

# Ferenc Mueller

## List of Publications by Year in descending order

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87  
papers

7,125  
citations

87723

38  
h-index

66788

78  
g-index

98  
all docs

98  
docs citations

98  
times ranked

14009  
citing authors

#	ARTICLE	IF	CITATIONS
1	An atlas of active enhancers across human cell types and tissues. <i>Nature</i> , 2014, 507, 455-461.	13.7	2,269
2	Pancreatic islet enhancer clusters enriched in type 2 diabetes risk-associated variants. <i>Nature Genetics</i> , 2014, 46, 136-143.	9.4	475
3	Zebrafish embryos as models for embryotoxic and teratological effects of chemicals. <i>Reproductive Toxicology</i> , 2009, 28, 245-253.	1.3	240
4	Germline mutations in DIS3L2 cause the Perlman syndrome of overgrowth and Wilms tumor susceptibility. <i>Nature Genetics</i> , 2012, 44, 277-284.	9.4	219
5	Prepatterning of Developmental Gene Expression by Modified Histones before Zygotic Genome Activation. <i>Developmental Cell</i> , 2011, 21, 993-1004.	3.1	188
6	Two independent transcription initiation codes overlap on vertebrate core promoters. <i>Nature</i> , 2014, 507, 381-385.	13.7	182
7	Transcriptional profiling reveals barcode-like toxicogenomic responses in the zebrafish embryo. <i>Genome Biology</i> , 2007, 8, R227.	13.9	166
8	Mutations in VIPAR cause an arthrogryposis, renal dysfunction and cholestasis syndrome phenotype with defects in epithelial polarization. <i>Nature Genetics</i> , 2010, 42, 303-312.	9.4	162
9	Loss-of-Function Mutations in RAB18 Cause Warburg Micro Syndrome. <i>American Journal of Human Genetics</i> , 2011, 88, 499-507.	2.6	158
10	Sonic hedgehog, secreted by amacrine cells, acts as a short-range signal to direct differentiation and lamination in the zebrafish retina. <i>Development (Cambridge)</i> , 2004, 131, 3849-3858.	1.2	128
11	Regulation and expression of transgenes in fish—a review. <i>Transgenic Research</i> , 1996, 5, 147-166.	1.3	127
12	Automated high-throughput mapping of promoter-enhancer interactions in zebrafish embryos. <i>Nature Methods</i> , 2009, 6, 911-916.	9.0	123
13	Dynamic regulation of the transcription initiation landscape at single nucleotide resolution during vertebrate embryogenesis. <i>Genome Research</i> , 2013, 23, 1938-1950.	2.4	119
14	A Multicassette Gateway Vector Set for High Throughput and Comparative Analyses in Ciona and Vertebrate Embryos. <i>PLoS ONE</i> , 2007, 2, e916.	1.1	113
15	Functional annotation of human long noncoding RNAs via molecular phenotyping. <i>Genome Research</i> , 2020, 30, 1060-1072.	2.4	109
16	TBP is not universally required for zygotic RNA polymerase II transcription in zebrafish. <i>Current Biology</i> , 2001, 11, 282-287.	1.8	102
17	Splicing Segregation: The Minor Spliceosome Acts outside the Nucleus and Controls Cell Proliferation. <i>Cell</i> , 2007, 131, 718-729.	13.5	97
18	The multicoloured world of promoter recognition complexes. <i>EMBO Journal</i> , 2004, 23, 2-8.	3.5	83

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19	Cloning and expression analysis of an inducible HSP70 gene from tilapia fish. <i>FEBS Letters</i> , 2000, 474, 5-10.	1.3	81
20	Search for enhancers: teleost models in comparative genomic and transgenic analysis of cisregulatory elements. <i>BioEssays</i> , 2002, 24, 564-572.	1.2	80
21	TBP2, a Vertebrate-Specific Member of the TBP Family, Is Required in Embryonic Development of Zebrafish. <i>Current Biology</i> , 2004, 14, 593-598.	1.8	80
22	New Problems in RNA Polymerase II Transcription Initiation: Matching the Diversity of Core Promoters with a Variety of Promoter Recognition Factors. <i>Journal of Biological Chemistry</i> , 2007, 282, 14685-14689.	1.6	80
23	Enhancers active in dopamine neurons are a primary link between genetic variation and neuropsychiatric disease. <i>Nature Neuroscience</i> , 2018, 21, 1482-1492.	7.1	79
24	Cooperation of sonic hedgehog enhancers in midline expression. <i>Developmental Biology</i> , 2007, 301, 578-589.	0.9	78
25	Developmental regulation of transcription initiation: more than just changing the actors. <i>Current Opinion in Genetics and Development</i> , 2010, 20, 533-540.	1.5	76
26	Chromatin and DNA sequences in defining promoters for transcription initiation. <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2014, 1839, 118-128.	0.9	69
27	Automated feature detection and imaging for high-resolution screening of zebrafish embryos. <i>BioTechniques</i> , 2011, 50, 319-324.	0.8	65
28	Optimisation of Embryonic and Larval ECG Measurement in Zebrafish for Quantifying the Effect of QT Prolonging Drugs. <i>PLoS ONE</i> , 2013, 8, e60552.	1.1	61
29	The TATA-binding protein regulates maternal mRNA degradation and differential zygotic transcription in zebrafish. <i>EMBO Journal</i> , 2007, 26, 3945-3956.	3.5	57
30	Developmental toxicity and estrogenic potency of zearalenone in zebrafish ( <i>Danio rerio</i> ). <i>Aquatic Toxicology</i> , 2013, 136-137, 13-21.	1.9	55
31	Activator effect of coinjected enhancers on the muscle-specific expression of promoters in zebrafish embryos. <i>Molecular Reproduction and Development</i> , 1997, 47, 404-412.	1.0	53
32	Synthesis of azetidines and pyrrolidines via iodocyclisation of homoallyl amines and exploration of activity in a zebrafish embryo assay. <i>Organic and Biomolecular Chemistry</i> , 2013, 11, 5083.	1.5	53
33	Targeted transgene integration overcomes variability of position effects in zebrafish. <i>Development (Cambridge)</i> , 2014, 141, 715-724.	1.2	53
34	Efficient transient expression system based on square pulse electroporation and in vivo luciferase assay of fertilized fish eggs. <i>FEBS Letters</i> , 1993, 324, 27-32.	1.3	47
35	Hedgehog signaling patterns the outgrowth of unpaired skeletal appendages in zebrafish. <i>BMC Developmental Biology</i> , 2007, 7, 75.	2.1	46
36	Characterization of zebrafish smad1, smad2 and smad5: the amino-terminus of Smad1 and Smad5 is required for specific function in the embryo. <i>Mechanisms of Development</i> , 1999, 88, 73-88.	1.7	43

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37	A Floor Plate Enhancer of the Zebrafish <i>netrin1</i> Gene Requires Cyclops (Nodal) Signalling and the Winged Helix Transcription Factor FoxA2. <i>Developmental Biology</i> , 2002, 252, 1-14.	0.9	42
38	A cell cycle-coordinated Polymerase II transcription compartment encompasses gene expression before global genome activation. <i>Nature Communications</i> , 2019, 10, 691.	5.8	42
39	Shuffling of cis-regulatory elements is a pervasive feature of the vertebrate lineage. <i>Genome Biology</i> , 2006, 7, R56.	13.9	41
40	Redefining the Initiation and Maintenance of Zebrafish Interrenal Steroidogenesis by Characterizing the Key Enzyme <i>Cyp11a2</i> . <i>Endocrinology</i> , 2013, 154, 2702-2711.	1.4	38
41	Transcriptional, post-transcriptional and chromatin-associated regulation of pri-miRNAs, pre-miRNAs and moRNAs. <i>Nucleic Acids Research</i> , 2016, 44, 3070-3081.	6.5	38
42	Dual-initiation promoters with intertwined canonical and TCT/TOP transcription start sites diversify transcript processing. <i>Nature Communications</i> , 2020, 11, 168.	5.8	37
43	High transgene activity in the yolk syncytial layer affects quantitative transient expression assays in zebrafish ( <i>Danio rerio</i> ) embryos. <i>Transgenic Research</i> , 1996, 5, 433-442.	1.3	34
44	SLIC-CAGE: high-resolution transcription start site mapping using nanogram-levels of total RNA. <i>Genome Research</i> , 2018, 28, 1943-1956.	2.4	33
45	Methyl Mercury Suppresses the Formation of the Tail Primordium in Developing Zebrafish Embryos. <i>Toxicological Sciences</i> , 2010, 115, 379-390.	1.4	31
46	Ferredoxin 1b ( <i>Fdx1b</i> ) Is the Essential Mitochondrial Redox Partner for Cortisol Biosynthesis in Zebrafish. <i>Endocrinology</i> , 2016, 157, 1122-1134.	1.4	29
47	Fish as Bioreactors: Transgene Expression of Human Coagulation Factor VII in Fish Embryos. <i>Marine Biotechnology</i> , 2004, 6, 485-492.	1.1	28
48	Novel SPG11 mutations in Asian kindreds and disruption of spatascin function in the zebrafish. <i>Neurogenetics</i> , 2010, 11, 379-389.	0.7	26
49	Multiomic atlas with functional stratification and developmental dynamics of zebrafish cis-regulatory elements. <i>Nature Genetics</i> , 2022, 54, 1037-1050.	9.4	26
50	Transposition and targeting of the prokaryotic mobile element IS30 in zebrafish. <i>FEBS Letters</i> , 2003, 550, 46-50.	1.3	25
51	The identification and functional characterisation of conserved regulatory elements in developmental genes. <i>Briefings in Functional Genomics &amp; Proteomics</i> , 2005, 3, 332-350.	3.8	24
52	Highly conserved elements discovered in vertebrates are present in non-syntenic loci of tunicates, act as enhancers and can be transcribed during development. <i>Nucleic Acids Research</i> , 2013, 41, 3600-3618.	6.5	24
53	Genetic Disruption of 21-Hydroxylase in Zebrafish Causes Interrenal Hyperplasia. <i>Endocrinology</i> , 2017, 158, 4165-4173.	1.4	24
54	TBPL2/TFIIA complex establishes the maternal transcriptome through oocyte-specific promoter usage. <i>Nature Communications</i> , 2020, 11, 6439.	5.8	23

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55	Two Lamprey Hedgehog Genes Share Non-Coding Regulatory Sequences and Expression Patterns with Gnathostome Hedgehogs. <i>PLoS ONE</i> , 2010, 5, e13332.	1.1	22
56	Liposome-mediated gene transfer in fish embryos. <i>Transgenic Research</i> , 1994, 3, 116-119.	1.3	20
57	Cyclops-independent floor plate differentiation in zebrafish embryos. <i>Developmental Dynamics</i> , 2003, 226, 59-66.	0.8	19
58	Transcriptional Regulation During Zygotic Genome Activation in Zebrafish and Other Anamniote Embryos. <i>Advances in Genetics</i> , 2016, 95, 161-194.	0.8	18
59	Germ cell differentiation requires Tdrd7-dependent chromatin and transcriptome reprogramming marked by germ plasm relocalization. <i>Developmental Cell</i> , 2021, 56, 641-656.e5.	3.1	18
60	Estrogen sensitive liver transgenic zebrafish ( <i>Danio rerio</i> ) line (Tg(vtg1:mCherry)) suitable for the direct detection of estrogenicity in environmental samples. <i>Aquatic Toxicology</i> , 2019, 208, 157-167.	1.9	17
61	Phenotypic effects in <i>Xenopus</i> and zebrafish suggest that one-eyed pinhead functions as antagonist of BMP signalling. <i>Mechanisms of Development</i> , 2000, 94, 37-46.	1.7	16
62	Functional diversification of sonic hedgehog paralog enhancers identified by phylogenomic reconstruction. <i>Genome Biology</i> , 2007, 8, R106.	13.9	15
63	Conservation of shh cis-regulatory architecture of the coelacanth is consistent with its ancestral phylogenetic position. <i>EvoDevo</i> , 2010, 1, 11.	1.3	15
64	The Tetraodon nigroviridis reference transcriptome: developmental transition, length retention and microsynteny of long non-coding RNAs in a compact vertebrate genome. <i>Scientific Reports</i> , 2016, 6, 33210.	1.6	14
65	Expression and activity profiling of the steroidogenic enzymes of glucocorticoid biosynthesis and the co-factors in zebrafish. <i>Journal of Neuroendocrinology</i> , 2018, 30, e12586.	1.2	14
66	The Development and Growth of Tissues Derived from Cranial Neural Crest and Primitive Mesoderm Is Dependent on the Ligand Status of Retinoic Acid Receptor $\beta$ : Evidence That Retinoic Acid Receptor $\beta$ Functions to Maintain Stem/Progenitor Cells in the Absence of Retinoic Acid. <i>Stem Cells and Development</i> , 2015, 24, 507-519.	1.1	13
67	Comparative Aspects of Alternative Laboratory Fish Models. <i>Zebrafish</i> , 2005, 2, 47-54.	0.5	12
68	Glucocorticoid deficiency causes transcriptional and post-transcriptional reprogramming of glutamine metabolism. <i>EBioMedicine</i> , 2018, 36, 376-389.	2.7	12
69	Cellular rearrangement of the prechordal plate contributes to eye degeneration in the cavefish. <i>Developmental Biology</i> , 2018, 441, 221-234.	0.9	12
70	A native, highly active <i>Tc1/mariner</i> transposon from zebrafish ( <i>ZB</i> ) offers an efficient genetic manipulation tool for vertebrates. <i>Nucleic Acids Research</i> , 2021, 49, 2126-2140.	6.5	11
71	Ancestrally Duplicated Conserved Noncoding Element Suggests Dual Regulatory Roles of HOTAIR in cis and trans. <i>Science</i> , 2020, 23, 101008.	1.9	9
72	Intraovarian transplantation of stage I-II follicles results in viable zebrafish embryos. <i>International Journal of Developmental Biology</i> , 2010, 54, 585-589.	0.3	8

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73	Description of Embryonic Development of Spotted Green Pufferfish ( <i>Tetraodon nigroviridis</i> ). Zebrafish, 2014, 11, 509-517.	0.5	8
74	Enhancer Trapping and Annotation in Zebrafish Mediated with Sleeping Beauty, piggyBac and Tol2 Transposons. Genes, 2018, 9, 630.	1.0	8
75	Embryonic tissue differentiation is characterized by transitions in cell cycle dynamic-associated core promoter regulation. Nucleic Acids Research, 2020, 48, 8374-8392.	6.5	8
76	TBP2 is a general transcription factor specialized for female germ cells. Journal of Biology, 2009, 8, 97.	2.7	7
77	3D Finite Element Electrical Model of Larval Zebrafish ECG Signals. PLoS ONE, 2016, 11, e0165655.	1.1	7
78	Testing of Cis-Regulatory Elements by Targeted Transgene Integration in Zebrafish Using PhiC31 Integrase. Methods in Molecular Biology, 2016, 1451, 81-91.	0.4	7
79	Identification of downstream effectors of retinoic acid specifying the zebrafish pancreas by integrative genomics. Scientific Reports, 2021, 11, 22717.	1.6	6
80	Minor splicing: Nuclear dogma still in question. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, E37.	3.3	3
81	Expression and knockdown of zebrafish folliculin suggests requirement for embryonic brain morphogenesis. BMC Developmental Biology, 2016, 16, 23.	2.1	3
82	Fish genomics: casting the net wide. Briefings in Functional Genomics, 2014, 13, 79-81.	1.3	2
83	Using Tg(Vtg1:mcherry) Zebrafish Embryos to Test the Estrogenic Effects of Endocrine Disrupting Compounds. Journal of Visualized Experiments, 2020, , .	0.2	1
84	Protocol for intelligent high-content screening of zebrafish embryos on a standard widefield screening microscope. BioTechniques, 2017, 62, xx.	0.8	0
85	A Method for Zebrafish Follicle Transplantation into Recipient Mothers for the Generation of Fertilizable Eggs and Viable Offspring. Methods in Molecular Biology, 2019, 1920, 343-352.	0.4	0
86	Visualization of Transcriptional Activity in Early Zebrafish Primordial Germ. Methods in Molecular Biology, 2021, 2218, 185-194.	0.4	0
87	Sequence Analyses to Study the Evolutionary History and Cis-Regulatory Elements of Hedgehog Genes. Methods in Molecular Biology, 2007, 397, 231-250.	0.4	0