Active Motif Epigenetic Services Publications

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Technique	Target	Journal	Year	Reference
ChIP-Seq and ChIP-qPCR	CBP, p300, H3K27Ac	Cell Reports	2019	Denise de Almeida Nagata *et al.* Regulation of Tumor-Associated Myeloid Cell Activity by CBP/EP300 Bromodomain Modulation of H3K27 Acetylation. *Cell Reports.* 27:269
ChIP-qPCR	ChREBP	Nature Metabolism	2019	Pauline Morigny *et al.* Interaction between hormone-sensitive lipase and ChREBP in fat cells controls insulin sensitivity. *Nature Metabolism.* 1: 133
ChIP-Seq	H3K4me1, H3K4me3, H3K27Ac	Cell Metabolism	2019	Liming Du et al. IGF-2 Preprograms Maturing Macrophages to Acquire Oxidative Phosphorylation-Dependent Antiinflammatory Properties. *Cell Metabolism.* 29: 1
ChIP-qPCR	SRF and CEBP	Nature Communications	2019	Shilpita Sarcar *et al.* Next-generation muscle-directed gene therapy by in silico vector design. *Nat Commun.* https://doi.org/10.1038/s41467-018-08283-7
ChIP-Seq	H3K27ac, H3K4me3, CTCF, and NR4A3	Nature Communications	2019	Florian Haller *et al.* Enhancer hijacking activates oncogenic transcription factor NR4A3 in acinic cell carcinomas of the salivary glands. *Nat Commun.* 10: 336
ChIP-Seq	Gfi1, Lsd1	Nature Communications	2019	Catherine Lee *et al.* Lsd1 as a therapeutic target in Gfi1-activated medulloblastoma. *Nat Commun.* 10: 332
ChIP-Seq	H3K27me3, H3K4me1, H3K4me3, H3K-9me3, H3K36me3, EZH2, SMARCB1, SUZ12,SMARCA4, REST	Cancer Cell	2019	Serap Erkek *et al.* Comprehensive Analysis of Chromatin States in Atypical Teratoid/Rhabdoid Tumors Identified Diverging Roles for SWI/SNF and Polycomb in Gene Regulation. *Cancer Cell.* 35: 95
ChIP-Seq	BRG1	The Journal of Clinical Investigation	2019	Yufeng Ding *et al.* Chromatin remodeling ATPase BRG1 and PTEN are synthetic lethal in prostate cancer. *J Clin Invest.* https://doi.org/10.1172/JCI123557
ChIP-Seq, RNA-Seq	GR	Journal of the American Heart Association	2019	Elena Severinova *et al.* Glucocorticoid Receptor Binding and Transcriptome Signature in Cardiomyocytes. *J Am Heart Assoc.* 8:e011484. DOI: 10. 1161/JAHA.118.011484.
ChIP-Seq	H4K20me1	Journal of Autoimmunity	2019	Keqi Fan *et al.* CRL4DCAF2is required for mature T-cell expansion via Aurora B-regulatedproteasome activity. *Journal of Autoimmunity.* 96:74
MeDIP-Seq	—	Epigenomes	2019	Liliana Ferreira *et al.* Uncovering Differentially Methylated Regions (DMRs) in a Salt-Tolerant Rice Variety under Stress: One Step towards New Regulatory Regions for Enhanced Salt Tolerance. *Epigenomes.* 3:4
ChIP-Seq	RNA Pol II, H3K27Ac	Journal of Neuro-Oncology	2019	Isabel Tegeder *et al.* Functional relevance of genes predicted to be affected by epigenetic alterations in atypical teratoid/rhabdoid tumors. *J. Neurooncol.* https://doi.org/10.1007/s11060-018-03018-6
ChIP-Seq	EGR1, H3K27Ac	Journal of Steroid Biochemsitry & Molecular Biology	2019	Maria Szwarc *et al.* Early growth response 1 transcriptionally primes the human endometrial stromal cell for decidualization. *J. Steroid Biochem. Mol. Biol.* https://doi.org/10.1016/j.jsbmb.2019.01.021

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ChIP-Seq	H3K27me3, H3K4me3	Acta Neuropathologica	2019	Andre Silveira *et al.* H3.3 K27M depletion increases diferentiation and extends latency of difuse intrinsic pontine glioma growth in vivo. *Acta Neuropathol*. https://doi.org/10.1007/s00401-019-01975-4
ChIP-Seq	RNA Pol II	Agronomy	2019	Sonja Klemme *et al.* Selection of Salicylic Acid Tolerant Epilines in *Brassica napus*. *Agronomy*. 9: 92
ChIP-Seq	LSD1, H3K4me2, H3K27Ac	Science Signaling	2019	Arnaud Augert *et al.* Targeting NOTCH activation in small cell lung cancer through LSD1 inhibition. *Sci.Signal*. 12, eaau2922
ChIP-qPCR	PPAR gamma	International Journal of Obesity	2019	Laura Butruille *et al.* Maternal high-fat diet during suckling programs visceral adiposity and epigenetic regulation of adipose tissue stearoyl-CoA desaturase-1 in offspring. *Int. J. Obes.* https://doi.org/10.1038/s41366-018-0310-z
ChIP-Seq	HA-tagged JP2NT, TBP, MEF2C	Science	2018	Ang Guo *et al.* E-C coupling structural protein junctophilin-2 encodes a stress-adaptive transcription regulator. *Science*. 362: 6421
ChIP-qPCR	p65/RelA	Human Molecular Genetics	2018	Jeffrey R. Gehlhausen *et al.* A proteasome-resistant fragment of NIK mediates oncogenic NF-B signaling in schwannomas. *Clin Cancer Res*. 25:1601. doi: 10.1093/hmg/ddy361
ChIP-Seq	Nrf2	Journal of Biological Chemsitry	2018	Junsheng Fu *et al.* Hyperactivity of the transcription factor Nrf2 causes metabolic reprogramming in mouse esophagus. *J. Biol. Chem.* 294: 327
ChIP-Seq	H3K36me2	Cell reports	2018	Jingjing Chen *et al.* Methyltransferase Nsd2 Ensures Germinal Center Selection by Promoting Adhesive Interactions between B Cells and Follicular Dendritic Cells. *Cell Reports*. 25: 3393
ChIP-Seq	H3K27Ac	Clinical Cancer Research	2018	Shori Saito *et al.* Eradication of central nervous system leukemia of T-cell origin with a brain-permeable LSD1 inhibitor. *Clin Cancer Res*. 25:1601
ChIP-Seq	RNA Pol II, Brd4, H3K9me3	Cell Reports	2018	Lu Want *et al.* Retinal Cell Type DNA Methylation and Histone Modifications Predict Reprogramming Efficiency and Retinogenesis in 3D Organoid Cultures. *Cell Reports*. 22: 2601
ATAC-Seq	NA	Cell Reports	2018	Xiaolong Zhang *et al.* OX40 Costimulation inhibits Foxp3 expresssion and Treg induction via BATF3-dependent and independent mechanisms. *Cell Reports*. 24: 607.
Mod Spec	NA	Acta Neuropathologica	2018	Leah Katz *et al.* Loss of histone H3K27me3 identifies a subset of meningiomas with increased risk of recurrence. *Acta Neuropathol*. 135: 6.
RIME	TRPS1	Cell Reports	2018	Robert M. Witwicki *et al.* TRPS1 is a lineage-specific transcriptional dependency in breast cancer. *Cell*. 175: 1.
ChIP-Seq	H3K27Ac	Cell	2018	Xun Huang *et al.* Targeting epigenetic crosstalk as a therapeutic strategy for EZH2-aberrant solid tumors. *Cell*. 175: 1.
ChIP-Seq	BRD4, RNA Pol II	Cancer Cell	2018	Elizabeth Stewart *et al.* Identification of therapeutic targets in rhabdomyosarcoma through integrated genomic, epigenomic, and proteomic analyses. *Cancer Cell*. 34: 1.
ChIP-Seq	Sox17	Nature Communications	2018	Xiaoqiu Wang *et al.* SOX17 regulates uterine epithelial-stromal cross-talk acting via a distal enhancer upstream of Ihh. *Nat Commun*. 9: 4421
ChIP-Seq, RNA-Seq	NRF2, RNA Pol II	Nature Communications	2018	David Olagnier *et al.* Nrf2 negatively regulates STING indicating a link between antiviral sensing and metabolic reprogramming. *Nat Commun*. 9:3506

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ChIP-Seq, RIME	p300, CBP, CDK9, BRD4, H3K27Ac, H3K4me1, H3K4me3, H3K18Ac	Cell Reports	2018	Ryan Raisner *et al.* Enhancer activity requires CBP/P300 bromodomain-dependent histone H3K27 acetylation. *Cell Reports.* 24: 1722.
ChIP-Seq, RNA-Seq	H3K27Ac, Med1, RNA Pol II	The Journal of Clinical Investigation	2018	Maria Florencia Martinez *et al.* Super-Enhancers maintain renin-expressing cell identity and memory to preserve multi-system homeostasis. *J. Clin Invest.* Advance Online Publication doi: 10.1172/JCI121361.
ChIP-Seq,	BRD4, H3K27Ac	JCI Insight	2018	Anisley Valenciaga *et al.* Transcriptional targeting of oncogene addiction in medullary thyroid cancer. *JCI Insight.* 3: e122225.
RNA-Seq	NA	PAIN	2018	Pradipta Ray *et al.* Comparative transcriptome profiling of the human and mouse dorsal root ganglia: an RNA-Seq based resource for pain and sensory neuroscience research. *Pain.* 159: 7.
ChIP-Seq	ELL2	The Journal Of Immunology	2018	Ashley M. Nelson *et al.* RNA splicing in the transition from B cells to antibody-secreting cells: The influences of ELL2, small nuclear RNA, and endoplasmic reticulum stress. *J Immunol.* Advance online publication doi: 10.4049/jimmunol
ChIP-Seq	MTA1	Cancer Medicine	2018	Nasir A. Butt *et al.* Targeting MTA1/HIF-1*a* signaling by pterostilbene in combination with histone deacetylase inhibitor attenuates prostate cancer progression. *Cancer Med.* 6: 2673
ChIP-Seq, Next-Gen Bisulfite-Seq	H3K4me3, RNA Pol II	Agronomy	2018	Martin Schmidt *et al.* Methylome and epialleles in rice epilines selected for energy use efficiency. *Agronomy.* 8: 163
ChIP-Seq	GATA2	The American Journal of Human Genetics	2018	Katelyn M. Mika *et al.* An ancientfecundability-associated polymorphism creates a GATA2 binding site in a distal enhancer of HLA-F. *Am J Hum Genet.* 103: 509
ChIP-Seq	FOXL2	Human Molecular Genetics	2018	Barbara Nicol *et al.* Genome-wide identifcation of FOXL2 binding and characterization of FOXL2 feminizing action in the fetal gonads. *Hum Mol Genet.* Advance online publication doi: 10.1093/hmg/ddy312
MeDIP-Seq	5-Methylcytosine	Scientific Reports	2018	Jose P. Silva *et al.* Analysis of diet-induced differential methylation, expression, and interactions of lncRNA and protein- coding genes in mouse liver. *Sci Rep.* 8: 11537
ChIP-Seq	ZFP24	Cell Reports	2018	Benayahu Elbaz *et al.* Phosphorylation state of ZFP24 controls oligodendrocyte differentiation. *Cell Reports.* 23: 2254
ChIP-Seq, Histone PTM Quantitation	H3K9me3, H3K27me3	Cell Reports	2018	Jessica Camacho *et al.* The memory of environmental chemical exposure in C. elegans is dependent on the jumonji demethylases jmjd-2 and jmjd-3/utx-1. *Cell Reports.* 23: 2392
ChIP-Seq	TET1, DNMT1	Nature Genetics	2018	Nipun Verma *et al.* TET proteins safeguard bivalent promoters from de novo methylation in human embryonic stem cells. *Nat Genet.* 50: 83
ChIP-Seq	EZH2	Journal of Experimental Medicine	2018	Xingli Zhang *et al.* Macrophage/microglial Ezh2 facilitates autoimmune inflammation through inhibition of Socs3. *J Exp Med.* 215: 1365
ChIP-Seq	STAT3,STAT5, ROCK2	Scientific Reports	2018	Wei Chen et al. ROCK2, but not ROCK1 interacts with phosphorylated STAT3 and co-occupies TH17/TFH gene promoters in TH17-activated human T cells. *Sci Rep.* 8: 16636
ChIP-Seq	Bhlhe40	PLoS One	2018	Kelly A. Hamilton *et al.* Mice lacking the transcriptional regulator Bhlhe40 have enhanced neuronal excitability and impaired synaptic plasticity in the hippocampus. *PLoS One.* 13: e0196223

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ChIP-Seq	RNA Pol II	PLoS One	2018	Liana Basova *et al.* Dopamine and its receptors play a role in the modulation of CCR5 expression in innate immune cells following exposure to Methamphetamine: Implications to HIV infection. *PLoS One.* 13: e0199861
ChIP-qPCR	Glucocorticoid Receptor, TIF1	Arthritis & Rheumatology	2018	Yanhua Hu *et al.* Development of a molecular signature to monitor pharmacodynamic response mediated by in vivo administration of glucocorticoids. *Arthritis Rheumatol.* 70: 1331
ChIP-Seq	H3K27me3	Acta Neuropathologica Communications	2018	David Castel *et al.* Transcriptomic and epigenetic profiling of 'diffuse midline gliomas, H3 K27M-mutant' discriminate two subgroups based on the type of histone H3 mutated and not supratentorial or infratentorial location. *Acta Neuropathol Commun.* 6: 17
ChIP-Seq	NRF2	Molecular Pharmacology	2018	Rance Nault *et al.* Comparison of hepatic NRF2 and AHR binding in 2,3,7,8- tetrachlorodibenzo-p-dioxin (TCDD) treated mice demonstrates NRF2- independent PKM2 induction. *Mol Pharmacol.* 94: 876
ChIP-Seq	HDAC2, H3K27Ac, H3K27me3	Cancer Immunology, Immunotherapy	2018	David Briere *et al.* The class I/IV HDAC inhibitor mocetinostat increases tumor antigen presentation, decreases immune suppressive cell types and augments checkpoint inhibitor therapy. *Cancer Immunol Immunother.* 67: 381
ChIP-Seq, ChIP-qPCR	TRRAP	Journal of Cell Biology	2018	Zhao Wang *et al.* TRRAP is a central regulator of human multiciliated cell formation. *J Cell Biol.* 217: 1941
ChIP-Seq	BRD4	Blood	2018	Rebecca Kohnken *et al.* Diminished microRNA-29b level is associated with BRD4 mediated activation of oncogenes in cutaneous T-cell lymphoma. *Blood.* 131: 771
RNA-Seq	NA	Cell Communication and Signaling	2018	Keri Callegari *et al.* Pharmacological inhibition of LSD1 activity blocks REST-dependent medulloblastoma cell migration. *Cell Commun Signal.* 16: 60
ChIP-Seq	MTA1	Molecular Oncology	2018	Avinash Kumar *et al.* MTA drives malignant progression and bone metatasis in prostate cancer
ChIP-Seq	RNA Pol II, H2A.Z, CDK9, ANP32e	Biochimica et Biophysica Acta	2018	Hyewon Shin *et al.* Transcriptional regulation mediated by H2A.Z via ANP32e-dependent inhibition of protein phosphatase 2A. *Biochim Biophys Acta.* 1861: 481
ChIP-Seq	H3K4me3, H3K9me3, H3K27me3	Proceedings of the Royal Society B	2018	Theresa K. Kelly *et al.* Epigenetic regulation of transcriptional plasticity associated with developmental song learning. Proc Biol Sci. 285: 20180160
ChIP-Seq	SIRT1	Journal of Neuroimmune Pharmacology	2018	Nikki Bortell *et al.* Sirtuin 1-chromatin-binding dynamics points to a common mechanism regulating inflammatory targets in SIV infection and in the aging brain. *J Neuroimmune Pharmacol.* 13: 163
ChIP-Seq	H3K27me3	G3 - Genes, Genomics Genetics	2018	James Ferguson *et al.* PRC2 is dispensable in vivo for b-catenin-mediated repression of chondrogenesis in the mouse embryonic cranial mesenchyme. *G3.* 8: 491
MeDIP-Seq	5-Methylcytosine	Nature	2017	Rama S. Akondy *et al.* Origin and differentiation of human memory CD8 T cells after vaccination. *Nature.* 552: 362
ChIP-Seq	10 different Histone mods plus CTCF, RNA Pol II, & BRD4	Nature	2017	Elizabeth Stewart *et al.* Orthotopic patient-derived xenografts of paediatric solid tumours. *Nature.* 549: 96

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ChIP-Seq	BACH2	Nature Communications	2017	Nicolas Hipp *et al.* IL-2 imprints human naive B cell fate towards plasma cell through ERK/ELK1-mediated BACH2 repression. *Nat Commun.* 8: 1443
ChIP-qPCR	H3K9Ac, RNA Pol II, ChREBP, PPARa	Cell Reports	2017	Alison Iroz *et al.* A specific ChREBP and PPARa cross-talk Is required for the glucose-mediated FGF21 response. *Cell Reports.* 21: 403
ChIP-Seq	H3K79me2	Molecular Cancer Therapy	2017	Carly T. Campbell *et al.* Mechanisms of Pinometostat (EPZ-5676) treatment–emergent resistance in MLL-rearranged leukemia. *Mol Cancer Ther.* 16: 1669
Next-Gen Bisulfite-Seq, ChIP-Seq	KLF4	Leukemia	2017	Y Shen *et al.* Inactivation of KLF4 promotes T-cell acute lymphoblastic leukemia and activates the MAP2K7 pathway. *Leukemia.* 31: 1314
ChIP-Seq	H3K27ac, H3K4me1	Epigenomics	2017	John P. Thomson *et al.* Defining baseline epigenetic landscapes in the rat liver. *Epigenomics.* 9: 1503
ChIP-Seq	H3K36me3	The Journal of Clinical Investigation	2017	Huairui Yuan *et al.* Histone methyltransferase SETD2 modulates alternative splicing to inhibit intestinal tumorigenesis. *J Clin Invest.* 127: 3375
RIME	Estrogen Receptor, Progesterone Receptor	Cancer Research	2017	Jessica Finlay-Schultz *et al.* Breast cancer suppression by progesterone receptors is mediated by their modulation of estrogen receptors and RNA polymerase III. *Cancer Res.* 77: 4934
ChIP-Seq	EZH2, H3K27me3, H3K4me3	Molecular Cancer Therapy	2017	Dorothy Brach *et al.* EZH2 inhibition by tazemetostat results in altered dependency on B-cell activation signaling in DLBCL. *Mol Cancer Ther.* 16:2586
ChIP-Seq Spike-in	p300, H3K27Ac, Androgen Receptor	Cancer Research	2017	Lingyan Jin *et al.* Therapeutic targeting of the CBP/p300 bromodomain blocks the growth of castration-resistant prostate cancer. *Cancer Res.* 77: 5564
ChIP-qPCR	IRF5, NFkB (p65)	Journal of Biological Chemistry	2017	Leah Cushing *et al.* IRAK4 kinase activity controls Toll-like receptor induced inflammation through the transcription factor IRF5 in primary human monocytes. *J Biol Chem.* 292: 18689
ChIP-Seq	RNA Pol II	Journal of Biological Chemistry	2017	Leah A. Gates *et al.* Acetylation on histone H3 lysine 9 mediates a switch from transcription initiation to elongation. *J Biol Chem.* 292: 14456
ChIP-Seq ChIP-qPCR	Estrogen Receptor	Endocrinology	2017	Sylvia C. Hewitt *et al.* Role of ERα in mediating female uterine transcriptional Responses to IGF1. *Endocrinology.* 158: 2427
ATAC-Seq ChIP-Seq	H3K4me3, H3K9me3	Cancer Cell	2017	Gulfem Dilek Guler *et al.* Repression of stress-induced LINE-1 expression protects cancer cell subpopulations from lethal drug exposure. *Cancer Cell.* 32: 1
ChIP-Seq	BRD4, H3K9me3, RNA Pol II	Neuron	2017	Issam Aldiri *et al.* The dynamic epigenetic landscape of the retina during development, reprogramming and tumorigenesis. *Neuron.* 94: 550
ChIP-Seq	Lmx1b	Development	2017	Endika Haro *et al.* Lmx1b-targeted cis-regulatory modules involved in limb dorsalization. *Dev.* 11: 2009
ChIP-Seq	EZH2	Proceedings of the Natl. Academy of Sciences	2017	Yongfeng Liu *et al.* Epithelial EZH2 serves as an epigenetic determinant in experimental colitis by inhibiting TNFα-mediated inflammation and apoptosis. *Proc Natl Acad Sci.* 114: E3796
ChIP-Seq	Estrogen Receptor	Endocrinology	2017	Laurel A. Coons *et al.* DNA sequence constraints define functionally active steroid nuclear receptor binding sites in chromatin. *Endocrinology.* 10: 3212

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ChIP-Seq	ETV5	Proceedings of the Natl. Academy of Sciences	2017	Zhen Zhang *et al.* Transcription factor Etv5 is essential for the maintenance of alveolar type II cells. *Proc Natl Acad Sci.* 114: E3903
ChIP-Seq	Androgen Receptor	Molecular and Cellular Endocrinology	2017	Claire Nash *et al.* Genome-wide analysis of AR binding and comparison with transcript expression in primary human fetal prostate fibroblasts and cancer associated fibroblasts. *Mol Cell Endocrinol.* 471: 1
ChIP-Seq Analysis	H3K27me3	Scientific Reports	2017	Brid O'Leary *et al.* Long non-coding RNA PARTICLE bridges histone and DNA methylation. *Sci Rep.* 7: 1790
ChIP-Seq	H3K36me2	The Journal of Clinical Investigation	2017	Ni Li *et al.* AKT-mediated stabilization of histone methyltransferase WHSC1 promotes prostate cancer metastasis. *J Clin Invest.* 127: 1284
ChIP-Seq ChIP-qPCR	EZH2, H3K27me3, H3K27ac	Cancer Cell	2017	Eric E. Gardner *et al.* Chemosensitive relapse in small cell lung cancer proceeds through an EZH2-SLFN11 axis. *Cancer Cell.* 31: 286
ChIP-Seq	TRIM28	Cell Reports	2017	Per Ludvik Brattas *et al.* TRIM28 controls a gene regulatory network based on endogenous retroviruses in human neural progenitor cells. *Cell Reports* 18: 1
ChIP-Seq	Cbx3/HP1	Scientific Reports	2017	Michael Sun *et al.* Cbx3/HP1γ deficiency confers enhanced tumor-killing capacity on CD8+ T cells. *Sci Rep.* 7: 42888
ChIP-Seq	H3K27Ac	Scientific Reports	2017	Yao Shen *et al.* Epigenetic and genetic dissections of UV-induced global gene dysregulation in skin cells through multi-omics analyses. *Sci Rep.* 7: 42646
ChIP-qPCR	SIRT1	Scientific Reports	2017	Jung-Yoon Yoo *et al.* KRAS activation and over- expression of SIRT1/BCL6 contributes to the pathogenesis of endometriosis and progesterone resistance. *Sci Rep.* 7: 6765
MeDIP-Seq	5-methylcytosine	Cerebral Cortex	2017	Daniela Grassi *et al.* Neuronal activity, TGF-Signaling and unpredictable chronic stress modulate transcription of Gadd45 family members and DNA methylation in the hippocampus. *Cereb Cortex.* 27: 4166
ChIP-Seq	SRC-2	PLoS Genetics	2017	Shruthy Suresh *et al.* SRC-2-mediated coactivation of anti-tumorigenic target genes supresses MYC-induced liver cancer. *PLoS Genet.* 13: e1006650
ChIP-Seq	SLY1	Cell Death and Differentiation	2017	Charlotte Moretti *et al.* SLY regulates genes involved in chromatin remodeling and interacts with TBL1XR1 during sperm differentiation. *Cell Death Differ.* 24: 1029
ChIP-Seq	G34R	Acta Neuropathologica Communications	2017	Farhana Haque *et al.* Evaluation of a novel antibody to define histone 3.3 G34R mutant brain tumours. *Acta Neruopathol Commun.* 5: 45
ChIP-Seq	SALL4	Journal of Hematology & Oncology	2017	Lina Yang *et al.* The stem cell factor SALL4 is an essential transcriptional regulator in mixed lineage leukemia-rearranged leukemogenesis. *J Hematol Oncol.* 10: 159
ChIP-qPCR	EZH2	Mechanisms of Ageing and Development	2017	Khyobeni Mozhui *et al.* Conserved effect of aging on DNA methylation and association with EZH2 polycomb protein in mice and humans. *Mech Ageing Dev.* 162: 27
ChIP-Seq Spike-in	H3K27me3, H3K4me3, H3K9me3	PLoS One	2016	Brian Egan *et al.* An alternative approach to ChIP-Seq normalization enables detection of genome- wide changes in histone H3 lysine 27 trimethylation upon EZH2 inhibition. *PLoS One.* 11: e0166438

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Low Cell ChIP-Seq	H3K27me3, CTCF	Nature Genetics	2016	Joachim Weischenfeldt *et al.* Pan-cancer analysis of somatic copy-number alterations implicates IRS4 and IGF2 in enhancer hijacking. *Nat Genet*. 49: 65
ChIP-Seq	OTX2, MITF, BRD4, H3K27Ac	Cancer Cell	2016	Pascal D. Johann *et al.* Atypical teratoid/rhabdoid tumors are comprised of three epigenetic subgroups with distinct enhancer landscapes. *Cancer Cell*. 29: 379
ChIP-Seq	GATA2, Progesterone Receptor	Cell Reports	2016	Cory A. Rubel *et al.* A Gata2-dependent transcription network regulates uterine progesterone responsiveness and endometrial function. *Cell Reports*. 17: 1414
ChIP-Seq	Androgen Receptor	Molecular Cell	2016	Boyu Zhang *et al.* Non-Cell-Autonomous Regulation of Prostate Epithelial Homeostasis by Androgen Receptor. *Mol Cell*. 63: 976
ChIP-Seq Spike-in	BRD4	The Journal of Clinical Investigation	2016	Mark L. McCleland *et al.* CCAT1 is an enhancer-templated RNA that predicts BET sensitivity in colorectal cancer. *J Clin Invest*. 126: 639
hMeDIP-Seq	5-Hydroxymethylcytosine	Nature Communications	2016	Carolina M. Greco *et al.* DNA hydroxymethylation controls cardiomyocyte gene expression in development and hypertrophy. *Nat Commun*. 7: 12418
ChIP-qPCR	Androgen Receptor	Cell Reports	2016	Karyn Schmidt *et al.* The lncRNA SLNCR1 mediates melanoma invasion through a conserved SRA1-like region. *Cell Rep*. 15: 2025
Low Cell ChIP-Seq	H3K4me3, H3K27Ac, H3K27me3	Biology of Reproduction	2016	Kazadi Mutoji *et al.* TSPAN8 expression distinguishes spermatogonial stem cells in the prepubertal mouse testis. *Biol Reprod*. 95: 17
ChIP-Seq	HDAC1, H3K27Ac	Cancer Discovery	2016	Anjali Mishra *et al.* Mechanism, consequences, and therapeutic targeting of abnormal IL15 signaling in cutaneous T-cell lymphoma. *Cancer Discov*. 6: 986
ChIP-Seq	V5-tagged DUX4	Human Molecular Genetics	2016	Jocelyn O. Eidahl *et al.* Mouse Dux is myotoxic and shares partial functional homology with its human paralog DUX4. *Hum Mol Genet*. 25: 4577
ChIP-Seq	H3K27me3	Molecular and Cellular Biology	2016	Kyung Hyun Yoo *et al.* Histone demethylase KDM6A controls the mammary luminal lineage through enzyme-independent mechanisms. *Mol Cell Biol.* 36: 2108
ChIP-Seq	H3K27Ac	Nature Genetics	2016	Pratiti Bandopadhayay *et al.* MYB-QKI rearrangements in angiocentric glioma drive tumorigenicity through a tripartite mechanism. *Nat Genet*. 48: 273
ChIP-Seq Spike-in	KDM5C	Cell Reports	2016	Shigeki Iwase *et al.* A mouse model of X-linked intellectual disability associated with impaired removal of histone methylation. *Cell Reports*. 14: 1
ChIP-Seq ChIP-qPCR	Wiz	eLIFE	2016	Luke Isbel *et al.* Wiz binds active promoters and CTCF-binding sites and is required for normal behaviour in the mouse. *Elife*. 5: e15082
ChIP-Seq	BRD4, H3K27Ac, H3K4me1, H3K27me3, HLX, LHX2, LMX1A	Nature	2016	Charles Y. Lin *et al.* Active medulloblastoma enhancers reveal subgroup specific cellular origins. *Nature*. 530: 57
ChIP-qPCR	H3K9me3 H3 pan-acetyl	Nucleic Acids Research	2016	Xiaoyu Chen *et al.* Probing the impact of chromatin conformation on genome editing tools. *Nucleic Acids Res*. 44: 6482
ChIP-Seq ChIP-qPCR	H3K9Ac	Stem Cells Translational Medicine	2016	Dalia Ali *et al.* Epigenetic library screen identifies abexinostat as novel regulator of adipocytic and osteoblastic differentiation of human skeletal (mesenchymal) stem cells. *Stem Cells Transl Med*. 5: 1036

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ChIP-Seq	H3K27me3	Cancer Research	2016	John P. Thomson *et. al.* Loss of Tet1 associated 5-hydroxymethylcytosine is concomitant with aberrant promoter hypermethylation in liver cancer. *Cancer Res.* 76: 3097
ChIP-Seq, ChIP-qPCR	H3K36me3	Oncogene	2016	Thai H. Ho *et al.* High-resolution profiling of histone h3 lysine 36 trimethylation in metastatic renal cell carcinoma. *Oncogene.* 35: 1565
ChIP-Seq	HDAC1, HDAC2, GATA2	PLoS One	2016	Jeffrey R. Shearstone *et al.* Chemical Inhibition of histone deacetylases 1 and 2 Induces fetal hemoglobin through activation of GATA2. *PLoS One.* 11: e0153767
ChIP-Seq	LL-37 (antimicrobial peptide)	Journal of Cancer	2016	Mindy Munoz *et al.* Antimicrobial peptide LL-37 participates in the transcriptional regulation of melanoma cells. *J Cancer.* 26: 2341
ChIP-Seq ChIP-qPCR	p53	Molecular Oncology	2016	Cheryl Chan *et al.* Global re-wiring of p53 transcription regulation by the hepatitis B virus X protein. *Mol Oncol.* 10: 1183
ChIP-Seq	PLZF, SALL4	Development	2016	Dawn L. Lovelace *et al.* The regulatory repertoire of PLZF and SALL4 in undifferentiated spermatogonia. *Development.* 143: 1893
ChIP-Seq	PLZF	PLoS Genetics	2016	Ramakrishna Kommagani *et al.* The Promyelocytic leukemia zinc finger transcription factor is critical for human endometrial stromal cell decidualization. *PLoS Genet.* 12: e1005937
ChIP libraries, sequencing and analysis	H3K27me3	Developmental Dynamics	2016	Oyvind Dahle *et al.* Inhibiting smad2/3 signaling in pluripotent mouse embryonic stem cells enhances endoderm formation by increasing transcriptional priming of lineage-specifying target genes. *Dev Dyn.* 245: 807.
ChIP-Seq	NR4A1	PLoS One	2016	Ryan P. Duren *et al.* Genome wide mapping of NR4A binding reveals cooperativity with ETS factors to promote epigenetic activation of distal enhancers in acute myeloid leukemia cells. *PLoS One.* 11: e0150450
ChIP-Seq	MTA1	Oncotarget	2016	Swati Dhar *et al.* Dietary pterostilbene is a novel MTA1-targeted chemopreventive and therapeutic agent in prostate cancer. *Oncotarget.* 7: 18469
ChIP-Seq	Myc-tagged Pet-1	The Journal of Neuroscience	2016	Steven C. Wyler *et al.* Pet-1 switches transcriptional targets postnatally to regulate maturation of serotonin neuron excitability. *J Neurosci.* 36:1758
ChIP-Seq	EGR2	Nature Communications	2015	Tomohisa Okamura *et al.* TGF-b3-expressing $CD4^{+}CD25^{-}LAG3^{+}$ regulatory T cells control humoral immune responses. *Nat Commun.* 6: 6329
ChIP-Seq	HA-tagged Shox2	Development	2015	Wenduo Ye *et al.* A common Shox-2-Nkx2-5 antagonistic mechanism primes the pacemaker cell fate in the pulmonary vein myocardium and sinoatrial node. *Development.* 142: 2521
ChIP-Seq	BA180/PBRM1	Molecular Cell	2015	Bokai Zhu *et al.* Coactivator-dependent oscillation of chromatin accessibility dictates circadian gene amplitude via REV-ERB loading. *Mol Cell.* 60: 769
ChIP-qPCR	EZH2, SUZ12, H3K-27me3, RNA Pol II	The EMBO Journal	2015	Stephen G. Dann *et al.* Reciprocal regulation of amino acid import and epigenetic state through Lat1 and EZH2. *EMBO J.* 34: 1773
ChIP-Seq	CCND1 (Cyclin D1)	Oncotarget	2015	Mathew C. Casimiro *et al.* Kinase-independent role of cyclin D1 in chromosomal instability and mammary tumorigenesis. *Oncotarget.* 6: 8524
ChIP-Seq	TRIM33	PLoS Genetics	2015	Luke Isbel *et al.* Trim33 binds and silences a class of young endogenous retroviruses in the mouse testis; a novel component of the arms race between retrotransposons and the host genome. *PLoS Genetics.* 11: e1005693

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ChIP-Seq	Progesterone Receptor	Cell Reports	2015	Vincent J. Lynch *et al.* Ancient transposable elements transformed the uterine regulatory landscape and transcriptome during the evolution of mammalian pregnancy. *Cell Rep.* 10: 551
ChIP-Seq	EGR1	PLoS One	2015	Anthony A. Portale *et al.* Characterization of FGF23-dependent Egr-1 cistrome in the mouse renal proximal tubule. *PLoS One.* 10: e0142924
ChIP-Seq	FOXM1	Breast Cancer Research and Treatment	2015	Christina Yau *et al.* FOXM1 cistrome predicts breast cancer metastatic outcome better than FOXM1 expression levels or tumor proliferation index. *Breast Cancer Res Treat.* 154: 23
ChIP-Seq	BRD4	Molecular Cancer Therapy	2015	Ryan Lenhart *et al.* Sensitivity of small cell lung cancer to BET inhibition is mediated by regulation of ASCL1 gene expression. *Mol Cancer Ther.* 14: 2167
Next-Gen Bisulfite Seq	NA	The Journal of Clinical Investigation	2015	Coralie Hoareau-Aveilla *et al.* Reversal of microRNA-150 silencing disadvantages crizotinib-resistant NPM-ALK(+) cell growth. *J Clin Invest.* 125:3505
ChIP-Seq ChIP-qPCR	β–Arrestin-1	Cancer Research	2015	Smitha Pillai *et al.* β-arrestin-1 mediates nicotine-induced metastasis through E2F1 target genes that modulate epithelial-mesenchymal transition. *Cancer Res.* 75: 1009
ChIP-Seq	H3K9ac, RNA Pol II, TFIIB	Circulation: Heart Failure	2015	Danish Sayed *et al.* Acute targeting of general transcription factor IIB restricts cardiac hypertrophy via selective inhibition of gene transcription. *Circ Heart Fail.* 8: 138
ChIP-Seq	H3K4me3	Journal of the National Cancer Institute	2015	Cristian Taccioli *et al.* Repression of esophageal neoplasia and inflammatory signaling by anti-miR-31 delivery in vivo. *J Natl Cancer Inst.* 107: djv220
ChIP-Seq	FLAG-tagged Twist2	American Journal of Human Genetics	2015	Shannon Marchegiani *et al.* Recurrent mutations in the basic domain of TWIST2 cause ablepharon macrostomia and barber-say syndromes. *Am J Hum Genet.* 97: 99
ChIP-Seq	LHX6	Human Molecular Genetics	2015	Jeffry M. Cesario *et al.* Lhx6 and Lhx8 promote palate development through negative regulation of a cell cycle inhibitor gene, $p57^{Kip2}$. *Hum Mol Genet.* 24: 5024
ChIP-Seq	SOX9	Nucleic Acids Research	2015	Zhongcheng Shi *et al.* Context-specific role of SOX9 in NF-Y mediated gene regulation in colorectal cancer cells. *Nucleic Acids Res.* 43: 6257
ChIP-Seq	H3K27me3	Nucleic Acids Research	2015	Kyung Hyun Yoo *et al.* Loss of EZH2 results in precocious mammary gland development and activation of STAT5-dependent genes. *Nucleic Acids Res.* 43: 8774
ChIP-Seq	H3K4me1	BMC Biology	2015	Sara K. Harten *et al.* The recently identified modifier of murine metastable epialleles, Rearranged L-Myc Fusion, is involved in maintaining epigenetic marks at CpG island shores and enhancers. *BMC Biol.* 13: 21
ChIP-Seq	ZNF384	Molecular Endocrinology	2015	Paul Childress *et al.* Genome-wide mapping and interrogation of the Nmp4 anti-anabolic bone axis. *Mol Endocrinol.* 29: 1269
ChIP-Seq	FOXO1, RNA Pol II	Molecular Endocrinology	2015	Yasmin M. Vasquez *et al.* FOXO1 is Required for Binding of PR on IRF4, Novel Transcriptional Regulator of Endometrial Stromal Decidualization. *Mol Endocrinol.* 29: 421
ChIP-Seq	Progesterone Receptor, FOSL2	Endocrinology	2015	Erik C. Mazur *et al.* Progesterone receptor transcriptome and cistrome in decidualized human endometrial stromal cells. *Endocrinology.* 156: 2239
ChIP-Seq	H3K27me3,H3K4me1, H3K4me3, H3K27Ac, p300, RARα	Immunity	2015	Chrysothemis C. Brown *et al.* Retinoic acid is essential for Th1 cell lineage stability and prevents transition to a Th17 cell program. *Immunity.* 42: 1

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ChIP-Seq	SIX1	Cancer Cell	2015	Jenny Wegert *et al.* Mutations in the SIX1/2 pathway and the DROSHA/DGCR8 miRNA microprocessor complex underlie high-risk blastemal type Wilms tumors. *Cancer Cell.* 27: 298
ChIP-Seq	H3K4me3, H3K9me3	Epigenetics	2015	Nioka C. Chisholm *et al.* Histone methylation patterns in astrocytes are influenced by age following ischemia. *Epigenetics.* 10: 142
ChIP-Seq Data Analysis	KDM3A	PLoS Biology	2015	Mo-bin Cheng *et al.* Specific phosphorylation of histone demethylase KDM3A determines target gene expression in response to heat shock. *PLOS Biol.* 12: e1002026
ChIP-Seq	SRC1	Molecular Endocrinology	2014	Mounia Tannour-Louet *et al.* Hepatic SRC-1 activity orchestrates transcriptional circuitries of amino acid pathways with potential relevance for human metabolic pathogenesis. *Mol Endocrinol.* 28: 1707
ChIP-Seq	Fish H3K4me3	Molecular Ecology Resources	2014	Claudius F. Kratochwil *et al.* Mapping active promoters by ChIP-seq profiling of H3K4me3 in cichlid fish – a first step to uncover cis-regulatory elements in ecological model teleosts. *Mol Ecol Resour.* 15: 761
ChIP-Seq	GATA2, FOXA1, p300, CBP, SRC1, SRC2, SRC3, Androgen Receptor	Proceedings of the National Acadamy of Sciences	2014	Bin He *et al.* GATA2 facilitates steroid receptor coactivator recruitment to the androgen receptor complex. *Proc Natl Acad Sci.* 111: 18261
Next-Gen Bisulfite Seq	NA	Cell Reports	2014	Deepak Kumar *et al.* Fibroblast growth factor maintains chondrogenic potential of limb bud mesenchymal cells by modulating DNMT3A recruitment. *Cell Rep.* 8: 1419
ChIP-Seq	PPARg	Biology of Reproduction	2014	Kelsey E. Brooks *et al.* Peroxisome proliferator activator receptor gamma (PPARG) regulates conceptus elongation in sheep. *Biol Reprod.* 92: 42.
ChIP-Seq, LightSwitch™	PXR, p300, H3K4me1, H3K27ac	PLoS Genetics	2014	Robin P. Smith *et al.* Genome-wide discovery of drug-dependent human liver regulatory elements. *PLOS Genet.* 10: e1004648
ChIP-Seq	H3K9me3, H3K36me3, H3K27me3	Genome Research	2014	Goran Kungulovski *et al.* Application of histone modification-specific interaction domains as an alternative to antibodies. *Genome Res.* 24: 1842
ChIP-Seq ChIP-qPCR	SRF	Physiological Genomics	2014	Sharolyn V. Kawakami-Schulz. Serum response factor: positive and negative regulation of an epithelial gene expression network in the destrin mutant cornea. *Physiol Genomics.* 46: 277
ChIP-Seq ChIP-qPCR	SRC3	The Journal of Clinical Investigation	2014	Jun Qin *et al.* Androgen deprivation–induced NCoA2 promotes metastatic and castration-resistant prostate cancer. *J Clin Invest.* 124: 5013
ChIP-Seq	H3K27Ac	Journal of Biological Chemistry	2014	Andre Landin Malt *et al.* Identification of a face enhancer reveals direct regulation of LIM homeobox 8 (Lhx8) by wingless-int (WNT)/β-catenin Signaling. *J Biol Chem.* 289: 30289
ChIP-Seq ChIP-qPCR	ZBTB20	Journal of Medical Genetics	2014	Malene B. Rasmussen *et al.* Neurodevelopmental disorders associated with dosage imbalance of ZBTB20 correlate with the morbidity spectrum of ZBTB20 candidate target genes. *J Med Genet.* 51: 605
ChIP-Seq	H3K9Ac, H3K27Ac	Nature	2014	Paul A. Northcott *et al.* Enhancer hijacking activates GFI1 family oncogenes in medulloblastoma. *Nature.* 511: 428
ChIP-Seq	H3K4me3, H3K27me3	Nature	2014	Volker Hovestadt *et al.* Decoding the regulatory landscape of medulloblastoma using DNA methylation sequencing. *Nature.* 510: 537
ChIP-qPCR	EZH2, Suz12, H3K-27me3	Molecular Cancer Therapy	2014	Sarah K. Knutson *et al.* Selective Inhibition of EZH2 by EPZ-6438 Leads to Potent Antitumor Activity in EZH2 Mutant Non-Hodgkin Lymphoma. *Mol Cancer Ther.* 13: 842

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ChIP-Seq	RAR-α, RAR-β, RXR-α	American Journal of Physiology – Gastrointestinal and Liver Physiology	2014	Yuqi He *et al.* Biological functional annotation of retinoic acid alpha and beta in mouse liver based on genome-wide binding. *Am J Physiol Gastrointest Liver Physiol.* 307: G205
ChIP-Seq	H3K4me3, H3K27me3	Nature Communications	2014	Brian C. Belyea *et al.* Identification of renin progenitors in the mouse bone marrow that give rise to B-cell leukaemia. *Nat Commun.* 5: 3273
ChIP-Seq	Estrogen Receptor	Molecular Endocrinology	2014	Sylvia C. Hewitt *et al.* Novel DNA motif binding activity observed In vivo with an estrogen receptor a mutant mouse. *Mol Endocrinol.* 26: 899
ChIP-Seq	H3K27me3	Bioinformatics	2014	Yanxiao Zhang *et al.* PePr: A peak-calling prioritization pipeline to identify consistent or differential peaks from replicated ChIP-Seq data. *Bioinformatics.* 30: 2568
ChIP-Seq	H3K36me3	Genome Research	2014	Jeremy M. Simon *et al.* Variation in chromatin accessibility in human kidney cancer links H3K36 methyltransferase loss with widespread RNA processing defects. *Genome Res.* 24: 241
ChIP-Seq	SRC-2	Cell Reports	2014	Erin Stashi *et al.* SRC-2 Is an Essential Coactivator for Orchestrating Metabolism and Circadian Rhythm. *Cell Rep.* 6: 663
Next-Gen Bisulfite Seq	—	Nature Neuroscience	2014	Brian G. Dias *et al.* Parental olfactory experience influences behavior and neural structure in subsequent generations. *Nat Neurosci.* 17: 89
ChIP-Seq ChIP-qPCR	Zbtb20	Cerebral Cortex	2014	Jakob V. Nielsen *et al.* Zbtb20 Defines a Hippocampal Neuronal Identity Through Direct Repression of Genes That Control Projection Neuron Development in the Isocortex. *Cereb Cortex.* 24: 1216.
ChIP-qPCR	H3K9me3, H3	Nature Immunology	2013	Patrick M. Gubser *et al.* Rapid effector function of memory CD8+ T cells requires an immediate-early glycolytic switch. *Nat Immunol.* 14: 1064
ChIP-Seq	RNA Pol II	PLoS One	2013	Jonathan P. Riley *et al.* PARP-14 Binds Specific DNA Sequences to Promote Th2 Cell Gene Expression. *PLoS One.* 8: e83127
Next-Gen Bisulfite Seq	—	Clinical Cancer Research	2013	David S. Shames *et al.* Loss of NAPRT1 Expression by Tumor-Specific Promoter Methylation Provides a Novel Predictive Biomarker for NAMPT Inhibitors. *Clin Cancer Res.* 19: 6912
ChIP-Seq	Cyclin D1	Cancer Research	2013	Xiaoming Ju *et al.* Identification of a cyclin D1 network in prostate cancer that antagonizes epithelial-mesenchymal restraint. *Cancer Res.* 74: 508
ChIP-Seq	H3K27me3	Cancer Cell	2013	Sebastian Bender *et al.* Reduced H3K27me3 and DNA Hypomethylation Are Major Drivers of Gene Expression in K27M Mutant Pediatric High-Grade Gliomas. *Cancer Cell.* 24: 660
ChIP-Seq	ASXL1	Journal of Experimental Medicine	2013	Omar Abdel-Wahab *et al.* Deletion of Asxl1 results in myelodysplasia and severe developmental defects in vivo. *J Exp Med.* 210: 2641
ChIP-Seq	COUP-TFII (NR2F2)	Molecular Endocrinology	2013	Xilong Li *et al.* COUP-TFII Regulates Human Endometrial Stromal Genes Involved in Inflammation. *Mol Endocrinol.* 27: 2041
ChIP-Seq	Gfi1	Nature Immunology	2013	Chauncey J. Spooner *et al.* Specification of type 2 innate lymphocytes by the transcriptional determinant Gfi1. *Nat Immunol.* 14: 1229
ChIP-Seq	RNA Pol II	Journal of Allergy and Clinical Immunology	2013	Purvi Mehrotra *et al.* PARP-14 and its enzyme activity regulates Th2 differentiation and allergic airway disease. *J Allergy Clin Immunol.* 131: 52
ChIP-Seq	Progesterone Receptor	Molecular Endocrinology	2013	Ashlee R. Lain *et al.* Research Resource: Progesterone Receptor Targetome Underlying Mammary Gland Branching Morphogenesis. *Mol Endocrinol.* 27: 1743

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ChIP-Seq	PPARα	Chemico-Biological Interactions	2013	Patrick D. McMullen *et al.* A map of the PPARα transcription regulatory network for primary human hepatocytes. *Chem Biol Interact.* 209: 104
ChIP-Seq	H3K36me3	Cancer Discovery	2013	Lynn Bjerke *et al.* Histone H3.3 Mutations Drive Pediatric Glioblastoma through Upregulation of MYCN. *Cancer Discov.* 3: 512
ChIP-Seq	p53	Physiological Genomics	2013	Yuwen Li *et al.* Genome-wide analysis of the p53 gene regulatory network in the developing mouse kidney. *Physiol Genomics.* 45: 948
ChIP-Seq	FOXA2	The FASEB Journal	2013	Justyna Filant *et al.* Integrated chromatin immunoprecipitation sequencing and microarray analysis identifies FOXA2 target genes in the glands of the mouse uterus. *FASEB J.* 28: 230
ChIP-Seq	p300	PLoS Genetics	2013	Aaron M. Wenger *et al.* The Enhancer Landscape during Early Neocortical Development Reveals Patterns of Dense Regulation and Co-option. *PLoS Genet.* 9: e1003728
ChIP-Seq	MYRF	PLoS Biology	2013	Helena Bujalka *et al.* MYRF is a membrane-associated transcription factor that autoproteolytically cleaves to directly activate myelin genes. *PLoS Biol.* 11: e1001625
ChIP-Seq ChIP-qPCR	RXR-α, RNA Pol II	PLoS One	2013	Astrid Kosters *et al.* Sexually Dimorphic Genome-Wide Binding of Retinoid X Receptor alpha (RXRα) Determines Male-Female Differences in the Expression of Hepatic Lipid Processing Genes in Mice. *PLoS One.* 8: e71538
ChIP-qPCR	NFκB (p52 & p65)	Journal of Biological Chemistry	2013	Sarah L. Doyle *et al.* Nuclear factor κB2 p52 protein has a role in antiviral immunity through IκB kinase epsilon-dependent induction of Sp1 protein and interleukin 15. *J Biol Chem.* 288: 25066
ChIP-chip ChIP-qPCR	C/EBPβ, RNA Pol II	Journal of Biological Chemistry	2013	Hana Vakili *et al.* CCAAT-enhancer-binding protein β (C/EBPβ) and downstream human placental growth hormone genes are targets for dysregulation in pregnancies complicated by maternal obesity. *J Biol Chem.* 288: 22849
ChIP-Seq	KDM2B	The Journal of Clinical Investigation	2013	Alexandros Tzatsos *et al.* KDM2B promotes pancreatic cancer via Polycomb-dependent and -independent transcriptional programs. *J Clin Invest.* 123: 727
ChIP-Seq	Vitamin D Receptor	BMC Medicine	2013	Adam E. Handel *et al.* Vitamin D receptor ChIP-seq in primary CD4+ cells: relationship to serum 25-hydroxyvitamin D levels and autoimmune disease. *BMC Med.* 11: 163
ChIP-Seq	COUP-TFII (NR2F2)	Developmental Cell	2013	San-pin Wu *et al.* Atrial Identity Is Determined by a *COUP-TFII* Regulatory Network. *Dev Cell.* 25: 417
ChIP-Seq	H3K4me3	PLoS One	2013	Alex Gutteridge *et al.* Novel Pancreatic Endocrine Maturation Pathways Identified by Genomic Profiling and Causal Reasoning. *PLoS One.* 8: e56024.
ChIP-Seq	Glucocorticoid Receptor	PLoS One	2012	Marie-José C. van Lierop *et al.* Org 214007-0: A Novel Non-Steroidal Selective Glucocorticoid Receptor Modulator with Full Anti- Inflammatory Properties and Improved Therapeutic Index. *PLoS One.* 7: e48385
ChIP-Seq	KDM1A (LSD1), H3K4me2	Molecular and Cellular Biology	2012	Venugopalan D. Nair *et al.* Involvement of Histone Demethylase LSD1 in Short-Time-Scale Gene Expression Changes during Cell Cycle Progression in Embryonic Stem Cells. *Mol Cell Biol.* 32: 4861
ChIP-Seq	EZH2	Nature	2012	Michael T. McCabe *et al.* EZH2 inhibition as a therapeutic strategy for lymphoma with EZH2-activating mutations. *Nature.* 492: 108

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Data Analysis	pTyr37 H2B	Nature Structural and Molecular Biology	2012	Kiran Mahajan *et al.* H2B Tyr37 phosphorylation suppresses expression of replication-dependent core histone genes. *Nat Struct Mol Biol.* 19: 930
ChIP-chip	MYBL2 (B-MYB)	PLoS One	2012	Ming Zhan *et al.* The B-MYB Transcriptional Network Guides Cell Cycle Progression and Fate Decisions to Sustain Self-Renewal and the Identity of Pluripotent Stem Cells. *PLoS One.* 7: e42350
ChIP-Seq	FLAG-BAP1, OGT, HCF1	Science	2012	Anwesha Dey *et al.* Loss of the Tumor Suppressor BAP1 Causes Myeloid Transformation. *Science.* 337: 1541
ChIP-Seq	RNA Pol II	Journal of Biological Chemistry	2012	Danish Sayed *et al.* Transcriptional regulation patterns revealed by high-resolution chromatin immunoprecipitation during cardiac hypertrophy. *J Biol Chem.* 288: 2546
ChIP-Seq	Androgen Receptor	BMC Genomics	2012	Zhou Zhu *et al.* Dose-dependent effects of small-molecule antagonists on the genomic landscape of androgen receptor binding. *BMC Genomics.* 13: 355
ChIP-Seq	Progesterone Receptor	Molecular Endocrinology	2012	Cory A. Rubel *et al.* Genome-Wide Profiling of Progesterone Receptor Binding in the Mouse Uterus. *Mol Endocrinol.* 26: 1428
ChIP-Seq	Estrogen Receptor, RNA Pol II	Molecular Endocrinology	2012	Sylvia C. Hewitt *et al.* Whole-Genome Estrogen Receptor α Binding in Mouse Uterine Tissue Revealed by ChIP-Seq. *Mol Endocrinol.* 26: 887
ChIP-qPCR	PDX1, TCF3	Chemistry & Biology	2012	Alice Kiselyuk *et al.* HNF4α antagonists discovered by a high-throughput screen for modulators of the human insulin promoter. *Chem Biol.* 19: 806
ChIP-chip	RORα	PLoS One	2012	Yongjun Wang *et al.* Regulation of p53 Stability and Apoptosis by a ROR Agonist. *PLoS One.* 7: e34921
ChIP-Seq	Nkx3.1	The Journal of Clinical Investigation	2012	Philip D. Anderson *et al.* Nkx3.1 and Myc crossregulate shared target genes in mouse and human prostate tumorigenesis. *J Clin Invest.* 122: 1907
ChIP-Seq	RNA Pol II	Cardiovascular Research	2012	Mingyue Han *et al.* GATA4 expression is primarily regulated via a miR-26b-dependent post-transcriptional mechanism during cardiac hypertrophy. *Cardiovasc Res.* 93: 645
ChIP-Seq	FXR (Farnesoid X Receptor)	The American Journal of Physiology: Gastrointestinal and Liver Physiology	2012	Julia Yue Cui *et al.* Bile acids via FXR initiate the expression of major transporters involved in the enterohepatic circulation of bile acids in newborn mice. *Am J Physiol Gastrointest Liver Physiol.* 302: G979
ChIP-Seq	Progesterone Receptor	PLoS One	2012	Ping Yin *et al.* Genome-wide progesterone receptor binding: cell type-specific and shared mechanisms in T47D breast cancer cells and primary leiomyoma cells. *PLoS One.* 7: e29021
ChIP-Seq	FLAG-CCND1 (Cyclin D1)	The Journal of Clinical Investigation	2012	Mathew C. Casimiro *et al.* ChIP sequencing of cyclin D1 reveals a transcriptional role in chromosomal instability in mice. *J Clin Invest.* 122: 833
ChIP-Seq	RBPJ	Stem Cells	2012	Yaochen Li *et al.* Genome-wide Analysis of N1ICD/RBPJ Targets In Vivo Reveals Direct Transcriptional Regulation of Wnt, SHH, and Hippo Pathway Effectors by Notch1. *Stem Cells.* 30: 741
ChIP-chip	Progesterone Receptor	The FASEB Journal	2011	Heather L. Franco *et al.* Epithelial progesterone receptor exhibits pleiotropic roles in uterine development and function. *FASEB J.* 26: 1218
ChIP-Seq	Msgn1	Nature Communication	2011	Ravindra B. Chalamalasetty *et al.* The Wnt3a/β-catenin target gene Mesogenin1 controls the segmentation clock by activating a Notch signalling program. *Nat Commun.* 2: 390

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ChIP-qPCR	IRF4, Myc	British Journal of Haematology	2011	Antonia Lopez-Girona *et al.* Lenalidomide downregulates the cell survival factor, interferon regulatory factor-4, providing a potential mechanistic link for predicting response. *Br J Haematol.* 154: 325
ChIP-chip	IRF4	Allergy	2011	Barrenas S. Bruhn *et al.* Increased expression of IRF4 and ETS1 in CD4+ cells from patients with intermittent allergic rhinitis. *Allergy.* 67: 33
ChIP-Seq	COBRA1 (Nelf-b)	Journal of Biological Chemistry	2011	Jianlong Sun *et al.* Genetic and Genomic Analyses of RNA Polymerase II-pausing Factor in Regulation of Mammalian Transcription and Cell Growth. *J Biol Chem.* 286: 36248
ChIP-Seq	Foxp3	Nucleic Acids Research	2011	Fabian Birzele *et al.* Next-generation insights into regulatory T cells: expression profiling and FoxP3 occupancy in Human. *Nucleic Acids Res.* 39: 7946
ChIP-qPCR	BRD4	Proceedings of the National Academy of Sciences	2011	Jennifer A. Mertz *et al.* Targeting MYC dependence in cancer by inhibiting BET bromodomains. *Proc Natl Acad Sci.* 108: 16669
ChIP-chip	C/EBPβ	Journal of Molecular Endocrinology	2011	Aristides Lytras *et al.* Identification of functional CCAAT/ enhancer-binding protein and Ets protein binding sites in the human chorionic somatomammotropin enhancer sequences. *J Mol Endocrinol.* 47: 179
ChIP-qPCR	Progesterone Receptor	Biochemical Pharmacology	2011	Mattew R. Yudt *et al.* Discovery of a novel mechanism of steroid receptor antagonism: WAY-255348 modulates progesterone receptor cellular localization and promoter interactions. *Biochem Pharmacol.* 82: 1709
ChIP-chip	RNA Pol II, H3K9Ac	Journal of Biological Chemistry	2011	Hong Hao *et al.* The Transcription Factor Neural Retina Leucine Zipper (NRL) Controls Photoreceptor-specific Expression of Myocyte Enhancer Factor Mef2c from an Alternative Promoter. *J Biol Chem.* 286: 34893
ChIP-chip	DAF-12	PLoS Genetics	2011	Daniel Hochbaum *et al.* DAF-12 Regulates a Connected Network of Genes to Ensure Robust Developmental Decisions. *PLoS Genet.* 7: e1002179
ChIP-qPCR	CTCF	DNA and Cell Biology	2011	Yan Jin *et al.* Enhancer-Blocking Activity Is Associated with Hypersensitive Site V Sequences in the Human Growth Hormone Locus Control Region. *DNA Cell Biol.* 30: 995
Bisulfite-Seq	Bisulfite Sequencing	PLoS One	2011	Michela Deleidi *et al.* Oct4-Induced Reprogramming Is Required for Adult Brain Neural Stem Cell Differentiation into Midbrain Dopaminergic Neurons. *PLoS ONE.* 6: e19926
MeDIP-chip	5-methylcytosine (5-mC)	BMC Biology	2011	Genevieve Lavoie *et al.* PKC isoforms interact with and phosphorylate DNMT1. *BMC Biol.* 9: 31
ChIP-qPCR	AHR (Aryl Hydrocarbon Receptor)	Toxicology and Applied Pharmacology	2011	K. Nadira De Abrew *et al.* Regulation of Bach2 by the aryl hydrocarbon receptor as a mechanism for suppression of B-cell differentiation by 2,3,7,8-tetrachlorodibenzo-p-dioxin. *Toxicol Appl Pharmacol.* 252: 150
ChIP-qPCR	PPARγ	Investigative Ophthalmology and Visual Science	2011	Gerard A. Rodrigues *et al.* Differential Effects of PPARγ Ligands on Oxidative Stress–Induced Death of Retinal Pigmented Epithelial Cells. *Invest Ophthalmol Vis Sci.* 52: 890
ChIP-chip	p73	Proceedings of the National Academy of Sciences	2011	Jennifer M. Rosenbluth *et al.* Differential regulation of the p73 cistrome by mammalian target of rapamycin reveals transcriptional programs of mesenchymal differentiation and tumorigenesis. *Proc Natl Acad Sci.* 108: 2076

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ChIP-chip	LXR, RXR	Journal of Biological Chemistry	2011	Qi Shen *et al.* Liver X Receptor-Retinoid X Receptor (LXR-RXR) Heterodimer Cistrome Reveals Coordination of LXR and AP1 Signaling in Keratinocytes. *J Biol Chem.* 286: 14554
ChIP-qPCR	AHR (Aryl Hydrocarbon Receptor), MAF	Nature Immunology	2010	Lionel Apetoh *et al.* The aryl hydrocarbon receptor interacts with c-Maf to promote the differentiation of type 1 regulatory T cells induced by IL-27. *Nat Immunol.* 11: 854
ChIP-qPCR	p73	Molecular Cancer	2010	Kathryn G. Eby *et al.* ISG20L1 is a p53 family target gene that modulates genotoxic stress-induced autophagy. *Mol Cancer.* 9: 95
ChIP-Seq	Vitamin D Receptor	Genome Research	2010	Sreeram V. Ramagopalan *et al.* A ChIP-seq defined genome-wide map of vitamin D receptor binding: associations with disease and evolution. *Genome Res.* 20: 1352
ChIP-chip	O-GlcNAc, RNA Pol II	Proceedings of the National Academy of Sciences	2010	Dona C. Love *et al.* Dynamic O-GlcNAc cycling at promoters of *Caenorhabditis elegans* genes regulating longevity, stress, and immunity. *Proc Natl Acad Sci.* 107: 7413
ChIP-Seq	PXR (Pregnane X Receptor)	Toxicological Sciences	2010	Julia Yue Cui *et al.* Genetic and Epigenetic Regulation and Expression Signatures of Glutathione S-Transferases in Developing Mouse Liver. *Toxicol Sci.* 116: 32
ChIP-Seq	SRC3, RNA Pol II	Molecular Endocrinology	2010	Rainer B. Lanz *et al.* Global Characterization of Transcriptional Impact of the SRC-3 Coregulator. *Mol Endocrinol.* 24: 859
ChIP-qPCR	Maf1, Rpc39, mTOR, Brf1, Raptor	Journal of Biological Chemistry	2010	Boris Shor *et al.* Requirement of the mTOR Kinase for the Regulation of Maf1 Phosphorylation and Control of RNA Polymerase III-dependent Transcription in Cancer Cells. *J Biol Chem.* 285: 15380
ChIP-chip	AHR (Aryl Hydrocarbon Receptor)	Toxicological Sciences	2010	K. Nadira De Abrew *et al.* An Integrated Genomic Analysis of Aryl Hydrocarbon Receptor-Mediated Inhibition of B-Cell Differentiation. *Toxicol Sci.* 118: 454
ChIP-qPCR	Myc-tagged-Pet-1	Nature Neuroscience	2010	Chen Liu *et al.* Pet-1 is required across different stages of life to regulate serotonergic function. *Nat Neurosci.* 13: 1190
ChIP-chip	Androgen Receptor	Molecular Endocrinology	2010	Anastasia Wyce *et al.* The Androgen Receptor Modulates Expression of Genes with Critical Roles in Muscle Development and Function. *Mol Endocrinol.* 24: 1665
ChIP-qPCR	NFκB (p50 and p65)	Cardiovascular Research	2010	Alina G. Sofronescu *et al.* FGF-16 is a target for adrenergic stimulation through NF-κB activation in postnatal cardiac cells and adult mouse heart. *Cardiovasc Res.* 87: 102
ChIP-Seq	PXR (Pregnane X Receptor)	Nucleic Acids Research	2010	Julia Yue Cui *et al.* ChIPing the cistrome of PXR in mouse liver. *Nucleic Acids Res.* 38: 7943
ChIP-chip	RNA Pol II	Molecular Vision	2010	Padmaja Tummala *et al.* Temporal ChIP-on-Chip of RNA-Polymerase-II to detect novel gene activation events during photoreceptor maturation. *Mol Vis.* 16: 252
ChIP-chip	STAT4	The Journal of Immunology	2009	Seth R. Good *et al.* Temporal Induction Pattern of STAT4 Target Genes Defines Potential for Th1 Lineage-Specific Programming. *J Immunol.* 183: 3839
ChIP-Seq	p73	Molecular and Cellular Biology	2008	Jennifer M. Rosenbluth *et al.* A Gene Signature-Based Approach Identifies mTOR as a Regulator of p73. *Mol Cell Biol.* 28: 5951
ChIP-qPCR	AHR (Aryl Hydrocarbon Receptor)	Nature	2008	Francisco J. Quintana *et al.* Control of Treg and TH17 cell differentiation by the aryl hydrocarbon receptor. *Nature.* 453: 65