

# Andrew K Dingwall

## List of Publications by Year in descending order

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23  
papers

916  
citations

430874

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642732

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26  
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26  
docs citations

26  
times ranked

1144  
citing authors

#	ARTICLE	IF	CITATIONS
1	The cancer COMPASS: navigating the functions of MLL complexes in cancer. <i>Cancer Genetics</i> , 2015, 208, 178-191.	0.4	122
2	COMPASS Ascending: Emerging clues regarding the roles of MLL3/KMT2C and MLL2/KMT2D proteins in cancer. <i>Cancer Letters</i> , 2019, 458, 56-65.	7.2	121
3	Cascade regulation of <i>Caulobacter</i> flagellar and chemotaxis genes. <i>Journal of Molecular Biology</i> , 1987, 194, 71-80.	4.2	81
4	Plasmid and chromosomal DNA replication and partitioning during the <i>Caulobacter crescentus</i> cell cycle. <i>Journal of Molecular Biology</i> , 1990, 212, 709-722.	4.2	65
5	The <i>Drosophila</i> Brahma (SWI/SNF) chromatin remodeling complex exhibits cell-type specific activation and repression functions. <i>Developmental Biology</i> , 2004, 267, 279-293.	2.0	54
6	SNR1 is an essential subunit in a subset of <i>drosophila</i> brm complexes, targeting specific functions during development. <i>Developmental Biology</i> , 2003, 253, 291-308.	2.0	48
7	Hormone-response Genes Are Direct in Vivo Regulatory Targets of Brahma (SWI/SNF) Complex Function. <i>Journal of Biological Chemistry</i> , 2006, 281, 35305-35315.	3.4	44
8	<i>Drosophila</i> cyclin E interacts with components of the Brahma complex. <i>EMBO Journal</i> , 2002, 21, 3377-3389.	7.8	42
9	Organization and ordered expression of <i>Caulobacter</i> genes encoding flagellar basal body rod and ring proteins. <i>Journal of Molecular Biology</i> , 1992, 228, 1147-1162.	4.2	41
10	The chromatin remodeling and mRNA splicing functions of the Brahma (SWI/SNF) complex are mediated by the SNR1/SNF5 regulatory subunit. <i>Nucleic Acids Research</i> , 2012, 40, 5975-5987.	14.5	40
11	The <i>Drosophila</i> SNR1 (SNF5/INI1) Subunit Directs Essential Developmental Functions of the Brahma Chromatin Remodeling Complex. <i>Molecular and Cellular Biology</i> , 2003, 23, 289-305.	2.3	35
12	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. <i>Development (Cambridge)</i> , 2012, 139, 1997-2008.	2.5	35
13	Identification of Cis and Trans-elements involved in the timed control of a <i>Caulobacter</i> flagellar gene. <i>Journal of Molecular Biology</i> , 1991, 217, 247-257.	4.2	30
14	DAXX Suppresses Tumor-Initiating Cells in Estrogen Receptor-Positive Breast Cancer Following Endocrine Therapy. <i>Cancer Research</i> , 2019, 79, 4965-4977.	0.9	27
15	Temporal regulation and overlap organization of two <i>Caulobacter</i> flagellar genes. <i>Journal of Molecular Biology</i> , 1989, 205, 71-83.	4.2	26
16	Histone lysine demethylases function as co-repressors of SWI/SNF remodeling activities during <i>Drosophila</i> wing development. <i>Developmental Biology</i> , 2011, 350, 534-547.	2.0	26
17	SNR1 (INI1/SNF5) Mediates Important Cell Growth Functions of the <i>Drosophila</i> Brahma (SWI/SNF) Chromatin Remodeling Complex. <i>Genetics</i> , 2004, 168, 199-214.	2.9	22
18	Congenital anomalies and rhabdoid tumor associated with 22q11 germline deletion and somatic inactivation of the <i>SMARCB1</i> tumor suppressor. <i>Genes Chromosomes and Cancer</i> , 2011, 50, 379-388.	2.8	20

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19	The Drosophila COMPASS-like Cmi-Trr coactivator complex regulates dpp/BMP signaling in pattern formation. <i>Developmental Biology</i> , 2013, 380, 185-198.	2.0	13
20	The Drosophila MLR COMPASS complex is essential for programming cis-regulatory information and maintaining epigenetic memory during development. <i>Nucleic Acids Research</i> , 2020, 48, 3476-3495.	14.5	8
21	Genetic and Molecular Analysis of Region 88E9;88F2 in <i>Drosophila melanogaster</i> , Including the ear Gene Related to Human Factors Involved in Lineage-Specific Leukemias. <i>Genetics</i> , 2002, 160, 1051-1065.	2.9	8
22	<i>Drosophila</i> LSD1/CoREST demethylase complex regulates DPP/TGF $\beta$ signaling during wing development. <i>Genesis</i> , 2013, 51, 16-31.	1.6	6
23	The <i>Drosophila</i> MLR COMPASS-like complex regulates bantam miRNA expression differentially in the context of cell fate. <i>Developmental Biology</i> , 2020, 468, 41-53.	2.0	2