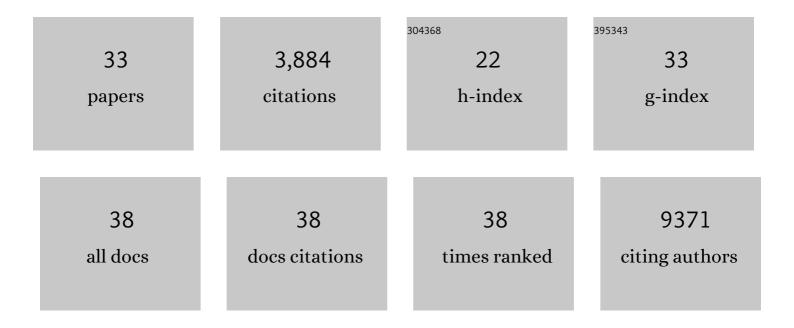
Kevin Lebrigand

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Persistent Properties of a Subpopulation of Cancer Cells Overexpressing the Hedgehog Receptor Patched. Pharmaceutics, 2022, 14, 988.	2.0	2
2	Evidence of early increased sialylation of airway mucins and defective mucociliary clearance in CFTR-deficient piglets. Journal of Cystic Fibrosis, 2021, 20, 173-182.	0.3	12
3	Single-cell RNA sequencing reveals intratumoral heterogeneity in primary uveal melanomas and identifies HES6 as a driver of the metastatic disease. Cell Death and Differentiation, 2021, 28, 1990-2000.	5.0	56
4	ldentification of oncolytic vaccinia restriction factors in canine high-grade mammary tumor cells using single-cell transcriptomics. PLoS Pathogens, 2020, 16, e1008660.	2.1	4
5	Agonist-induced functional analysis and cell sorting associated with single-cell transcriptomics characterizes cell subtypes in normal and pathological brain. Genome Research, 2020, 30, 1633-1642.	2.4	7
6	Transcriptomic and Ultrastructural Signatures of K+-Induced Aggregation in Phytophthora parasitica Zoospores. Microorganisms, 2020, 8, 1012.	1.6	7
7	High throughput error corrected Nanopore single cell transcriptome sequencing. Nature Communications, 2020, 11, 4025.	5.8	124
8	A Single-Cell Atlas of the Human Healthy Airways. American Journal of Respiratory and Critical Care Medicine, 2020, 202, 1636-1645.	2.5	282
9	Identification of a Repair-Supportive Mesenchymal Cell Population during Airway Epithelial Regeneration. Cell Reports, 2020, 33, 108549.	2.9	28
10	The nuclear hypoxia-regulated NLUCAT1 long non-coding RNA contributes to an aggressive phenotype in lung adenocarcinoma through regulation of oxidative stress. Oncogene, 2019, 38, 7146-7165.	2.6	75
11	CD4 ⁺ T Cells Affect the Thyroid Hormone Transport at the Choroid Plexus in Mice Raised in Enriched Environment. NeuroImmunoModulation, 2019, 26, 59-66.	0.9	2
12	The Long Noncoding RNA DNM3OS Is a Reservoir of FibromiRs with Major Functions in Lung Fibroblast Response to TGF-β and Pulmonary Fibrosis. American Journal of Respiratory and Critical Care Medicine, 2019, 200, 184-198.	2.5	78
13	HITS-CLIP in various brain areas reveals new targets and new modalities of RNA binding by fragile X mental retardation protein. Nucleic Acids Research, 2018, 46, 6344-6355.	6.5	124
14	Genome evolution across 1,011 Saccharomyces cerevisiae isolates. Nature, 2018, 556, 339-344.	13.7	952
15	CD8+ T cells are essential for the effects of enriched environment on hippocampus-dependent behavior, hippocampal neurogenesis and synaptic plasticity. Brain, Behavior, and Immunity, 2018, 69, 235-254.	2.0	44
16	Comparative Transcriptome Profiling of Virulent and Attenuated Ehrlichia ruminantium Strains Highlighted Strong Regulation of map1- and Metabolism Related Genes. Frontiers in Cellular and Infection Microbiology, 2018, 8, 153.	1.8	9
17	CD4+ T Cells Have a Permissive Effect on Enriched Environment-Induced Hippocampus Synaptic Plasticity. Frontiers in Synaptic Neuroscience, 2018, 10, 14.	1.3	11
18	Characterizing isomiR variants within the micro <scp>RNA</scp> â€34/449 family. FEBS Letters, 2017, 591, 693-705	1.3	48

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#	Article	IF	CITATIONS
19	A cost effective 5΄, selective single cell transcriptome profiling approach with improved UMI design. Nucleic Acids Research, 2017, 45, e48-e48.	6.5	27
20	Characterization of microRNAs from <i>Arabidopsis</i> galls highlights a role for miR159 in the plant response to the rootâ€knot nematode <i>Meloidogyne incognita</i> . New Phytologist, 2017, 216, 882-896.	3.5	71
21	Sequential activation and distinct functions for distal and proximal modules within the IgH 3′ regulatory region. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 1618-1623.	3.3	24
22	Comparative Genomic Analysis of Drechmeria coniospora Reveals Core and Specific Genetic Requirements for Fungal Endoparasitism of Nematodes. PLoS Genetics, 2016, 12, e1006017.	1.5	45
23	miR-193b/365a cluster controls progression of epidermal squamous cell carcinoma. Carcinogenesis, 2014, 35, 1110-1120.	1.3	66
24	Distinct epithelial gene expression phenotypes in childhood respiratory allergy. European Respiratory Journal, 2012, 39, 1197-1205.	3.1	64
25	A synonymous variant in IRGM alters a binding site for miR-196 and causes deregulation of IRGM-dependent xenophagy in Crohn's disease. Nature Genetics, 2011, 43, 242-245.	9.4	523
26	The human TTAGGG repeat factors 1 and 2 bind to a subset of interstitial telomeric sequences and satellite repeats. Cell Research, 2011, 21, 1028-1038.	5.7	123
27	Can the microRNA signature distinguish between thyroid tumors of uncertain malignant potential and other well-differentiated tumors of the thyroid gland?. Endocrine-Related Cancer, 2011, 18, 579-594.	1.6	31
28	Coxiella burnetii Transcriptional Analysis Reveals Serendipity Clusters of Regulation in Intracellular Bacteria. PLoS ONE, 2010, 5, e15321.	1.1	7
29	Identification of Keratinocyte Growth Factor as a Target of microRNA-155 in Lung Fibroblasts: Implication in Epithelial-Mesenchymal Interactions. PLoS ONE, 2009, 4, e6718.	1.1	192
30	Gene expression profiling of imatinib and PD166326-resistant CML cell lines identifies Fyn as a gene associated with resistance to BCR-ABL inhibitors. Molecular Cancer Therapeutics, 2009, 8, 1924-1933.	1.9	71
31	Global analysis of DNA methylation and transcription of human repetitive sequences Epigenetics, 2009, 4, 339-350.	1.3	28
32	Suppression of MicroRNA-Silencing Pathway by HIV-1 During Virus Replication. Science, 2007, 315, 1579-1582.	6.0	608
33	Transcriptional Signature of Epidermal Keratinocytes Subjected to in Vitro Scratch Wounding Reveals Selective Roles for ERK1/2, p38, and Phosphatidylinositol 3-Kinase Signaling Pathways. Journal of Biological Chemistry, 2007, 282, 15090, 15102	1.6	107