# Michael R Stallcup

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/7701243/publications.pdf

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|---|---|---|---|
| 35 papers | 4,593 citations | 249298 26 h-index | 406436 35 g-index |
| 36 all docs | 36 docs citations | 36 times ranked | 5060 citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | How Protein Methylation Regulates Steroid Receptor Function. Endocrine Reviews, 2022, 43, 160-197. | 8.9 | 13 |
| 2 | Gene-Specific Actions of Transcriptional Coregulators Facilitate Physiological Plasticity: Evidence for a Physiological Coregulator Code. Trends in Biochemical Sciences, 2020, 45, 497-510. | 3.7 | 35 |
| 3 | Relapse-associated AURKB blunts the glucocorticoid sensitivity of B cell acute lymphoblastic leukemia. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 3052-3061. | 3.3 | 32 |
| 4 | Increasing G9a automethylation sensitizes B acute lymphoblastic leukemia cells to glucocorticoid-induced death. Cell Death and Disease, 2018, 9, 1038. | 2.7 | 23 |
| 5 | Aberrant expression of SETD1A promotes survival and migration of estrogen receptor Î±â€positive breast cancer cells. International Journal of Cancer, 2018, 143, 2871-2883. | 2.3 | 32 |
| 6 | Different chromatin and DNA sequence characteristics define glucocorticoid receptor binding sites that are blocked or not blocked by coregulator Hic-5. PLoS ONE, 2018, 13, e0196965. | 1.1 | 8 |
| 7 | A postâ€translational modification switch controls coactivator function of histone methyltransferases G9a and GLP. EMBO Reports, 2017, 18, 1442-1459. | 2.0 | 32 |
| 8 | Glucocorticoid receptor binding to chromatin is selectively controlled by the coregulator Hic-5 and chromatin remodeling enzymes. Journal of Biological Chemistry, 2017, 292, 9320-9334. | 1.6 | 17 |
| 9 | Identifying differential transcription factor binding in ChIP-seq. Frontiers in Genetics, 2015, 6, 169. | 1.1 | 29 |
| 10 | Selective Coregulator Function and Restriction of Steroid Receptor Chromatin Occupancy by Hic-5. Molecular Endocrinology, 2015, 29, 716-729. | 3.7 | 19 |
| 11 | Distinct, Genome-Wide, Gene-Specific Selectivity Patterns of Four Glucocorticoid Receptor Coregulators. Nuclear Receptor Signaling, 2014, 12, nrs.12002. | 1.0 | 26 |
| 12 | Hic-5 is a transcription coregulator that acts before and/or after glucocorticoid receptor genome occupancy in a gene-selective manner. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 4007-4012. | 3.3 | 48 |
| 13 | Coregulator Cell Cycle and Apoptosis Regulator 1 (CCAR1) Positively Regulates Adipocyte Differentiation through the Glucocorticoid Signaling Pathway. Journal of Biological Chemistry, 2014, 289, 17078-17086. | 1.6 | 32 |
| 14 | Role of distinct surfaces of the G9a ankyrin repeat domain in histone and DNA methylation during embryonic stem cell self-renewal and differentiation. Epigenetics and Chromatin, 2014, 7, 27. | 1.8 | 17 |
| 15 | Gene-Specific Patterns of Coregulator Requirements by Estrogen Receptor-Î± in Breast Cancer Cells. Molecular Endocrinology, 2012, 26, 955-966. | 3.7 | 62 |
| 16 | G9a functions as a molecular scaffold for assembly of transcriptional coactivators on a subset of Glucocorticoid Receptor target genes. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 19673-19678. | 3.3 | 119 |
| 17 | Recruitment of coregulator G9a by Runx2 for selective enhancement or suppression of transcription. Journal of Cellular Biochemistry, 2012, 113, 2406-2414. | 1.2 | 43 |
| 18 | Recognition of enhancer elementâ€“specific histone methylation by TIP60 in transcriptional activation. Nature Structural and Molecular Biology, 2011, 18, 1358-1365. | 3.6 | 124 |

| # | Article | IF | Citations |
|---|---|---|---|
| 19 | A Distinct Mechanism for Coactivator versus Corepressor Function by Histone Methyltransferase G9a in Transcriptional Regulation. Journal of Biological Chemistry, 2011, 286, 41963-41971. | 1.6 | 74 |
| 20 | Regulated recruitment of tumor suppressor BRCA1 to the p21 gene by coactivator methylation. Genes and Development, 2011, 25, 176-188. | 2.7 | 60 |
| 21 | Minireview: Protein Arginine Methylation of Nonhistone Proteins in Transcriptional Regulation. Molecular Endocrinology, 2009, 23, 425-433. | 3.7 | 187 |
| 22 | The ankyrin repeats of G9a and GLP histone methyltransferases are mono- and dimethyllysine binding modules. Nature Structural and Molecular Biology, 2008, 15, 245-250. | 3.6 | 250 |
| 23 | CCAR1, a Key Regulator of Mediator Complex Recruitment to Nuclear Receptor Transcription Complexes. Molecular Cell, 2008, 31, 510-519. | 4.5 | 133 |
| 24 | Transcriptional Intermediary Factor 1Î± Mediates Physical Interaction and Functional Synergy between the Coactivator-Associated Arginine Methyltransferase 1 and Glucocorticoid Receptor-Interacting Protein 1 Nuclear Receptor Coactivators. Molecular Endocrinology, 2006, 20, 1276-1286. | 3.7 | 44 |
| 25 | Histone H3 Lysine 9 Methyltransferase G9a Is a Transcriptional Coactivator for Nuclear Receptors*. Journal of Biological Chemistry, 2006, 281, 8476-8485. | 1.6 | 168 |
| 26 | Role of Protein Methylation in Regulation of Transcription. Endocrine Reviews, 2005, 26, 147-170. | 8.9 | 392 |
| 27 | Regulation of coactivator complex assembly and function by protein arginine methylation and demethylimination. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 3611-3616. | 3.3 | 209 |
| 28 | Synergistic Effects of Coactivators GRIP1 and Î²-Catenin on Gene Activation. Journal of Biological Chemistry, 2004, 279, 4212-4220. | 1.6 | 64 |
| 29 | CoCoA, a Nuclear Receptor Coactivator which Acts through an N-Terminal Activation Domain of p160 Coactivators. Molecular Cell, 2003, 12, 1537-1549. | 4.5 | 116 |
| 30 | Methylation of Histone H3 by Coactivator-Associated Arginine Methyltransferase 1. Biochemistry, 2001, 40, 5747-5756. | 1.2 | 297 |
| 31 | Mouse Zac1, a Transcriptional Coactivator and Repressor for Nuclear Receptors. Molecular and Cellular Biology, 2000, 20, 1855-1867. | 1.1 | 144 |
| 32 | Interaction of the Ï„2 Transcriptional Activation Domain of Glucocorticoid Receptor with a Novel Steroid Receptor Coactivator, Hic-5, Which Localizes to Both Focal Adhesions and the Nuclear Matrix. Molecular Biology of the Cell, 2000, 11, 2007-2018. | 0.9 | 122 |
| 33 | Regulation of Transcription by a Protein Methyltransferase. Science, 1999, 284, 2174-2177. | 6.0 | 1,083 |
| 34 | Nuclear Receptor-Binding Sites of Coactivators Glucocorticoid Receptor Interacting Protein 1 (GRIP1) and Steroid Receptor Coactivator 1 (SRC-1): Multiple Motifs with Different Binding Specificities. Molecular Endocrinology, 1998, 12, 302-313. | 3.7 | 379 |
| 35 | Localization of the mouse glucocorticoid receptor-interacting protein 1 gene (Grip1) to proximal Chromosome 1 by linkage analysis. Mammalian Genome, 1997, 8, 620-621. | 1.0 | 1 |