Manfred Kunz

List of Publications by Year in descending order

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MANEDED KUNZ

#	Article	IF	CITATIONS
1	A Transcriptome-Wide Isoform Landscape of Melanocytic Nevi and Primary Melanomas Identifies Gene Isoforms Associated with Malignancy. International Journal of Molecular Sciences, 2021, 22, 7165.	4.1	7
2	Single-cell trajectories of melanoma cell resistance to targeted treatment. Cancer Biology and Medicine, 2021, 18, 0-0.	3.0	6
3	Melanoma development: stage-dependent cancer competence of the melanocytic lineage. Signal Transduction and Targeted Therapy, 2021, 6, 433.	17.1	0
4	Nanoparticle-complexed antimiRs for inhibiting tumor growth and metastasis in prostate carcinoma and melanoma. Journal of Nanobiotechnology, 2020, 18, 173.	9.1	17
5	Mycobacterium marinum infection in an immunocompromised patient with infliximab. European Journal of Dermatology, 2020, 30, 436-437.	0.6	5
6	Psoriasis: Obesity and Fatty Acids. Frontiers in Immunology, 2019, 10, 1807.	4.8	52
7	Erythematous scaling lesions of the face, dorsal fingers, elbows, and knees together with symmetrical muscle weakness in a child. Clinical Case Reports (discontinued), 2019, 7, 1347-1349.	0.5	0
8	MEK inhibitor cobimetinib rescues a dR af mutant lethal phenotype in Drosophila melanogaster. Experimental Dermatology, 2019, 28, 1079-1082.	2.9	1
9	Modelling of Protein Kinase Signaling Pathways in Melanoma and Other Cancers. Cancers, 2019, 11, 465.	3.7	14
10	Mitogenâ€activated protein kinase pathway inhibitors rescue lethal phenotypes in a <scp>BRAF</scp> gainâ€ofâ€function <i>Drosophila melanogaster</i> model. Pigment Cell and Melanoma Research, 2018, 31, 545-548.	3.3	2
11	Nonhealing Crusted Scalp Lesions in a 4-Year-Old Boy. JAMA Dermatology, 2018, 154, 607.	4.1	0
12	Expression signatures of early-stage and advanced medaka melanomas. Comparative Biochemistry and Physiology Part - C: Toxicology and Pharmacology, 2018, 208, 20-28.	2.6	11
13	Pseudotime Dynamics in Melanoma Single-Cell Transcriptomes Reveals Different Mechanisms of Tumor Progression. Biology, 2018, 7, 23.	2.8	16
14	RNA-seq analysis identifies different transcriptomic types and developmental trajectories of primary melanomas. Oncogene, 2018, 37, 6136-6151.	5.9	91
15	Mapping heterogeneity in patient-derived melanoma cultures by single-cell RNA-seq. Oncotarget, 2017, 8, 846-862.	1.8	87
16	Extrakranielle kutane Manifestation einer Arteriitis temporalis - ein Fall erythematöser Plaques an den Unterschenkeln. JDDG - Journal of the German Society of Dermatology, 2016, 14, 66-71.	0.8	0
17	Tumor heterogeneity, clonality and single cells. Experimental Dermatology, 2016, 25, 857-858.	2.9	3
18	Extracranial cutaneous manifestation of temporal arteritis - a case of erythematous plaques on the legs. JDDG - Journal of the German Society of Dermatology, 2016, 14, 66-71.	0.8	1

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19	Genomeâ€wide association study identifies new susceptibility loci for cutaneous lupus erythematosus. Experimental Dermatology, 2015, 24, 510-515.	2.9	66
20	Reduced Adolescent-Age Spatial Learning Ability Associated with Elevated Juvenile-Age Superoxide Levels in Complex I Mouse Mutants. PLoS ONE, 2015, 10, e0123863.	2.5	8
21	Chk1 and Wee1 control genotoxic-stress induced G2–M arrest in melanoma cells. Cellular Signalling, 2015, 27, 951-960.	3.6	33
22	Polymorphisms in the mitochondrially encoded <scp>ATP</scp> synthase 8 gene are associated with susceptibility to bullous pemphigoid in the German population. Experimental Dermatology, 2015, 24, 715-717.	2.9	24
23	Genome-wide Association Analysis of Psoriatic Arthritis and Cutaneous Psoriasis Reveals Differences in Their Genetic Architecture. American Journal of Human Genetics, 2015, 97, 816-836.	6.2	245
24	miR-638 promotes melanoma metastasis and protects melanoma cells from apoptosis and autophagy. Oncotarget, 2015, 6, 2966-2980.	1.8	72
25	A mutation in the NADH-dehydrogenase subunit 2 suppresses fibroblast aging. Oncotarget, 2015, 6, 8552-8566.	1.8	12
26	The Genetic Basis of New Treatment Modalities in Melanoma. Current Drug Targets, 2015, 16, 233-248.	2.1	8
27	Genomewide <scp>RNA</scp> i screen identifies protein kinase <scp>C</scp> β and new members of mitogenâ€activated protein kinase pathway as regulators of melanoma cell growth and metastasis. Pigment Cell and Melanoma Research, 2014, 27, 418-430.	3.3	12
28	Oncogenes in melanoma: An update. European Journal of Cell Biology, 2014, 93, 1-10.	3.6	51
29	Lupus erythematosus. Part I: epidemiology, genetics and immunology. JDDG - Journal of the German Society of Dermatology, 2013, 11, 709-720.	0.8	8
30	Highâ€ŧhroughput sequencing of the melanoma genome. Experimental Dermatology, 2013, 22, 10-17.	2.9	33
31	MicroRNAs in Melanoma Biology. Advances in Experimental Medicine and Biology, 2013, 774, 103-120.	1.6	60
32	A Systems' Biology Approach to Study MicroRNA-Mediated Gene Regulatory Networks. BioMed Research International, 2013, 2013, 1-15.	1.9	32
33	New developments in dermatological oncogenetics. JDDG - Journal of the German Society of Dermatology, 2013, 11, 831-836.	0.8	1
34	Non-Major Histocompatibility Complex Rheumatoid Arthritis Susceptibility Genes. Critical Reviews in Immunology, 2011, 31, 99-114.	0.5	12
35	E2F1 in Melanoma Progression and Metastasis. Journal of the National Cancer Institute, 2010, 102, 127-133.	6.3	108
36	MicroRNA let-7b targets important cell cycle molecules in malignant melanoma cells and interferes with anchorage-independent growth. Cell Research, 2008, 18, 549-557.	12.0	425

#	Article	IF	CITATIONS
37	Gene Expression Signatures for Tumor Progression, Tumor Subtype, and Tumor Thickness in Laser-Microdissected Melanoma Tissues. Clinical Cancer Research, 2007, 13, 806-815.	7.0	205