

# Kang-Hoon Lee

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3162334/publications.pdf>

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