Peter VrtaÄnik

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5325846/publications.pdf

Version: 2024-02-01

1040056 1281871 12 618 9 11 citations h-index g-index papers 12 12 12 1299 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	The many faces of estrogen signaling. Biochemia Medica, 2014, 24, 329-342.	2.7	296
2	Somatic mutagenesis in satellite cells associates with human skeletal muscle aging. Nature Communications, 2018, 9, 800.	12.8	94
3	MiR-148a the epigenetic regulator of bone homeostasis is increased in plasma of osteoporotic postmenopausal women. Wiener Klinische Wochenschrift, 2016, 128, 519-526.	1.9	63
4	Whole genome DNA sequencing provides an atlas of somatic mutagenesis in healthy human cells and identifies a tumor-prone cell type. Genome Biology, 2019, 20, 285.	8.8	46
5	Epigenetic enzymes influenced by oxidative stress and hypoxia mimetic in osteoblasts are differentially expressed in patients with osteoporosis and osteoarthritis. Scientific Reports, 2018, 8, 16215.	3.3	31
6	Rare progerin-expressing preadipocytes and adipocytes contribute to tissue depletion over time. Scientific Reports, 2017, 7, 4405.	3.3	24
7	Hypoxia mimetic deferoxamine influences the expression of histone acetylation- and DNA methylation-associated genes in osteoblasts. Connective Tissue Research, 2015, 56, 228-235.	2.3	18
8	Influence of trypsinization and alternative procedures for cell preparation before RNA extraction on RNA integrity. Analytical Biochemistry, 2014, 463, 38-44.	2.4	16
9	VEGF-A is associated with early degenerative changes in cartilage and subchondral bone. Growth Factors, 2018, 36, 263-273.	1.7	14
10	Healthy skeletal muscle aging: The role of satellite cells, somatic mutations and exercise. International Review of Cell and Molecular Biology, 2019, 346, 157-200.	3.2	10
11	Analysis of somatic mutations identifies signs of selection during in vitro aging of primary dermal fibroblasts. Aging Cell, 2019, 18, e13010.	6.7	6
12	SO042WHOLE GENOME SEQUENCING OF HUMAN KIDNEY PROGENITORS IDENTIFIES A MUTATION-PRONE CELL TYPE IN THE PROXIMAL TUBULE. Nephrology Dialysis Transplantation, 2020, 35, .	0.7	0