

Kang-Hoon Lee

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

277
citations

1039880

9
h-index

940416

16
g-index

18
all docs

18
docs citations

18
times ranked

447
citing authors

#	ARTICLE	IF	CITATIONS
1	The Role of Nitrosamine (NNK) in Breast Cancer Carcinogenesis. <i>Journal of Mammary Gland Biology and Neoplasia</i> , 2017, 22, 159-170.	1.0	35
2	Somatic Mutation of PIK3CA (H1047R) Is a Common Driver Mutation Hotspot in Canine Mammary Tumors as Well as Human Breast Cancers. <i>Cancers</i> , 2019, 11, 2006.	1.7	35
3	Methylation of LINE-1 in cell-free DNA serves as a liquid biopsy biomarker for human breast cancers and dog mammary tumors. <i>Scientific Reports</i> , 2019, 9, 175.	1.6	30
4	Transplantation of hMSCs Genome Edited with LEF1 Improves Cardio-Protective Effects in Myocardial Infarction. <i>Molecular Therapy - Nucleic Acids</i> , 2020, 19, 1186-1197.	2.3	25
5	Epigenetic Mechanisms of LncRNAs Binding to Protein in Carcinogenesis. <i>Cancers</i> , 2020, 12, 2925.	1.7	21
6	Upregulation of Complement Factor H by SOCS-1/STAT4 in Lung Cancer. <i>Cancers</i> , 2019, 11, 471.	1.7	20
7	Transcriptome Signatures of Canine Mammary Gland Tumors and Its Comparison to Human Breast Cancers. <i>Cancers</i> , 2018, 10, 317.	1.7	19
8	Genome-Wide Methylation Profiling in Canine Mammary Tumor Reveals miRNA Candidates Associated with Human Breast Cancer. <i>Cancers</i> , 2019, 11, 1466.	1.7	16
9	Alternative methylation of intron motifs is associated with cancer-related gene expression in both canine mammary tumor and human breast cancer. <i>Clinical Epigenetics</i> , 2020, 12, 110.	1.8	15
10	ANK2 Hypermethylation in Canine Mammary Tumors and Human Breast Cancer. <i>International Journal of Molecular Sciences</i> , 2020, 21, 8697.	1.8	11
11	Transposable element-mediated structural variation analysis in dog breeds using whole-genome sequencing. <i>Mammalian Genome</i> , 2019, 30, 289-300.	1.0	9
12	Long Non-Coding RNA Associated with Cholesterol Homeostasis and Its Involvement in Metabolic Diseases. <i>International Journal of Molecular Sciences</i> , 2020, 21, 8337.	1.8	9
13	Spatiotemporal expression of long noncoding RNA <i>Moshe</i> modulates heart cell lineage commitment. <i>RNA Biology</i> , 2021, 18, 640-654.	1.5	9
14	CXCL5 secreted from macrophages during cold exposure mediates white adipose tissue browning. <i>Journal of Lipid Research</i> , 2021, 62, 100117.	2.0	8
15	METTL8 mRNA Methyltransferase Enhances Cancer Cell Migration via Direct Binding to ARID1A. <i>International Journal of Molecular Sciences</i> , 2021, 22, 5432.	1.8	7
16	Analysis of opposing histone modifications H3K4me3 and H3K27me3 reveals candidate diagnostic biomarkers for TNBC and gene set prediction combination. <i>BMB Reports</i> , 2020, 53, 266-271.	1.1	5
17	Concise approach for screening long non-coding RNAs functionally linked to human breast cancer associated genes. <i>Experimental and Molecular Pathology</i> , 2019, 108, 89-96.	0.9	3