## Daechan Park

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Molecular-level analysis of the serum antibody repertoire in young adults before and after seasonal influenza vaccination. Nature Medicine, 2016, 22, 1456-1464.	30.7	271
2	Large-scale sequence and structural comparisons of human naive and antigen-experienced antibody repertoires. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E2636-45.	7.1	179
3	Stem cell and neurogenic gene-expression profiles link prostate basal cells to aggressive prostate cancer. Nature Communications, 2016, 7, 10798.	12.8	166
4	Widespread Misinterpretable ChIP-seq Bias in Yeast. PLoS ONE, 2013, 8, e83506.	2.5	126
5	Simultaneous mapping of transcript ends at single-nucleotide resolution and identification of widespread promoter-associated non-coding RNA governed by TATA elements. Nucleic Acids Research, 2014, 42, 3736-3749.	14.5	93
6	Global Transcriptional Profiling Reveals Distinct Functions of Thymic Stromal Subsets and Age-Related Changes during Thymic Involution. Cell Reports, 2014, 9, 402-415.	6.4	87
7	RORα is crucial for attenuated inflammatory response to maintain intestinal homeostasis. Proceedings of the United States of America, 2019, 116, 21140-21149.	7.1	52
8	Determinants governing T cell receptor α/β-chain pairing in repertoire formation of identical twins. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 532-540.	7.1	42
9	Subtype-specific addiction of the activated B-cell subset of diffuse large B-cell lymphoma to FOXP1. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E577-E586.	7.1	36
10	The ATP-dependent chromatin remodeler Chd1 is recruited by transcription elongation factors and maintains H3K4me3/H3K36me3 domains at actively transcribed and spliced genes. Nucleic Acids Research, 2017, 45, 7180-7190.	14.5	35
11	The histone variant H2A.Z in yeast is almost exclusively incorporated into the +1 nucleosome in the direction of transcription. Nucleic Acids Research, 2020, 48, 157-170.	14.5	28
12	Poly(A)-specific ribonuclease sculpts the $3\hat{e}^2$ ends of microRNAs. Rna, 2019, 25, 388-405.	3.5	28
13	Chd1 co-localizes with early transcription elongation factors independently of H3K36 methylation and releases stalled RNA polymerase II at introns. Epigenetics and Chromatin, 2014, 7, 32.	3.9	25
14	CTCF cooperates with CtIP to drive homologous recombination repair of double-strand breaks. Nucleic Acids Research, 2019, 47, 9160-9179.	14.5	23
15	Dual-targeting of EGFR and Neuropilin-1 attenuates resistance to EGFR-targeted antibody therapy in KRAS-mutant non-small cell lung cancer. Cancer Letters, 2019, 466, 23-34.	7.2	23
16	iNrich, Rapid and Robust Method to Enrich N-Terminal Proteome in a Highly Multiplexed Platform. Analytical Chemistry, 2020, 92, 6462-6469.	6.5	18
17	Deep sequencing of B cell receptor repertoire. BMB Reports, 2019, 52, 540-547.	2.4	17
18	Beyond Autoantibodies: Biologic Roles of Human Autoreactive B Cells in Rheumatoid Arthritis Revealed by RNA‧equencing. Arthritis and Rheumatology, 2019, 71, 529-541.	5.6	17

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19	The chromatin-binding protein PHF6 functions as an E3 ubiquitin ligase of H2BK120 via H2BK12Ac recognition for activation of trophectodermal genes. Nucleic Acids Research, 2020, 48, 9037-9052.	14.5	15
20	Influenza vaccination in the elderly boosts antibodies against conserved viral proteins and egg-produced glycans. Journal of Clinical Investigation, 2021, 131, .	8.2	12
21	Genomic analysis of the telomeric length effect on organismic lifespan in Caenorhabditis elegans. Biochemical and Biophysical Research Communications, 2010, 396, 382-387.	2.1	8
22	Selection of self-priming molecular replicators. Nucleic Acids Research, 2019, 47, 2169-2176.	14.5	7
23	Diffusion-weighted MR imaging in pancreatic ductal adenocarcinoma: prediction of next-generation sequencing-based tumor cellularity and prognosis after surgical resection. Abdominal Radiology, 2021, 46, 4787-4799.	2.1	5
24	Telomeres reforged with non-telomeric sequences in mouse embryonic stem cells. Nature Communications, 2021, 12, 1097.	12.8	3
25	Deep RNA-Seq analysis reveals unexpected features of human prostate basal epithelial cells. Genomics Data, 2016, 7, 318-320.	1.3	Ο