Nicoletta Bobola

List of Publications by Year in descending order

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623734 580821 27 735 14 25 citations g-index h-index papers 35 35 35 1376 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Hoxa2 Selectively Enhances Meis Binding to Change a Branchial Arch Ground State. Developmental Cell, 2015, 32, 265-277.	7. O	76
2	Mouse <i>Hoxa2</i> mutations provide a model for microtia and auricle duplication. Development (Cambridge), 2013, 140, 4386-4397.	2.5	75
3	Genome-wide occupancy links Hoxa2 to Wnt–β-catenin signaling in mouse embryonic development. Nucleic Acids Research, 2012, 40, 3990-4001.	14.5	71
4	An integrative transcriptomic atlas of organogenesis in human embryos. ELife, 2016, 5, .	6.0	61
5	Homeodomain proteins in action: similar DNA binding preferences, highly variable connectivity. Current Opinion in Genetics and Development, 2017, 43, 1-8.	3. 3	41
6	Mesenchymal patterning by Hoxa2 requires blocking Fgf-dependent activation of Ptx1. Development (Cambridge), 2003, 130, 3403-3414.	2.5	40
7	Hoxa2 downregulates Six2 in the neural crest-derived mesenchyme. Development (Cambridge), 2005, 132, 469-478.	2.5	40
8	Six2 functions redundantly immediately downstream of Hoxa2. Development (Cambridge), 2008, 135, 1463-1470.	2.5	39
9	Inactivation of Six2 in mouse identifies a novel genetic mechanism controlling development and growth of the cranial base. Developmental Biology, 2010, 344, 720-730.	2.0	38
10	Laser Capture and Deep Sequencing Reveals the Transcriptomic Programmes Regulating the Onset of Pancreas and Liver Differentiation in Human Embryos. Stem Cell Reports, 2017, 9, 1387-1394.	4.8	37
11	TALE factors use two distinct functional modes to control an essential zebrafish gene expression program. ELife, 2018, 7, .	6.0	31
12	HOX paralogs selectively convert binding of ubiquitous transcription factors into tissue-specific patterns of enhancer activation. PLoS Genetics, 2020, 16, e1009162.	3.5	23
13	Transient Activation of Meox1 Is an Early Component of the Gene Regulatory Network Downstream of Hoxa2. Molecular and Cellular Biology, 2011, 31, 1301-1308.	2.3	20
14	Epimorphin Alters the Inhibitory Effects of SOX9 on Mmp13 in Activated Hepatic Stellate Cells. PLoS ONE, 2014, 9, e100091.	2.5	19
15	Dynamic changes in the epigenomic landscape regulate human organogenesis and link to developmental disorders. Nature Communications, 2020, 11, 3920.	12.8	17
16	Diabetes Inhibits Gr-1+Myeloid Cell Maturation via Cebpa Deregulation. Diabetes, 2015, 64, 4184-4197.	0.6	14
17	A distal 594bp ECR specifies $\langle i\rangle$ Hmx $1\langle i\rangle$ expression in pinna and lateral facial morphogenesis and is regulated by Hox-Pbx-Meis. Development (Cambridge), 2016, 143, 2582-92.	2.5	13
18	Uncovering tissue-specific binding features from differential deep learning. Nucleic Acids Research, 2020, 48, e27-e27.	14.5	13

#	Article	IF	CITATIONS
19	The spatial phenotype of genotypically distinct meningiomas demonstrate potential implications of the embryology of the meninges. Oncogene, 2021, 40, 875-884.	5.9	13
20	A tissue-specific, Gata6-driven transcriptional program instructs remodeling of the mature arterial tree. ELife, $2017, 6, .$	6.0	13
21	Expressing Hoxa2 across the entire endochondral skeleton alters the shape of the skeletal template in a spatially restricted fashion. Differentiation, 2010, 79, 194-202.	1.9	11
22	Combinatorial action of NF–Y and TALE at embryonic enhancers defines distinct gene expression programs during zygotic genome activation in zebrafish. Developmental Biology, 2020, 459, 161-180.	2.0	8
23	IGFBP5 is a potential regulator of craniofacial skeletogenesis. Genesis, 2008, 46, 52-59.	1.6	6
24	Chromatin Immunoprecipitation and Chromatin Immunoprecipitation with Massively Parallel Sequencing on Mouse Embryonic Tissue. Methods in Molecular Biology, 2014, 1196, 231-239.	0.9	5
25	Differential Distribution of the Ca (2+) Regulator Pcp4 in the Branchial Arches Is Regulated by Hoxa2. PLoS ONE, 2013, 8, e63160.	2.5	2
26	Molecular Characterization of HOXA2 and HOXA3 Binding Properties. Journal of Developmental Biology, 2021, 9, 55.	1.7	2
27	From DNA binding to transcriptional activation: Is the TALE complete?. Journal of Cell Biology, 2017, 216, 2603-2605.	5.2	1