Murat Iskar

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1726523/publications.pdf Version: 2024-02-01



MIIDAT ISKAD

#	Article	lF	CITATIONS
1	Cell typeâ€specific nuclear pores: a case in point for contextâ€dependent stoichiometry of molecular machines. Molecular Systems Biology, 2013, 9, 648.	7.2	277
2	The landscape of viral associations in human cancers. Nature Genetics, 2020, 52, 320-330.	21.4	261
3	IL4I1 Is a Metabolic Immune Checkpoint that Activates the AHR and Promotes Tumor Progression. Cell, 2020, 182, 1252-1270.e34.	28.9	259
4	Tumor-derived exosomes modulate PD-L1 expression in monocytes. Science Immunology, 2017, 2, .	11.9	236
5	Integrated Transcriptome and Proteome Analyses Reveal Organ-Specific Proteome Deterioration in Old Rats. Cell Systems, 2015, 1, 224-237.	6.2	176
6	Novel Drug Candidates for the Treatment of Metastatic Colorectal Cancer through Global Inverse Gene-Expression Profiling. Cancer Research, 2014, 74, 5690-5699.	0.9	142
7	Drug-Induced Regulation of Target Expression. PLoS Computational Biology, 2010, 6, e1000925.	3.2	120
8	Spatiotemporal variation of mammalian protein complex stoichiometries. Genome Biology, 2016, 17, 47.	8.8	115
9	Characterization of drugâ€induced transcriptional modules: towards drug repositioning and functional understanding. Molecular Systems Biology, 2013, 9, 662.	7.2	110
10	Identification and Analyses of Extra-Cranial and Cranial Rhabdoid Tumor Molecular Subgroups Reveal Tumors with Cytotoxic T Cell Infiltration. Cell Reports, 2019, 29, 2338-2354.e7.	6.4	74
11	Interleukin-10 receptor signaling promotes the maintenance of a PD-1int TCF-1+ CD8+ TÂcell population that sustains anti-tumor immunity. Immunity, 2021, 54, 2825-2841.e10.	14.3	57
12	The endosomal transcriptional regulator RNF11 integrates degradation and transport of EGFR. Journal of Cell Biology, 2016, 215, 543-558.	5.2	51
13	Molecular subgrouping of primary pineal parenchymal tumors reveals distinct subtypes correlated with clinical parameters and genetic alterations. Acta Neuropathologica, 2020, 139, 243-257.	7.7	50
14	Linking aberrant chromatin features in chronic lymphocytic leukemia to transcription factor networks. Molecular Systems Biology, 2019, 15, e8339.	7.2	39
15	DECIPHER pooled shRNA library screen identifies PP2A and FGFR signaling as potential therapeutic targets for diffuse intrinsic pontine gliomas. Neuro-Oncology, 2019, 21, 867-877.	1.2	24
16	CD8 ⁺ T-cells of CLL-bearing mice acquire a transcriptional program of T-cell activation and exhaustion. Leukemia and Lymphoma, 2020, 61, 351-356.	1.3	17
17	Pilocytic astrocytoma demethylation and transcriptional landscapes link bZIP transcription factors to immune response. Neuro-Oncology, 2020, 22, 1327-1338.	1.2	10
18	Discovery and validation of the antimetastatic activity of citalopram in colorectal cancer. Molecular and Cellular Oncology, 2015, 2, e975080.	0.7	6

Murat Iskar

#	Article	IF	CITATIONS
19	A synergistic interaction between HDAC―and PARP inhibitors in childhood tumors with chromothripsis. International Journal of Cancer, 2022, 151, 590-606.	5.1	5
20	CLL Exosome-Derived Y RNA hY4 Induces TLR7/8-Mediated Inflammation and PD-L1 Expression in Monocytes. Blood, 2016, 128, 3217-3217.	1.4	1
21	Immune Suppression in CLL Is Mediated By the L-Amino Acid Oxidase IL4I1, a Reason for the Treatment Failure of IDO1 Inhibitors. Blood, 2020, 136, 34-34.	1.4	0
22	Proteogenomic Subtyping of Chronic Lymphocytic Leukemia Identifies a Novel Poor Outcome Subgroup with a Distinct Drug Response Profile. Blood, 2020, 136, 10-11.	1.4	0