

# Gunes Gundem

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2428335/publications.pdf>

Version: 2024-02-01

18  
papers

3,288  
citations

687363

13  
h-index

752698

20  
g-index

21  
all docs

21  
docs citations

21  
times ranked

6208  
citing authors

#	ARTICLE	IF	CITATIONS
1	Bone Marrow Surveillance of Pediatric Cancer Survivors Identifies Clones that Predict Therapy-Related Leukemia. <i>Clinical Cancer Research</i> , 2022, 28, 1614-1627.	7.0	4
2	Feasibility of whole genome and transcriptome profiling in pediatric and young adult cancers. <i>Nature Communications</i> , 2022, 13, 2485.	12.8	31
3	Molecular International Prognostic Scoring System for Myelodysplastic Syndromes. , 2022, 1, .		259
4	Comprehensive Molecular Profiling of Desmoplastic Small Round Cell Tumor. <i>Molecular Cancer Research</i> , 2021, 19, 1146-1155.	3.4	14
5	Association of <i>BRAF V600E</i> mutations with vasoactive intestinal peptide syndrome in <i>MYCN</i> -amplified neuroblastoma. <i>Pediatric Blood and Cancer</i> , 2021, 68, e29265.	1.5	7
6	Prostate cancer evolution from multilineage primary to single lineage metastases with implications for liquid biopsy. <i>Nature Communications</i> , 2020, 11, 5070.	12.8	44
7	Accelerated single cell seeding in relapsed multiple myeloma. <i>Nature Communications</i> , 2020, 11, 3617.	12.8	41
8	Implications of TP53 allelic state for genome stability, clinical presentation and outcomes in myelodysplastic syndromes. <i>Nature Medicine</i> , 2020, 26, 1549-1556.	30.7	372
9	Cancer therapy shapes the fitness landscape of clonal hematopoiesis. <i>Nature Genetics</i> , 2020, 52, 1219-1226.	21.4	367
10	Revealing the Impact of Structural Variants in Multiple Myeloma. <i>Blood Cancer Discovery</i> , 2020, 1, 258-273.	5.0	81
11	Isabl Platform, a digital biobank for processing multimodal patient data. <i>BMC Bioinformatics</i> , 2020, 21, 549.	2.6	26
12	Patient-Driven Discovery, Therapeutic Targeting, and Post-Clinical Validation of a Novel <i>AKT1</i> Fusion-Driven Cancer. <i>Cancer Discovery</i> , 2019, 9, 605-616.	9.4	11
13	Abnormal oxidative metabolism in a quiet genomic background underlies clear cell papillary renal cell carcinoma. <i>ELife</i> , 2019, 8, .	6.0	31
14	Sequencing of prostate cancers identifies new cancer genes, routes of progression and drug targets. <i>Nature Genetics</i> , 2018, 50, 682-692.	21.4	182
15	A recurrent novel <i>MGA</i> - <i>NUTM1</i> fusion identifies a new subtype of high-grade spindle cell sarcoma. <i>Journal of Physical Education and Sports Management</i> , 2018, 4, a003194.	1.2	32
16	Mutational signatures of ionizing radiation in second malignancies. <i>Nature Communications</i> , 2016, 7, 12605.	12.8	214
17	Analysis of the genetic phylogeny of multifocal prostate cancer identifies multiple independent clonal expansions in neoplastic and morphologically normal prostate tissue. <i>Nature Genetics</i> , 2015, 47, 367-372.	21.4	380
18	The evolutionary history of lethal metastatic prostate cancer. <i>Nature</i> , 2015, 520, 353-357.	27.8	1,185