Giulia Carini

List of Publications by Year
in descending order

10 papers	204 citations	1040056 9 h-index	1372567 10 g-index
10 all docs	10 docs citations	10 times ranked	324 citing authors

#	Article	IF	Citations
1	Transcriptome-Wide Expression Profiling in Skin Fibroblasts of Patients with Joint Hypermobility Syndrome/Ehlers-Danlos Syndrome Hypermobility Type. PLoS ONE, 2016, 11, e0161347.	2.5	40
2	Transcriptome analysis of skin fibroblasts with dominant negative COL3A1 mutations provides molecular insights into the etiopathology of vascular Ehlers-Danlos syndrome. PLoS ONE, 2018, 13, e0191220.	2.5	31
3	Modulation by chronic stress and ketamine of ionotropic AMPA/NMDA and metabotropic glutamate receptors in the rat hippocampus. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2021, 104, 110033.	4.8	24
4	Molecular insights in the pathogenesis of classical Ehlers-Danlos syndrome from transcriptome-wide expression profiling of patientsâ€™ skin fibroblasts. PLoS ONE, 2019, 14, e0211647.	2.5	22
5	miR-9-5p is involved in the rescue of stress-dependent dendritic shortening of hippocampal pyramidal neurons induced by acute antidepressant treatment with ketamine. Neurobiology of Stress, 2021, 15, 100381.	4.0	20
6	Acute Ketamine Facilitates Fear Memory Extinction in a Rat Model of PTSD Along With Restoring Glutamatergic Alterations and Dendritic Atrophy in the Prefrontal Cortex. Frontiers in Pharmacology, 2022, 13, 759626.	3.5	17
7	GLUT10â€”Lacking in Arterial Tortuosity Syndromeâ€”Is Localized to the Endoplasmic Reticulum of Human Fibroblasts. International Journal of Molecular Sciences, 2017, 18, 1820.	4.1	15
8	Ehlersâ€“Danlos syndrome with lethal cardiac valvular dystrophy in males carrying a novel splice mutation in FLNA. American Journal of Medical Genetics, Part A, 2017, 173, 169-176.	1.2	13
9	The Potential Role of miRNAs in Cognitive Frailty. Frontiers in Aging Neuroscience, 2021, 13, 763110.	3.4	12
10	miRNome Profiling Detects miR-101-3p and miR-142-5p as Putative Blood Biomarkers of Frailty Syndrome. Genes, 2022, 13, 231.	2.4	10