

# François Le Dily

## List of Publications by Year in descending order

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Version: 2024-02-01

35  
papers

2,206  
citations

331670

21  
h-index

361022

35  
g-index

43  
all docs

43  
docs citations

43  
times ranked

3814  
citing authors

#	ARTICLE	IF	CITATIONS
1	Chromatin topology defines estradiol-primed progesterone receptor and PAX2 binding in endometrial cancer cells. <i>ELife</i> , 2022, 11, .	6.0	10
2	Coordinated changes in gene expression, H1 variant distribution and genome 3D conformation in response to H1 depletion. <i>Nucleic Acids Research</i> , 2022, 50, 3892-3910.	14.5	10
3	In vivo temporal resolution of acute promyelocytic leukemia progression reveals a role of <i>Klf4</i> in suppressing early leukemic transformation. <i>Genes and Development</i> , 2022, 36, 451-467.	5.9	1
4	OUP accepted manuscript. <i>Nucleic Acids Research</i> , 2021, 49, 11005-11021.	14.5	14
5	The impact of chromosomal fusions on 3D genome folding and recombination in the germ line. <i>Nature Communications</i> , 2021, 12, 2981.	12.8	34
6	A set of accessible enhancers enables the initial response of breast cancer cells to physiological progestin concentrations. <i>Nucleic Acids Research</i> , 2021, 49, 12716-12731.	14.5	13
7	CTCF is dispensable for immune cell transdifferentiation but facilitates an acute inflammatory response. <i>Nature Genetics</i> , 2020, 52, 655-661.	21.4	98
8	90 YEARS OF PROGESTERONE: Molecular mechanisms of progesterone receptor action on the breast cancer genome. <i>Journal of Molecular Endocrinology</i> , 2020, 65, T65-T79.	2.5	9
9	C/EBP $\beta$ mediates the growth inhibitory effect of progestins on breast cancer cells. <i>EMBO Journal</i> , 2019, 38, e101426.	7.8	15
10	Three-Dimensional Genomic Structure and Cohesin Occupancy Correlate with Transcriptional Activity during Spermatogenesis. <i>Cell Reports</i> , 2019, 28, 352-367.e9.	6.4	112
11	ATP, Mg <sup>2+</sup> , Nuclear Phase Separation, and Genome Accessibility. <i>Trends in Biochemical Sciences</i> , 2019, 44, 565-574.	7.5	37
12	Rapid reversible changes in compartments and local chromatin organization revealed by hyperosmotic shock. <i>Genome Research</i> , 2019, 29, 18-28.	5.5	40
13	Arginine Citrullination at the C-Terminal Domain Controls RNA Polymerase II Transcription. <i>Molecular Cell</i> , 2019, 73, 84-96.e7.	9.7	50
14	Hormone-control regions mediate steroid receptor-dependent genome organization. <i>Genome Research</i> , 2019, 29, 29-39.	5.5	49
15	OneD: increasing reproducibility of Hi-C samples with abnormal karyotypes. <i>Nucleic Acids Research</i> , 2018, 46, e49-e49.	14.5	50
16	Transcription factors orchestrate dynamic interplay between genome topology and gene regulation during cell reprogramming. <i>Nature Genetics</i> , 2018, 50, 238-249.	21.4	295
17	Unliganded Progesterone Receptor Governs Estrogen Receptor Gene Expression by Regulating DNA Methylation in Breast Cancer Cells. <i>Cancers</i> , 2018, 10, 371.	3.7	15
18	Promoter bivalency favors an open chromatin architecture in embryonic stem cells. <i>Nature Genetics</i> , 2018, 50, 1452-1462.	21.4	113

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19	Signaling by Steroid Hormones in the 3D Nuclear Space. <i>International Journal of Molecular Sciences</i> , 2018, 19, 306.	4.1	49
20	Lamin B1 mapping reveals the existence of dynamic and functional euchromatin lamin B1 domains. <i>Nature Communications</i> , 2018, 9, 3420.	12.8	66
21	Distinct roles of cohesin-SA1 and cohesin-SA2 in 3D chromosome organization. <i>Nature Structural and Molecular Biology</i> , 2018, 25, 496-504.	8.2	128
22	Parallel sequencing lives, or what makes large sequencing projects successful. <i>GigaScience</i> , 2017, 6, 1-6.	6.4	4
23	ADP-ribose-derived nuclear ATP synthesis by NUDIX5 is required for chromatin remodeling. <i>Science</i> , 2016, 352, 1221-1225.	12.6	141
24	TADs as modular and dynamic units for gene regulation by hormones. <i>FEBS Letters</i> , 2015, 589, 2885-2892.	2.8	20
25	On the demultiplexing of chromosome capture conformation data. <i>FEBS Letters</i> , 2015, 589, 3005-3013.	2.8	23
26	Distinct structural transitions of chromatin topological domains correlate with coordinated hormone-induced gene regulation. <i>Genes and Development</i> , 2014, 28, 2151-2162.	5.9	270
27	Nucleosome-Driven Transcription Factor Binding and Gene Regulation. <i>Molecular Cell</i> , 2013, 49, 67-79.	9.7	129
28	Unliganded progesterone receptor-mediated targeting of an RNA-containing repressive complex silences a subset of hormone-inducible genes. <i>Genes and Development</i> , 2013, 27, 1179-1197.	5.9	76
29	CDK2-dependent activation of PARP-1 is required for hormonal gene regulation in breast cancer cells. <i>Genes and Development</i> , 2012, 26, 1972-1983.	5.9	107
30	Differential Estrogen-Regulation of CXCL12 Chemokine Receptors, CXCR4 and CXCR7, Contributes to the Growth Effect of Estrogens in Breast Cancer Cells. <i>PLoS ONE</i> , 2011, 6, e20898.	2.5	91
31	Involvement of COUP-TFs in Cancer Progression. <i>Cancers</i> , 2011, 3, 700-715.	3.7	14
32	Nuclear Factor 1 Synergizes with Progesterone Receptor on the Mouse Mammary Tumor Virus Promoter Wrapped around a Histone H3/H4 Tetramer by Facilitating Access to the Central Hormone-responsive Elements. <i>Journal of Biological Chemistry</i> , 2010, 285, 2622-2631.	3.4	22
33	Two Chromatin Remodeling Activities Cooperate during Activation of Hormone Responsive Promoters. <i>PLoS Genetics</i> , 2009, 5, e1000567.	3.5	47
34	COUP-TFI modulates estrogen signaling and influences proliferation, survival and migration of breast cancer cells. <i>Breast Cancer Research and Treatment</i> , 2008, 110, 69-83.	2.5	30
35	Loss of E-cadherin-mediated cell contacts reduces estrogen receptor alpha (ER $\alpha$ ) transcriptional efficiency by affecting the respective contribution exerted by AF1 and AF2 transactivation functions. <i>Biochemical and Biophysical Research Communications</i> , 2008, 365, 304-309.	2.1	10