

Enni Markkanen

List of Publications by Year in descending order

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Version: 2024-02-01

32
papers

1,585
citations

393982

19
h-index

433756

31
g-index

39
all docs

39
docs citations

39
times ranked

3080
citing authors

#	ARTICLE	IF	CITATIONS
1	Defining the molecular landscape of cancer-associated stroma in cutaneous squamous cell carcinoma. <i>Journal of Investigative Dermatology</i> , 2022, , .	0.3	3
2	Measuring DNA Damage Using the Alkaline Comet Assay in Cultured Cells. <i>Bio-protocol</i> , 2021, 11, e4119.	0.2	5
3	Cross-Reactivity and Functionality of Approved Human Immune Checkpoint Blockers in Dogs. <i>Cancers</i> , 2021, 13, 785.	1.7	15
4	Identification of disease-promoting stromal components by comparative proteomic and transcriptomic profiling of canine mammary tumors using laser-capture microdissected FFPE tissue. <i>Neoplasia</i> , 2021, 23, 400-412.	2.3	9
5	Measurement of DNA Damage Using the Neutral Comet Assay in Cultured Cells. <i>Bio-protocol</i> , 2021, 11, e4226.	0.2	0
6	Molecular homology between canine spontaneous oral squamous cell carcinomas and human head-and-neck squamous cell carcinomas reveals disease drivers and therapeutic vulnerabilities. <i>Neoplasia</i> , 2020, 22, 778-788.	2.3	10
7	Differential stromal reprogramming in benign and malignant naturally occurring canine mammary tumours identifies disease-modulating stromal components. <i>Scientific Reports</i> , 2020, 10, 5506.	1.6	20
8	Persistent DNA damage triggers activation of the integrated stress response to promote cell survival under nutrient restriction. <i>BMC Biology</i> , 2020, 18, 36.	1.7	24
9	Know Thy Model: Charting Molecular Homology in Stromal Reprogramming Between Canine and Human Mammary Tumors. <i>Frontiers in Cell and Developmental Biology</i> , 2019, 7, 348.	1.8	10
10	Next-generation RNA sequencing of FFPE subsections reveals highly conserved stromal reprogramming between canine and human mammary carcinoma. <i>DMM Disease Models and Mechanisms</i> , 2019, 12, .	1.2	20
11	MicroRNA Expression Profiling in the Prefrontal Cortex: Putative Mechanisms for the Cognitive Effects of Adolescent High Fat Feeding. <i>Scientific Reports</i> , 2018, 8, 8344.	1.6	14
12	Persistent DNA strand breaks induce a CAF-like phenotype in normal fibroblasts. <i>Oncotarget</i> , 2018, 9, 13666-13681.	0.8	20
13	Not breathing is not an option: How to deal with oxidative DNA damage. <i>DNA Repair</i> , 2017, 59, 82-105.	1.3	140
14	Impaired oxidative stress response characterizes HUWE1-promoted X-linked intellectual disability. <i>Scientific Reports</i> , 2017, 7, 15050.	1.6	21
15	Analysis of Gene Expression Signatures in Cancer-Associated Stroma from Canine Mammary Tumours Reveals Molecular Homology to Human Breast Carcinomas. <i>International Journal of Molecular Sciences</i> , 2017, 18, 1101.	1.8	35
16	An optimised protocol for isolation of RNA from small sections of laser-capture microdissected FFPE tissue amenable for next-generation sequencing. <i>BMC Molecular Biology</i> , 2017, 18, 22.	3.0	44
17	DNA Damage and Repair in Schizophrenia and Autism: Implications for Cancer Comorbidity and Beyond. <i>International Journal of Molecular Sciences</i> , 2016, 17, 856.	1.8	66
18	Targeting BRCA1 and BRCA2 Deficiencies with G-Quadruplex-Interacting Compounds. <i>Molecular Cell</i> , 2016, 61, 449-460.	4.5	185

#	ARTICLE	IF	CITATIONS
19	Abstract B40: WEE1 inhibition selectively kills histone H3K36me3-deficient cancers by dNTP starvation. , 2016, , .		0
20	Cells deficient in base-excision repair reveal cancer hallmarks originating from adjustments to genetic instability. <i>Nucleic Acids Research</i> , 2015, 43, 3667-3679.	6.5	39
21	Inhibiting WEE1 Selectively Kills Histone H3K36me3-Deficient Cancers by dNTP Starvation. <i>Cancer Cell</i> , 2015, 28, 557-568.	7.7	244
22	Gap-Directed Translesion DNA Synthesis of an Abasic Site on Circular DNA Templates by a Human Replication Complex. <i>PLoS ONE</i> , 2014, 9, e93908.	1.1	2
23	DNA polymerase ϵ -interacting protein 2 is a processivity factor for DNA polymerase δ during 8-oxo-7,8-dihydroguanine bypass. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 18850-18855.	3.3	44
24	MUTYH DNA glycosylase: the rationale for removing undamaged bases from the DNA. <i>Frontiers in Genetics</i> , 2013, 4, 18.	1.1	64
25	A switch between DNA polymerases ϵ and δ promotes error-free bypass of 8-oxo-G lesions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 20401-20406.	3.3	40
26	Base Excision Repair in Physiology and Pathology of the Central Nervous System. <i>International Journal of Molecular Sciences</i> , 2012, 13, 16172-16222.	1.8	22
27	Regulation of oxidative DNA damage repair: The adenine:8-oxo-guanine problem. <i>Cell Cycle</i> , 2012, 11, 1070-1075.	1.3	38
28	Regulation of oxidative DNA damage repair by DNA polymerase δ and MutYH by cross-talk of phosphorylation and ubiquitination. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 437-442.	3.3	67
29	Ubiquitylation of DNA polymerase δ . <i>FEBS Letters</i> , 2011, 585, 2826-2830.	1.3	16
30	In Vitro Gap-directed Translesion DNA Synthesis of an Abasic Site Involving Human DNA Polymerases μ , δ , and ϵ . <i>Journal of Biological Chemistry</i> , 2011, 286, 32094-32104.	1.6	27
31	Oxygen as a friend and enemy: How to combat the mutational potential of 8-oxo-guanine. <i>DNA Repair</i> , 2010, 9, 604-616.	1.3	272
32	The human checkpoint sensor and alternative DNA clamp Rad9/Rad1/Hus1 modulates the activity of DNA ligase I, a component of the long-patch base excision repair machinery. <i>Biochemical Journal</i> , 2005, 389, 13-17.	1.7	67