Xiaoyu Liang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1339192/publications.pdf

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| | | 1478505 | 1720034 | |
|----------|----------------|--------------|----------------|--|
| 8 | 107 | 6 | 7 | |
| papers | citations | h-index | g-index | |
| | | | | |
| | | | | |
| 11 | 11 | 11 | 219 | |
| all docs | docs citations | times ranked | citing authors | |
| | | | | |

| # | Article | IF | CITATIONS |
|---|---|-----|-----------|
| 1 | A new monocyte epigenetic clock reveals nonlinear effects of alcohol consumption on biological aging in three independent cohorts ($\langle i \rangle N \langle i \rangle \hat{A} = \hat{A} 2242$). Alcoholism: Clinical and Experimental Research, 2022, 46, 736-748. | 2.4 | 9 |
| 2 | DNA methylation signature on phosphatidylethanol, not onÂself-reported alcohol consumption, predicts hazardous alcohol consumption in two distinct populations. Molecular Psychiatry, 2021, 26, 2238-2253. | 7.9 | 20 |
| 3 | i2d: an R package for simulating data from images and the implications in biomedical research. Bioinformatics, 2021, 37, 2497-2498. | 4.1 | О |
| 4 | Single-cell Transcriptome Mapping Identifies Common and Cell-type Specific Genes Affected by Acute Delta9-tetrahydrocannabinol in Humans. Scientific Reports, 2020, 10, 3450. | 3.3 | 17 |
| 5 | Gene selection for optimal prediction of cell position in tissues from single-cell transcriptomics data. Life Science Alliance, 2020, 3, e202000867. | 2.8 | 20 |
| 6 | A hierarchical clustering method for dimension reduction in joint analysis of multiple phenotypes. Genetic Epidemiology, 2018, 42, 344-353. | 1.3 | 13 |
| 7 | Joint analysis of multiple phenotypes in association studies using allele-based clustering approach for non-normal distributions. Annals of Human Genetics, 2018, 82, 389-395. | 0.8 | 1 |
| 8 | An Adaptive Fisher's Combination Method for Joint Analysis of Multiple Phenotypes in Association Studies. Scientific Reports, 2016, 6, 34323. | 3.3 | 25 |