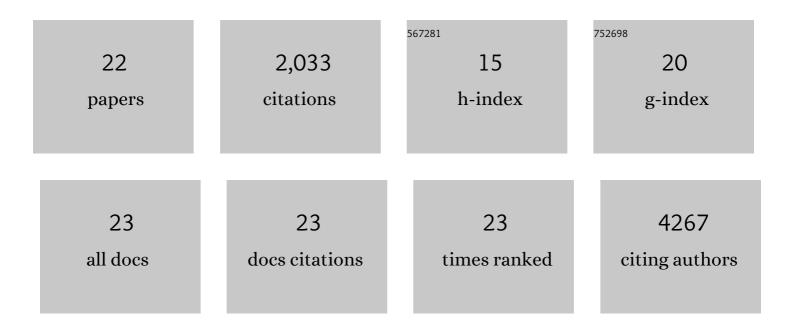
## Murat Iskar

List of Publications by Year in descending order

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



Μιίρλτ Ιςκλά

#	Article	IF	CITATIONS
1	A synergistic interaction between HDAC―and PARP inhibitors in childhood tumors with chromothripsis. International Journal of Cancer, 2022, 151, 590-606.	5.1	5
2	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
3	CD8 <sup>+</sup> 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
4	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
5	IL4I1 Is a Metabolic Immune Checkpoint that Activates the AHR and Promotes Tumor Progression. Cell, 2020, 182, 1252-1270.e34.	28.9	259
6	Pilocytic astrocytoma demethylation and transcriptional landscapes link bZIP transcription factors to immune response. Neuro-Oncology, 2020, 22, 1327-1338.	1.2	10
7	The landscape of viral associations in human cancers. Nature Genetics, 2020, 52, 320-330.	21.4	261
8	Immune Suppression in CLL Is Mediated By the L-Amino Acid Oxidase IL411, a Reason for the Treatment Failure of IDO1 Inhibitors. Blood, 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. Blood, 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. Cell Reports, 2019, 29, 2338-2354.e7.	6.4	74
11	Linking aberrant chromatin features in chronic lymphocytic leukemia to transcription factor networks. Molecular Systems Biology, 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. Neuro-Oncology, 2019, 21, 867-877.	1.2	24
13	Tumor-derived exosomes modulate PD-L1 expression in monocytes. Science Immunology, 2017, 2, .	11.9	236
14	The endosomal transcriptional regulator RNF11 integrates degradation and transport of EGFR. Journal of Cell Biology, 2016, 215, 543-558.	5.2	51
15	Spatiotemporal variation of mammalian protein complex stoichiometries. Genome Biology, 2016, 17, 47.	8.8	115
16	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
17	Discovery and validation of the antimetastatic activity of citalopram in colorectal cancer. Molecular and Cellular Oncology, 2015, 2, e975080.	0.7	6
18	Integrated Transcriptome and Proteome Analyses Reveal Organ-Specific Proteome Deterioration in Old Rats. Cell Systems, 2015, 1, 224-237.	6.2	176

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#	Article	IF	CITATIONS
19	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
20	Characterization of drugâ€induced transcriptional modules: towards drug repositioning and functional understanding. Molecular Systems Biology, 2013, 9, 662.	7.2	110
21	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
22	Drug-Induced Regulation of Target Expression. PLoS Computational Biology, 2010, 6, e1000925.	3.2	120