

Rui Li

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7208054/publications.pdf>

Version: 2024-02-01

15
papers

1,584
citations

759233

12
h-index

940533

16
g-index

16
all docs

16
docs citations

16
times ranked

3514
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Enhancer connectome in primary human cells identifies target genes of disease-associated DNA elements. <i>Nature Genetics</i> , 2017, 49, 1602-1612. | 21.4 | 419 |
| 2 | Leukemia-Associated Cohesin Mutants Dominantly Enforce Stem Cell Programs and Impair Human Hematopoietic Progenitor Differentiation. <i>Cell Stem Cell</i> , 2015, 17, 675-688. | 11.1 | 177 |
| 3 | Cpr124 is essential for blood-brain barrier integrity in central nervous system disease. <i>Nature Medicine</i> , 2017, 23, 450-460. | 30.7 | 177 |
| 4 | Chromatin Accessibility Landscape of Cutaneous T Cell Lymphoma and Dynamic Response to HDAC Inhibitors. <i>Cancer Cell</i> , 2017, 32, 27-41.e4. | 16.8 | 136 |
| 5 | Assessment of the Genetic Basis of Rosacea by Genome-Wide Association Study. <i>Journal of Investigative Dermatology</i> , 2015, 135, 1548-1555. | 0.7 | 129 |
| 6 | Individuality and Variation of Personal Regulomes in Primary Human T Cells. <i>Cell Systems</i> , 2015, 1, 51-61. | 6.2 | 128 |
| 7 | Transcript-indexed ATAC-seq for precision immune profiling. <i>Nature Medicine</i> , 2018, 24, 580-590. | 30.7 | 124 |
| 8 | ecDNA hubs drive cooperative intermolecular oncogene expression. <i>Nature</i> , 2021, 600, 731-736. | 27.8 | 123 |
| 9 | TFAP2C- and p63-Dependent Networks Sequentially Rearrange Chromatin Landscapes to Drive Human Epidermal Lineage Commitment. <i>Cell Stem Cell</i> , 2019, 24, 271-284.e8. | 11.1 | 76 |
| 10 | Chromatin accessibility landscapes of skin cells in systemic sclerosis nominate dendritic cells in disease pathogenesis. <i>Nature Communications</i> , 2020, 11, 5843. | 12.8 | 22 |
| 11 | PIRCh-seq: functional classification of non-coding RNAs associated with distinct histone modifications. <i>Genome Biology</i> , 2019, 20, 292. | 8.8 | 20 |
| 12 | Chromatin Landscape Underpinning Human Dendritic Cell Heterogeneity. <i>Cell Reports</i> , 2020, 32, 108180. | 6.4 | 18 |
| 13 | Paired Transcriptomic and Proteomic Analysis Implicates IL-1 β in the Pathogenesis of Papulopustular Rosacea Explants. <i>Journal of Investigative Dermatology</i> , 2021, 141, 800-809. | 0.7 | 12 |
| 14 | Novel Gene Expression Profile of Women with Intrinsic Skin Youthfulness by Whole Transcriptome Sequencing. <i>PLoS ONE</i> , 2016, 11, e0165913. | 2.5 | 11 |
| 15 | Alterations of Immune and Keratinization Gene Expression in Papulopustular Rosacea by Whole Transcriptome Analysis. <i>Journal of Investigative Dermatology</i> , 2020, 140, 1100-1103.e4. | 0.7 | 10 |