## Andrew K Dingwall

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The Drosophila MLR COMPASS-like complex regulates bantam miRNA expression differentially in the context of cell fate. Developmental Biology, 2020, 468, 41-53.	2.0	2
2	The Drosophila MLR COMPASS complex is essential for programming cis-regulatory information and maintaining epigenetic memory during development. Nucleic Acids Research, 2020, 48, 3476-3495.	14.5	8
3	DAXX Suppresses Tumor-Initiating Cells in Estrogen Receptor–Positive Breast Cancer Following Endocrine Therapy. Cancer Research, 2019, 79, 4965-4977.	0.9	27
4	COMPASS Ascending: Emerging clues regarding the roles of MLL3/KMT2C and MLL2/KMT2D proteins in cancer. Cancer Letters, 2019, 458, 56-65.	7.2	121
5	The cancer COMPASS: navigating the functions of MLL complexes in cancer. Cancer Genetics, 2015, 208, 178-191.	0.4	122
6	The Drosophila COMPASS-like Cmi-Trr coactivator complex regulates dpp/BMP signaling in pattern formation. Developmental Biology, 2013, 380, 185-198.	2.0	13
7	Drosophila LSD1 oREST demethylase complex regulates DPP/TGFβ signaling during wing development. Genesis, 2013, 51, 16-31.	1.6	6
8	Histone recognition and nuclear receptor co-activator functions of <i>Drosophila</i> Cara Mitad, a homolog of the N-terminal portion of mammalian MLL2 and MLL3. Development (Cambridge), 2012, 139, 1997-2008.	2.5	35
9	The chromatin remodeling and mRNA splicing functions of the Brahma (SWI/SNF) complex are mediated by the SNR1/SNF5 regulatory subunit. Nucleic Acids Research, 2012, 40, 5975-5987.	14.5	40
10	Histone lysine demethylases function as co-repressors of SWI/SNF remodeling activities during Drosophila wing development. Developmental Biology, 2011, 350, 534-547.	2.0	26
11	Congenital anomalies and rhabdoid tumor associated with 22q11 germline deletion and somatic inactivation of the <i>SMARCB1</i> tumor suppressor. Genes Chromosomes and Cancer, 2011, 50, 379-388.	2.8	20
12	Hormone-response Genes Are Direct in Vivo Regulatory Targets of Brahma (SWI/SNF) Complex Function. Journal of Biological Chemistry, 2006, 281, 35305-35315.	3.4	44
13	SNR1 (INI1/SNF5) Mediates Important Cell Growth Functions of the Drosophila Brahma (SWI/SNF) Chromatin Remodeling Complex. Genetics, 2004, 168, 199-214.	2.9	22
14	The Drosophila Brahma (SWI/SNF) chromatin remodeling complex exhibits cell-type specific activation and repression functions. Developmental Biology, 2004, 267, 279-293.	2.0	54
15	SNR1 is an essential subunit in a subset of drosophila brm complexes, targeting specific functions during development. Developmental Biology, 2003, 253, 291-308.	2.0	48
16	The Drosophila SNR1 (SNF5/INI1) Subunit Directs Essential Developmental Functions of the Brahma Chromatin Remodeling Complex. Molecular and Cellular Biology, 2003, 23, 289-305.	2.3	35
17	Drosophila cyclin E interacts with components of the Brahma complex. EMBO Journal, 2002, 21, 3377-3389.	7.8	42
18	Genetic and Molecular Analysis of Region 88E9;88F2 in Drosophila melanogaster, Including the ear Gene Related to Human Factors Involved in Lineage-Specific Leukemias. Genetics, 2002, 160, 1051-1065.	2.9	8

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#	Article	IF	CITATIONS
19	Organization and ordered expression of Caulobacter genes encoding flagellar basal body rod and ring proteins. Journal of Molecular Biology, 1992, 228, 1147-1162.	4.2	41
20	Identification of Cis and Trans-elements involved in the timed control of a Caulobacter flagellar gene. Journal of Molecular Biology, 1991, 217, 247-257.	4.2	30
21	Plasmid and chromosomal DNA replication and partitioning during the Caulobacter crescentus cell cycle. Journal of Molecular Biology, 1990, 212, 709-722.	4.2	65
22	Temporal regulation and overlap organization of two Caulobacter flagellar genes. Journal of Molecular Biology, 1989, 205, 71-83.	4.2	26
23	Cascade regulation of Caulobacter flagellar and chemotaxis genes. Journal of Molecular Biology, 1987, 194, 71-80.	4.2	81