

# Hellen Kuasne

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12074640/publications.pdf>

Version: 2024-02-01

25  
papers

1,018  
citations

516710

16  
h-index

610901

24  
g-index

25  
all docs

25  
docs citations

25  
times ranked

1658  
citing authors

#	ARTICLE	IF	CITATIONS
1	Multi-omics data integration analysis identifies the spliceosome as a key regulator of DNA double-strand break repair. <i>NAR Cancer</i> , 2022, 4, zcac013.	3.1	5
2	Penile Cancer-Derived Cells Molecularly Characterized as Models to Guide Targeted Therapies. <i>Cells</i> , 2021, 10, 814.	4.1	9
3	Circulating mRNA signature as a marker for high-risk prostate cancer. <i>Carcinogenesis</i> , 2020, 41, 139-145.	2.8	12
4	CD44 Promotes PD-L1 Expression and Its Tumor-Intrinsic Function in Breast and Lung Cancers. <i>Cancer Research</i> , 2020, 80, 444-457.	0.9	88
5	GLUT1 inhibition blocks growth of RB1-positive triple negative breast cancer. <i>Nature Communications</i> , 2020, 11, 4205.	12.8	130
6	Epigenetic Switchâ€œInduced Viral Mimicry Evasion in Chemotherapy-Resistant Breast Cancer. <i>Cancer Discovery</i> , 2020, 10, 1312-1329.	9.4	84
7	DNA Methylation-Based Method to Differentiate Malignant from Benign Thyroid Lesions. <i>Thyroid</i> , 2019, 29, 1244-1254.	4.5	19
8	E-Cadherin Downregulation is Mediated by Promoter Methylation in Canine Prostate Cancer. <i>Frontiers in Genetics</i> , 2019, 10, 1242.	2.3	6
9	Deregulation of E-cadherin, $\beta$ -catenin, APC and Caveolin-1 expression occurs in canine prostate cancer and metastatic processes. <i>Research in Veterinary Science</i> , 2018, 118, 254-261.	1.9	20
10	Loss of DNA methylation is related to increased expression of miR-21 and miR-146b in papillary thyroid carcinoma. <i>Clinical Epigenetics</i> , 2018, 10, 144.	4.1	27
11	<sc>SHLD</sc> 2/ <sc>FAM</sc> 35A coâ€œoperates with <sc>REV</sc> 7 to coordinate <sc>DNA</sc> doubleâ€œstrand break repair pathway choice. <i>EMBO Journal</i> , 2018, 37, .	7.8	111
12	Nuclear loss and cytoplasmic expression of androgen receptor in penile carcinomas: role as a driver event and as a prognosis factor. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2018, 473, 607-614.	2.8	7
13	Integrated data analysis reveals potential drivers and pathways disrupted by DNA methylation in papillary thyroid carcinomas. <i>Clinical Epigenetics</i> , 2017, 9, 45.	4.1	68
14	Multidimensional integrative analysis uncovers driver candidates and biomarkers in penile carcinoma. <i>Scientific Reports</i> , 2017, 7, 6707.	3.3	35
15	Prognostic Classifier Based on Genome-Wide DNA Methylation Profiling in Well-Differentiated Thyroid Tumors. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2017, 102, 4089-4099.	3.6	45
16	Circulating mRNAs and miRNAs as candidate markers for the diagnosis and prognosis of prostate cancer. <i>PLoS ONE</i> , 2017, 12, e0184094.	2.5	95
17	Integrative miRNA and mRNA analysis in penile carcinomas reveals markers and pathways with potential clinical impact. <i>Oncotarget</i> , 2017, 8, 15294-15306.	1.8	39
18	A comprehensive characterization of cell cultures and xenografts derived from a human verrucous penile carcinoma. <i>Tumor Biology</i> , 2016, 37, 11375-11384.	1.8	16

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19	Genomic Profiling of Human Penile Carcinoma Predicts Worse Prognosis and Survival. <i>Cancer Prevention Research</i> , 2015, 8, 149-156.	1.5	52
20	Genome-wide methylation and transcriptome analysis in penile carcinoma: uncovering new molecular markers. <i>Clinical Epigenetics</i> , 2015, 7, 46.	4.1	48
21	Epigenetic Mechanisms in Penile Carcinoma. <i>International Journal of Molecular Sciences</i> , 2013, 14, 10791-10808.	4.1	17
22	Evaluation of the influence of polymorphic variants CYP1A1*2B, CYP1B1*2, CYP3A4*1B, GSTM1*0, and GSTT1*0 in prostate cancer. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2011, 29, 654-663.	1.6	28
23	Polymorphisms in the AR and PSA Genes as Markers of Susceptibility and Aggressiveness in Prostate Cancer. <i>Cancer Investigation</i> , 2010, 28, 917-924.	1.3	8
24	Association between polymorphisms in the biometabolism genes CYP1A1, GSTM1, GSTT1 and GSTP1 in bladder cancer. <i>Clinical and Experimental Medicine</i> , 2009, 9, 21-28.	3.6	47
25	Interplay Between Immune and Cancer-Associated Fibroblasts: A Path to Target Metalloproteinases in Penile Cancer. <i>Frontiers in Oncology</i> , 0, 12, .	2.8	2