

# Murat Iskar

## List of Publications by Year in descending order

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Version: 2024-02-01

22  
papers

2,033  
citations

567281

15  
h-index

752698

20  
g-index

23  
all docs

23  
docs citations

23  
times ranked

4267  
citing authors

#	ARTICLE	IF	CITATIONS
1	A synergistic interaction between HDAC and PARP inhibitors in childhood tumors with chromothripsis. <i>International Journal of Cancer</i> , 2022, 151, 590-606.	5.1	5
2	Interleukin-10 receptor signaling promotes the maintenance of a PD-1 <sup>int</sup> TCF-1 <sup>+</sup> CD8 <sup>+</sup> T cell population that sustains anti-tumor immunity. <i>Immunity</i> , 2021, 54, 2825-2841.e10.	14.3	57
3	CD8 <sup>+</sup> T-cells of CLL-bearing mice acquire a transcriptional program of T-cell activation and exhaustion. <i>Leukemia and Lymphoma</i> , 2020, 61, 351-356.	1.3	17
4	Molecular subgrouping of primary pineal parenchymal tumors reveals distinct subtypes correlated with clinical parameters and genetic alterations. <i>Acta Neuropathologica</i> , 2020, 139, 243-257.	7.7	50
5	IL4I1 Is a Metabolic Immune Checkpoint that Activates the AHR and Promotes Tumor Progression. <i>Cell</i> , 2020, 182, 1252-1270.e34.	28.9	259
6	Pilocytic astrocytoma demethylation and transcriptional landscapes link bZIP transcription factors to immune response. <i>Neuro-Oncology</i> , 2020, 22, 1327-1338.	1.2	10
7	The landscape of viral associations in human cancers. <i>Nature Genetics</i> , 2020, 52, 320-330.	21.4	261
8	Immune Suppression in CLL Is Mediated By the L-Amino Acid Oxidase IL4I1, a Reason for the Treatment Failure of IDO1 Inhibitors. <i>Blood</i> , 2020, 136, 34-34.	1.4	0
9	Proteogenomic Subtyping of Chronic Lymphocytic Leukemia Identifies a Novel Poor Outcome Subgroup with a Distinct Drug Response Profile. <i>Blood</i> , 2020, 136, 10-11.	1.4	0
10	Identification and Analyses of Extra-Cranial and Cranial Rhabdoid Tumor Molecular Subgroups Reveal Tumors with Cytotoxic T Cell Infiltration. <i>Cell Reports</i> , 2019, 29, 2338-2354.e7.	6.4	74
11	Linking aberrant chromatin features in chronic lymphocytic leukemia to transcription factor networks. <i>Molecular Systems Biology</i> , 2019, 15, e8339.	7.2	39
12	DECIPHER pooled shRNA library screen identifies PP2A and FGFR signaling as potential therapeutic targets for diffuse intrinsic pontine gliomas. <i>Neuro-Oncology</i> , 2019, 21, 867-877.	1.2	24
13	Tumor-derived exosomes modulate PD-L1 expression in monocytes. <i>Science Immunology</i> , 2017, 2, .	11.9	236
14	The endosomal transcriptional regulator RNF11 integrates degradation and transport of EGFR. <i>Journal of Cell Biology</i> , 2016, 215, 543-558.	5.2	51
15	Spatiotemporal variation of mammalian protein complex stoichiometries. <i>Genome Biology</i> , 2016, 17, 47.	8.8	115
16	CLL Exosome-Derived Y RNA hY4 Induces TLR7/8-Mediated Inflammation and PD-L1 Expression in Monocytes. <i>Blood</i> , 2016, 128, 3217-3217.	1.4	1
17	Discovery and validation of the antimetastatic activity of citalopram in colorectal cancer. <i>Molecular and Cellular Oncology</i> , 2015, 2, e975080.	0.7	6
18	Integrated Transcriptome and Proteome Analyses Reveal Organ-Specific Proteome Deterioration in Old Rats. <i>Cell Systems</i> , 2015, 1, 224-237.	6.2	176

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19	Novel Drug Candidates for the Treatment of Metastatic Colorectal Cancer through Global Inverse Gene-Expression Profiling. <i>Cancer Research</i> , 2014, 74, 5690-5699.	0.9	142
20	Characterization of drug-induced transcriptional modules: towards drug repositioning and functional understanding. <i>Molecular Systems Biology</i> , 2013, 9, 662.	7.2	110
21	Cell type-specific nuclear pores: a case in point for context-dependent stoichiometry of molecular machines. <i>Molecular Systems Biology</i> , 2013, 9, 648.	7.2	277
22	Drug-Induced Regulation of Target Expression. <i>PLoS Computational Biology</i> , 2010, 6, e1000925.	3.2	120