, 2016, 9, 1183-1192.	6.0	24
34	IL-7 Abrogates the Immunosuppressive Function of Human Double-Negative T Cells by Activating Akt/mTOR Signaling. Journal of Immunology, 2015, 195, 3139-3148.	0.8	16
35	Gene network analysis: from heart development to cardiac therapy. Thrombosis and Haemostasis, 2015, 113, 521-531.	3.4	7
36	Probabilistic Modelling with Bayesian Networks. , 2014, , 257-280.		4

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37	Regeneration-associated WNT Signaling Is Activated in Long-term Reconstituting AC133bright Acute Myeloid Leukemia Cells. Neoplasia, 2012, 14, 1236-IN45.	5.3	26
38	TGFÎ²1-Induced Baf60c Regulates both Smooth Muscle Cell Commitment and Quiescence. PLoS ONE, 2012, 7, e47629.	2.5	12
39	Inferring cell cycle feedback regulation from gene expression data. Journal of Biomedical Informatics, 2011, 44, 565-575.	4.3	9
40	Predictive data mining in clinical medicine: a focus on selected methods and applications. Wiley Interdisciplinary Reviews: Data Mining and Knowledge Discovery, 2011, 1, 416-430.	6.8	73
41	Nephronectin regulates atrioventricular canal differentiation via Bmp4-Has2 signaling in zebrafish. Development (Cambridge), 2011, 138, 4499-4509.	2.5	56
42	R Engine Cell: integrating R into the i2b2 software infrastructure. Journal of the American Medical Informatics Association: JAMIA, 2011, 18, 314-317.	4.4	17
43	TWEAK is a positive regulator of cardiomyocyte proliferation. Cardiovascular Research, 2010, 85, 681-690.	3.8	90
44	Cardiac Deletion of Smyd2 Is Dispensable for Mouse Heart Development. PLoS ONE, 2010, 5, e9748.	2.5	63
45	Phenotype forecasting with SNPs data through gene-based Bayesian networks. BMC Bioinformatics, 2009, 10, S7.	2.6	11
46	A stability-based algorithm to validate hierarchical clusters of genes. International Journal of Knowledge Engineering and Soft Data Paradigms, 2009, 1, 318.	0.0	2
47	TimeClust: a clustering tool for gene expression time series. Bioinformatics, 2008, 24, 430-432.	4.1	50
48	Bayesian approaches to reverse engineer cellular systems: a simulation study on nonlinear Gaussian networks. BMC Bioinformatics, 2007, 8, S2.	2.6	38
49	Inferring gene regulatory networks by integrating static and dynamic data. International Journal of Medical Informatics, 2007, 76, S462-S475.	3.3	7
50	Can we use linear Gaussian networks to model dynamic interactions among genes? Results from a simulation study. , 2006, , .		19
51	Dynamic Bayesian Networks in Modelling Cellular Systems: a Critical Appraisal on Simulated Data. , 2006, , .		2
52	Inferring gene expression networks via static and dynamic data integration. Studies in Health Technology and Informatics, 2006, 124, 119-24.	0.3	3
53	Random Walk Models for Bayesian Clustering of Gene Expression Profiles. Applied Bioinformatics, 2005, 4, 263-276.	1.6	14