

Anne-Valerie Gendrel

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9340698/publications.pdf>

Version: 2024-02-01

21
papers

3,652
citations

430874

18
h-index

713466

21
g-index

23
all docs

23
docs citations

23
times ranked

5509
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Role of transposable elements in heterochromatin and epigenetic control. <i>Nature</i> , 2004, 430, 471-476. | 27.8 | 1,103 |
| 2 | Profiling histone modification patterns in plants using genomic tiling microarrays. <i>Nature Methods</i> , 2005, 2, 213-218. | 19.0 | 521 |
| 3 | Dependence of Heterochromatic Histone H3 Methylation Patterns on the <i>Arabidopsis</i> Gene <i>DDM1</i> . <i>Science</i> , 2002, 297, 1871-1873. | 12.6 | 417 |
| 4 | SmcHD1, containing a structural-maintenance-of-chromosomes hinge domain, has a critical role in X inactivation. <i>Nature Genetics</i> , 2008, 40, 663-669. | 21.4 | 305 |
| 5 | Noncoding RNAs and Epigenetic Mechanisms During X-Chromosome Inactivation. <i>Annual Review of Cell and Developmental Biology</i> , 2014, 30, 561-580. | 9.4 | 195 |
| 6 | Smchd1-Dependent and -Independent Pathways Determine Developmental Dynamics of CpG Island Methylation on the Inactive X Chromosome. <i>Developmental Cell</i> , 2012, 23, 265-279. | 7.0 | 160 |
| 7 | Changes in 5S rDNA Chromatin Organization and Transcription during Heterochromatin Establishment in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2003, 15, 2929-2939. | 6.6 | 120 |
| 8 | Profiling DNA methylation patterns using genomic tiling microarrays. <i>Nature Methods</i> , 2005, 2, 219-224. | 19.0 | 119 |
| 9 | Developmental Dynamics and Disease Potential of Random Monoallelic Gene Expression. <i>Developmental Cell</i> , 2014, 28, 366-380. | 7.0 | 118 |
| 10 | Epigenetic Functions of Smchd1 Repress Gene Clusters on the Inactive X Chromosome and on Autosomes. <i>Molecular and Cellular Biology</i> , 2013, 33, 3150-3165. | 2.3 | 99 |
| 11 | Landscape of monoallelic DNA accessibility in mouse embryonic stem cells and neural progenitor cells. <i>Nature Genetics</i> , 2017, 49, 377-386. | 21.4 | 76 |
| 12 | MOF-associated complexes ensure stem cell identity and Xist repression. <i>ELife</i> , 2014, 3, e02024. | 6.0 | 76 |
| 13 | Fifty years of X-inactivation research. <i>Development (Cambridge)</i> , 2011, 138, 5049-5055. | 2.5 | 73 |
| 14 | Independent Mechanisms Target SMCHD1 to Trimethylated Histone H3 Lysine 9-Modified Chromatin and the Inactive X Chromosome. <i>Molecular and Cellular Biology</i> , 2015, 35, 4053-4068. | 2.3 | 66 |
| 15 | <i>Arabidopsis</i> epigenetics: when RNA meets chromatin. <i>Current Opinion in Plant Biology</i> , 2005, 8, 142-147. | 7.1 | 55 |
| 16 | Unusual chromatin status and organization of the inactive X chromosome in murine trophoblast giant cells. <i>Development (Cambridge)</i> , 2013, 140, 861-872. | 2.5 | 45 |
| 17 | Random monoallelic expression of genes on autosomes: Parallels with X-chromosome inactivation. <i>Seminars in Cell and Developmental Biology</i> , 2016, 56, 100-110. | 5.0 | 44 |
| 18 | The influence of DNA methylation on monoallelic expression. <i>Essays in Biochemistry</i> , 2019, 63, 663-676. | 4.7 | 39 |

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|----|--|------|-----------|
| 19 | The tandem repeat modules of <i>Xist</i> lncRNA: a swiss army knife for the control of X-chromosome inactivation. <i>Biochemical Society Transactions</i> , 2021, 49, 2549-2560. | 3.4 | 9 |
| 20 | Locus specific epigenetic modalities of random allelic expression imbalance. <i>Nature Communications</i> , 2021, 12, 5330. | 12.8 | 7 |
| 21 | X-Chromosome Inactivation and Autosomal Random Monoallelic Expression as "Faux Amis". <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 740937. | 3.7 | 1 |