

Steve Bilodeau

List of Publications by Year in descending order

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Version: 2024-02-01

32
papers

5,294
citations

361413

20
h-index

377865

34
g-index

36
all docs

36
docs citations

36
times ranked

9894
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Subversion of infiltrating prostate macrophages to a mixed immunosuppressive tumor-associated macrophage phenotype. <i>Clinical and Translational Medicine</i> , 2022, 12, e581. | 4.0 | 9 |
| 2 | Cis-regulatory hubs: a new 3D model of complex disease genetics with an application to schizophrenia. <i>Life Science Alliance</i> , 2022, 5, e202101156. | 2.8 | 4 |
| 3 | The gut-liver axis: host microbiota interactions shape hepatocarcinogenesis. <i>Trends in Cancer</i> , 2022, 8, 583-597. | 7.4 | 22 |
| 4 | Control of adipogenic commitment by a STAT3-VSTM2A axis. <i>American Journal of Physiology - Endocrinology and Metabolism</i> , 2021, 320, E259-E269. | 3.5 | 8 |
| 5 | Proximity-dependent Mapping of the Androgen Receptor Identifies Kruppel-like Factor 4 as a Functional Partner. <i>Molecular and Cellular Proteomics</i> , 2021, 20, 100064. | 3.8 | 11 |
| 6 | Modulating HSF1 levels impacts expression of the estrogen receptor β and antiestrogen response. <i>Life Science Alliance</i> , 2021, 4, e202000811. | 2.8 | 7 |
| 7 | ZNF768 links oncogenic RAS to cellular senescence. <i>Nature Communications</i> , 2021, 12, 4841. | 12.8 | 11 |
| 8 | Defining the Transcriptional Ecosystem. <i>Molecular Cell</i> , 2018, 72, 920-924. | 9.7 | 18 |
| 9 | Connected Gene Communities Underlie Transcriptional Changes in Cornelia de Lange Syndrome. <i>Genetics</i> , 2017, 207, 139-151. | 2.9 | 23 |
| 10 | FOXA and master transcription factors recruit Mediator and Cohesin to the core transcriptional regulatory circuitry of cancer cells. <i>Scientific Reports</i> , 2016, 6, 34962. | 3.3 | 40 |
| 11 | metagene Profiles Analyses Reveal Regulatory Element's Factor-Specific Recruitment Patterns. <i>PLoS Computational Biology</i> , 2016, 12, e1004751. | 3.2 | 12 |
| 12 | Mutant cohesin affects RNA polymerase II regulation in Cornelia de Lange syndrome. <i>Scientific Reports</i> , 2015, 5, 16803. | 3.3 | 35 |
| 13 | ZFH4 Interacts with the NuRD Core Member CHD4 and Regulates the Glioblastoma Tumor-Initiating Cell State. <i>Cell Reports</i> , 2014, 6, 313-324. | 6.4 | 106 |
| 14 | Multiple Structural Maintenance of Chromosome Complexes at Transcriptional Regulatory Elements. <i>Stem Cell Reports</i> , 2013, 1, 371-378. | 4.8 | 113 |
| 15 | A Chromatin Switch for Chromosome Condensation. <i>Developmental Cell</i> , 2012, 23, 1127-1128. | 7.0 | 4 |
| 16 | X-linked H3K27me3 demethylase Utx is required for embryonic development in a sex-specific manner. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 13004-13009. | 7.1 | 179 |
| 17 | Enhancer decommissioning by LSD1 during embryonic stem cell differentiation. <i>Nature</i> , 2012, 482, 221-225. | 27.8 | 527 |
| 18 | Master Transcription Factors Determine Cell-Type-Specific Responses to TGF- β Signaling. <i>Cell</i> , 2011, 147, 565-576. | 28.9 | 536 |

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 19 | The histone methyltransferase SETDB1 is recurrently amplified in melanoma and accelerates its onset. <i>Nature</i> , 2011, 471, 513-517. | 27.8 | 506 |
| 20 | Mediator and cohesin connect gene expression and chromatin architecture. <i>Nature</i> , 2010, 467, 430-435. | 27.8 | 1,707 |
| 21 | Stem Cells, Differentiation and Cell Cycle Control in Pituitary. <i>Frontiers of Hormone Research</i> , 2010, 38, 15-24. | 1.0 | 20 |
| 22 | Cooperation between Cyclin E and p27Kip1 in Pituitary Tumorigenesis. <i>Molecular Endocrinology</i> , 2010, 24, 1835-1845. | 3.7 | 76 |
| 23 | SetDB1 contributes to repression of genes encoding developmental regulators and maintenance of ES cell state. <i>Genes and Development</i> , 2009, 23, 2484-2489. | 5.9 | 292 |
| 24 | Distinct Developmental Roles of Cell Cycle Inhibitors p57 ^{Kip2} and p27 ^{Kip1} Distinguish Pituitary Progenitor Cell Cycle Exit from Cell Cycle Reentry of Differentiated Cells. <i>Molecular and Cellular Biology</i> , 2009, 29, 1895-1908. | 2.3 | 113 |
| 25 | Wnt Signaling Promotes Reprogramming of Somatic Cells to Pluripotency. <i>Cell Stem Cell</i> , 2008, 3, 132-135. | 11.1 | 396 |
| 26 | Expression and mutation analysis of Tpit in the canine pituitary gland and corticotroph adenomas. <i>Domestic Animal Endocrinology</i> , 2008, 34, 217-222. | 1.6 | 8 |
| 27 | Of old and new diseases: genetics of pituitary ACTH excess (Cushing) and deficiency. <i>Clinical Genetics</i> , 2007, 72, 175-182. | 2.0 | 42 |
| 28 | Role of Brg1 and HDAC2 in GR <i>trans</i> -repression of the pituitary <i>POMC</i> gene and misexpression in Cushing disease. <i>Genes and Development</i> , 2006, 20, 2871-2886. | 5.9 | 213 |
| 29 | Rb Enhances p160/SRC Coactivator-dependent Activity of Nuclear Receptors and Hormone Responsiveness. <i>Journal of Biological Chemistry</i> , 2005, 280, 19746-19756. | 3.4 | 42 |
| 30 | Retinoblastoma and the Related Pocket Protein p107 Act as Coactivators of NeuroD1 to Enhance Gene Transcription. <i>Journal of Biological Chemistry</i> , 2005, 280, 16088-16095. | 3.4 | 32 |
| 31 | Protein-Protein Interactions and Transcriptional Antagonism between the Subfamily of NGFI-B/Nur77 Orphan Nuclear Receptors and Glucocorticoid Receptor. <i>Molecular Endocrinology</i> , 2005, 19, 885-897. | 3.7 | 106 |
| 32 | The T-box Factor Tpit Recruits SRC/p160 Co-activators and Mediates Hormone Action. <i>Journal of Biological Chemistry</i> , 2003, 278, 46523-46532. | 3.4 | 65 |