Walter Schaffner

List of Publications by Citations

Source: https://exaly.com/author-pdf/11492238/walter-schaffner-publications-by-citations.pdf

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

16,625 128 58 142 h-index g-index citations papers 6.08 146 15.2 17,300 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
142	Rapid detection of octamer binding proteins with Rinini-extracts prepared from a small number of cells. <i>Nucleic Acids Research</i> , 1989 , 17, 6419	20.1	3646
141	Expression of a beta-globin gene is enhanced by remote SV40 DNA sequences. <i>Cell</i> , 1981 , 27, 299-308	56.2	1459
140	A lymphocyte-specific cellular enhancer is located downstream of the joining region in immunoglobulin heavy chain genes. <i>Cell</i> , 1983 , 33, 729-40	56.2	1434
139	Enhancers and eukaryotic gene transcription. <i>Trends in Genetics</i> , 1985 , 1, 224-230	8.5	563
138	A cloned octamer transcription factor stimulates transcription from lymphoid-specific promoters in non-B cells. <i>Nature</i> , 1988 , 336, 544-51	50.4	506
137	A lymphocyte-specific enhancer in the mouse immunoglobulin kappa gene. <i>Nature</i> , 1984 , 307, 80-2	50.4	429
136	A small segment of polyoma virus DNA enhances the expression of a cloned beta-globin gene over a distance of 1400 base pairs. <i>Nucleic Acids Research</i> , 1981 , 9, 6251-64	20.1	354
135	A B-cell coactivator of octamer-binding transcription factors. <i>Nature</i> , 1995 , 373, 360-2	50.4	280
134	OVEC, a versatile system to study transcription in mammalian cells and cell-free extracts. <i>Nucleic Acids Research</i> , 1987 , 15, 6787-98	20.1	276
133	Polyoma virus DNA replication requires an enhancer. <i>Nature</i> , 1984 , 312, 242-6	50.4	259
132	An SV40 "enhancer trap" incorporates exogenous enhancers or generates enhancers from its own sequences. <i>Cell</i> , 1984 , 36, 983-92	56.2	225
131	Thionein (apometallothionein) can modulate DNA binding and transcription activation by zinc finger containing factor Sp1. <i>FEBS Letters</i> , 1991 , 279, 310-2	3.8	215
130	The taste of heavy metals: gene regulation by MTF-1. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2012 , 1823, 1416-25	4.9	214
129	Activity of metal-responsive transcription factor 1 by toxic heavy metals and H2O2 in vitro is modulated by metallothionein. <i>Molecular and Cellular Biology</i> , 2003 , 23, 8471-85	4.8	197
128	Transcriptional RenhancersRfrom SV40 and polyoma virus show a cell type preference. <i>Nucleic Acids Research</i> , 1982 , 10, 7965-76	20.1	187
127	Cloning, chromosomal mapping and characterization of the human metal-regulatory transcription factor MTF-1. <i>Nucleic Acids Research</i> , 1994 , 22, 3167-73	20.1	180
126	An enhancer stimulates transcription in trans when attached to the promoter via a protein bridge. <i>Cell</i> , 1989 , 58, 767-77	56.2	180

(2006-2013)

125	A recent evolutionary change affects a regulatory element in the human FOXP2 gene. <i>Molecular Biology and Evolution</i> , 2013 , 30, 844-52	8.3	163
124	Copper homeostasis in eukaryotes: teetering on a tightrope. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2006 , 1763, 737-46	4.9	162
123	Astrocytes and glioblastoma cells express novel octamer-DNA binding proteins distinct from the ubiquitous Oct-1 and B cell type Oct-2 proteins. <i>Nucleic Acids Research</i> , 1990 , 18, 5495-503	20.1	148
122	Functional domains of the heavy metal-responsive transcription regulator MTF-1. <i>Nucleic Acids Research</i> , 1995 , 23, 2277-86	20.1	146
121	RNA polymerase II C-terminal domain required for enhancer-driven transcription. <i>Nature</i> , 1995 , 374, 660-2	50.4	143
120	Enhancer sequences and the regulation of gene transcription. FEBS Journal, 1988, 176, 485-95		134
119	How do different transcription factors binding the same DNA sequence sort out their jobs?. <i>Trends in Genetics</i> , 1989 , 5, 37-9	8.5	131
118	Regulation of metallothionein transcription by the metal-responsive transcription factor MTF-1: identification of signal transduction cascades that control metal-inducible transcription. <i>Journal of Biological Chemistry</i> , 2002 , 277, 20438-45	5.4	129
117	The Drosophila homolog of mammalian zinc finger factor MTF-1 activates transcription in response to heavy metals. <i>Molecular and Cellular Biology</i> , 2001 , 21, 4505-14	4.8	126
116	Transcriptome response to heavy metal stress in Drosophila reveals a new zinc transporter that confers resistance to zinc. <i>Nucleic Acids Research</i> , 2006 , 34, 4866-77	20.1	122
115	Molecular analysis of the histone gene cluster of psammechinus miliaris: II. The arrangement of the five histone-coding and spacer sequences. <i>Cell</i> , 1976 , 8, 471-8	56.2	122
114	Metal-responsive transcription factor (MTF-1) handles both extremes, copper load and copper starvation, by activating different genes. <i>Genes and Development</i> , 2005 , 19, 891-6	12.6	121
113	B lymphocytes are impaired in mice lacking the transcriptional co-activator Bob1/OCA-B/OBF1. <i>European Journal of Immunology</i> , 1996 , 26, 3214-8	6.1	121
112	Eukaryotic expression vectors for the analysis of mutant proteins. <i>Nucleic Acids Research</i> , 1989 , 17, 641	820.1	118
111	Two major branches of anti-cadmium defense in the mouse: MTF-1/metallothioneins and glutathione. <i>Nucleic Acids Research</i> , 2005 , 33, 5715-27	20.1	116
110	During B-cell differentiation enhancer activity and transcription rate of immunoglobulin heavy chain genes are high before mRNA accumulation. <i>Cell</i> , 1986 , 45, 45-52	56.2	115
109	Knockout of Rmetal-responsive transcription factor RMTF-1 in Drosophila by homologous recombination reveals its central role in heavy metal homeostasis. <i>EMBO Journal</i> , 2003 , 22, 100-8	13	110
108	A family knockout of all four Drosophila metallothioneins reveals a central role in copper homeostasis and detoxification. <i>Molecular and Cellular Biology</i> , 2006 , 26, 2286-96	4.8	107

107	Nucleo-cytoplasmic trafficking of metal-regulatory transcription factor 1 is regulated by diverse stress signals. <i>Journal of Biological Chemistry</i> , 2001 , 276, 25487-95	5.4	105
106	Octamer transcription factors and the cell type-specificity of immunoglobulin gene expression. <i>FASEB Journal</i> , 1990 , 4, 1444-9	0.9	99
105	High frequency of homologous recombination in mammalian cells between endogenous and introduced SV40 genomes. <i>Cell</i> , 1985 , 43, 695-703	56.2	99
104	A novel expression selection approach allows precise mapping of the hepatitis B virus enhancer. <i>Nucleic Acids Research</i> , 1985 , 13, 7457-72	20.1	94
103	The four members of the Drosophila metallothionein family exhibit distinct yet overlapping roles in heavy metal homeostasis and detoxification. <i>Genes To Cells</i> , 2006 , 11, 647-58	2.3	88
102	Role of amyloid-beta glycine 33 in oligomerization, toxicity, and neuronal plasticity. <i>Journal of Neuroscience</i> , 2009 , 29, 7582-90	6.6	87
101	Promoters with the octamer DNA motif (ATGCAAAT) can be ubiquitous or cell type-specific depending on binding affinity of the octamer site and Oct-factor concentration. <i>Nucleic Acids Research</i> , 1991 , 19, 237-42	20.1	86
100	Simian virus 40 enhancer increases RNA polymerase density within the linked gene. <i>Nature</i> , 1985 , 315, 75-7	50.4	79
99	Molecular analysis of the histone gene cluster of Psammechinus miliaris: I. Fractionation and identification of five individual histone mRNAs. <i>Cell</i> , 1976 , 8, 455-69	56.2	79
98	Metal-responsive transcription factor-1 (MTF-1) is essential for embryonic liver development and heavy metal detoxification in the adult liver. <i>FASEB Journal</i> , 2004 , 18, 1071-9	0.9	78
97	cDNA cloning of human N-Oct3, a nervous-system specific POU domain transcription factor binding to the octamer DNA motif. <i>Nucleic Acids Research</i> , 1993 , 21, 253-8	20.1	76
96	Rapid test for in vivo stability and DNA binding of mutated octamer binding proteins with Rnini-extracts Rprepared from transfected cells. <i>Nucleic Acids Research</i> , 1989 , 17, 6420	20.1	73
95	NF-kappaB contributes to transcription of placenta growth factor and interacts with metal responsive transcription factor-1 in hypoxic human cells. <i>Biological Chemistry</i> , 2005 , 386, 865-72	4.5	69
94	Evidence for erosion of mouse CpG islands during mammalian evolution. <i>Somatic Cell and Molecular Genetics</i> , 1993 , 19, 543-55		68
93	Silencing of RNA polymerases II and III-dependent transcription by the KRAB protein domain of KOX1, a Krppel-type zinc finger factor. <i>Biological Chemistry</i> , 1997 , 378, 669-77	4.5	67
92	Tissue-specific expression of a FMR1/beta-galactosidase fusion gene in transgenic mice. <i>Human Molecular Genetics</i> , 1995 , 4, 359-66	5.6	66
91	Transcriptional enhancers can act in trans. <i>Trends in Genetics</i> , 1990 , 6, 300-4	8.5	66
90	DNA looping induced by a transcriptional enhancer in vivo. <i>Nucleic Acids Research</i> , 2005 , 33, 3743-50	20.1	64

[1996-1988]

89	Redundancy of information in enhancers as a principle of mammalian transcription control. <i>Journal of Molecular Biology</i> , 1988 , 201, 81-90	6.5	63	
88	Molecular analysis of the histone gene cluster of Psammechinus miliaris: III. Polarity and asymmetry of the histone-coding sequences. <i>Cell</i> , 1976 , 8, 479-84	56.2	62	
87	Copper homeostasis in Drosophila by complex interplay of import, storage and behavioral avoidance. <i>EMBO Journal</i> , 2007 , 26, 1035-44	13	61	
86	Open reading frame UL26 of human cytomegalovirus encodes a novel tegument protein that contains a strong transcriptional activation domain. <i>Journal of Virology</i> , 2002 , 76, 4836-47	6.6	59	
85	Partial denaturation mapping of cloned histone DNA from the sea urchin Psammechinus miliaris. <i>Nature</i> , 1976 , 264, 31-4	50.4	58	
84	Heavy metal ions in transcription factors from HeLa cells: Sp1, but not octamer transcription factor requires zinc for DNA binding and for activator function. <i>Nucleic Acids Research</i> , 1988 , 16, 5771-81	20.1	57	
83	Conservation of glutamine-rich transactivation function between yeast and humans. <i>Molecular and Cellular Biology</i> , 2000 , 20, 2774-82	4.8	53	
82	Homologous recombination and DNA-end joining reactions in zygotes and early embryos of zebrafish (Danio rerio) and Drosophila melanogaster. <i>Biological Chemistry</i> , 1998 , 379, 673-81	4.5	52	
81	Enhancers, enhancers - from their discovery to todayß universe of transcription enhancers. <i>Biological Chemistry</i> , 2015 , 396, 311-27	4.5	51	
80	Metal-responsive transcription factor (MTF-1) and heavy metal stress response in Drosophila and mammalian cells: a functional comparison. <i>Biological Chemistry</i> , 2004 , 385, 597-603	4.5	50	
79	Heat and heavy metal stress synergize to mediate transcriptional hyperactivation by metal-responsive transcription factor MTF-1. <i>Journal of Biological Chemistry</i> , 2003 , 278, 31879-83	5.4	47	
78	Direct interaction rescue, a novel filamentous phage technique to study protein-protein interactions. <i>Nucleic Acids Research</i> , 1994 , 22, 5761-2	20.1	46	
77	Evidence for transient requirement of the IgH enhancer. <i>Nucleic Acids Research</i> , 1985 , 13, 8901-12	20.1	45	
76	A novel cysteine cluster in human metal-responsive transcription factor 1 is required for heavy metal-induced transcriptional activation in vivo. <i>Journal of Biological Chemistry</i> , 2004 , 279, 4515-22	5.4	44	
75	Sp1 sites in the noncoding control region of BK polyomavirus are key regulators of bidirectional viral early and late gene expression. <i>Journal of Virology</i> , 2015 , 89, 3396-411	6.6	43	
74	Extended lifespan of Drosophila parkin mutants through sequestration of redox-active metals and enhancement of anti-oxidative pathways. <i>Neurobiology of Disease</i> , 2010 , 40, 82-92	7.5	43	
73	The nucleotide sequence and a first generation gene transfer vector of species B human adenovirus serotype 3. <i>Virology</i> , 2005 , 343, 283-98	3.6	43	
72	Two versatile eukaryotic vectors permitting epitope tagging, radiolabelling and nuclear localisation of expressed proteins. <i>Gene</i> , 1996 , 168, 165-7	3.8	43	

71	Upstream box/TATA box order is the major determinant of the direction of transcription. <i>Nucleic Acids Research</i> , 1991 , 19, 6699-704	20.1	43
70	The distal short consensus repeats 1 and 2 of the membrane cofactor protein CD46 and their distance from the cell membrane determine productive entry of species B adenovirus serotype 35. Journal of Virology, 2005 , 79, 10013-22	6.6	42
69	Synergistic activation of transcription by multiple binding sites for NF-kappa B even in absence of co-operative factor binding to DNA. <i>Journal of Molecular Biology</i> , 1990 , 214, 373-80	6.5	42
68	Toxicity of Alzheimerß disease-associated Alþeptide is ameliorated in a Drosophila model by tight control of zinc and copper availability. <i>Biological Chemistry</i> , 2011 , 392, 919-26	4.5	41
67	Activation of gene expression by metal-responsive signal transduction pathways. <i>Environmental Health Perspectives</i> , 2002 , 110 Suppl 5, 813-7	8.4	41
66	The CpG-specific methylase SssI has topoisomerase activity in the presence of Mg2+. <i>Nucleic Acids Research</i> , 1994 , 22, 5354-9	20.1	39
65	Copper sensing function of Drosophila metal-responsive transcription factor-1 is mediated by a tetranuclear Cu(I) cluster. <i>Nucleic Acids Research</i> , 2008 , 36, 3128-38	20.1	37
64	Metal-responsive transcription factor-1 (MTF-1) selects different types of metal response elements at low vs. high zinc concentration. <i>Biological Chemistry</i> , 2004 , 385, 623-32	4.5	37
63	Distorted copper homeostasis with decreased sensitivity to cisplatin upon chaperone Atox1 deletion in Drosophila. <i>BioMetals</i> , 2011 , 24, 445-53	3.4	36
62	Characterization of MtnE, the fifth metallothionein member in Drosophila. <i>Journal of Biological Inorganic Chemistry</i> , 2011 , 16, 1047-56	3.7	36
61	Transcription factors interacting with herpes simplex virus alpha gene promoters in sensory neurons. <i>Nucleic Acids Research</i> , 1995 , 23, 4978-85	20.1	36
60	The parkin mutant phenotype in the fly is largely rescued by metal-responsive transcription factor (MTF-1). <i>Molecular and Cellular Biology</i> , 2011 , 31, 2151-61	4.8	35
59	Polyglutamine tracts as modulators of transcriptional activation from yeast to mammals. <i>Biological Chemistry</i> , 2012 , 393, 63-70	4.5	33
58	Dramatic changes in the ratio of homologous recombination to nonhomologous DNA-end joining in oocytes and early embryos of Xenopus laevis. <i>Biological Chemistry Hoppe-Seyler</i> , 1996 , 377, 239-50		33
57	C-terminal domain (CTD) of RNA-polymerase II and N-terminal segment of the human TATA binding protein (TBP) can mediate remote and proximal transcriptional activation, respectively. <i>Nucleic Acids Research</i> , 1993 , 21, 5609-15	20.1	33
56	Metal-responsive transcription factor 1 (MTF-1) activity is regulated by a nonconventional nuclear localization signal and a metal-responsive transactivation domain. <i>Molecular and Cellular Biology</i> , 2009 , 29, 6283-93	4.8	31
55	Analysis of the heavy metal-responsive transcription factor MTF-1 from human and mouse. <i>Somatic Cell and Molecular Genetics</i> , 1995 , 21, 289-97		29
54	A conserved cysteine cluster, essential for transcriptional activity, mediates homodimerization of human metal-responsive transcription factor-1 (MTF-1). <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2012 , 1823, 476-83	4.9	28

53	Zinc supplement greatly improves the condition of parkin mutant Drosophila. <i>Biological Chemistry</i> , 2010 , 391, 513-8	4.5	28	
52	Positive and negative regulation at the herpes simplex virus ICP4 and ICP0 TAATGARAT motifs. <i>Virology</i> , 1995 , 207, 107-16	3.6	28	
51	Loss of metal transcription factor-1 suppresses tumor growth through enhanced matrix deposition. <i>FASEB Journal</i> , 2004 , 18, 1176-84	0.9	27	
50	Differential sensitivity of zinc finger transcription factors MTF-1, Sp1 and Krox-20 to CpG methylation of their binding sites. <i>Biological Chemistry Hoppe-Seyler</i> , 1996 , 377, 47-56		27	
49	A minimal transcription activation domain consisting of a specific array of aspartic acid and leucine residues. <i>Biological Chemistry Hoppe-Seyler</i> , 1994 , 375, 463-70		27	
48	Expression in mammalian cells of a cloned gene encoding murine DNA methyltransferase. <i>Gene</i> , 1991 , 109, 259-63	3.8	26	
47	An efficient method to generate chromosomal rearrangements by targeted DNA double-strand breaks in Drosophila melanogaster. <i>Genome Research</i> , 2004 , 14, 1382-93	9.7	24	
46	Complex demethylation patterns at Sp1 binding sites in F9 embryonal carcinoma cells. <i>FEBS Letters</i> , 1995 , 370, 170-4	3.8	24	
45	The Drosophila copper transporter Ctr1C functions in male fertility. <i>Journal of Biological Chemistry</i> , 2010 , 285, 17089-97	5.4	23	
44	Human copper transporter Ctr1 is functional in Drosophila, revealing a high degree of conservation between mammals and insects. <i>Journal of Biological Inorganic Chemistry</i> , 2010 , 15, 107-13	3.7	21	
43	A novel SR-related protein specifically interacts with the carboxy-terminal domain (CTD) of RNA polymerase II through a conserved interaction domain. <i>Biological Chemistry</i> , 1997 , 378, 565-71	4.5	21	
42	Predisposition to mouse thymic lymphomas in response to ionizing radiation depends on variant alleles encoding metal-responsive transcription factor-1 (Mtf-1). <i>Oncogene</i> , 2005 , 24, 399-406	9.2	21	
41	Pathological, physiological, and evolutionary aspects of short unstable DNA repeats in the human genome. <i>Biological Chemistry Hoppe-Seyler</i> , 1995 , 376, 201-11		20	
40	Two closely spaced promoters are equally activated by a remote enhancer: evidence against a scanning model for enhancer action. <i>Nucleic Acids Research</i> , 1989 , 17, 8931-47	20.1	20	
39	Dissection of Drosophila MTF-1 reveals a domain for differential target gene activation upon copper overload vs. copper starvation. <i>International Journal of Biochemistry and Cell Biology</i> , 2012 , 44, 404-11	5.6	18	
38	Use of the two-hybrid system and random sonicated DNA to identify the interaction domain of a protein. <i>BioTechniques</i> , 1996 , 21, 430-2	2.5	17	
37	Short introns interrupting the Oct-2 POU domain may prevent recombination between POU family genes without interfering with potential POU domain RhufflingRin evolution. <i>Biological Chemistry Hoppe-Seyler</i> , 1994 , 375, 675-83		17	
36	Specific transcriptional activation in vitro by the herpes simplex virus protein VP16. <i>Nucleic Acids Research</i> , 1993 , 21, 5570-6	20.1	16	

35	Characterization of the mouse gene for the heavy metal-responsive transcription factor MTF-1. <i>Cell Stress and Chaperones</i> , 2000 , 5, 196-206	4	16
34	Different potential of cellular and viral activators of transcription revealed in oocytes and early embryos of Xenopus laevis. <i>Biological Chemistry Hoppe-Seyler</i> , 1994 , 375, 105-12		13
33	Mercury and cadmium trigger expression of the copper importer Ctr1B, which enables Drosophila to thrive on heavy metal-loaded food. <i>Biological Chemistry</i> , 2009 , 390, 109-13	4.5	12
32	Drosophila bloom helicase maintains genome integrity by inhibiting recombination between divergent DNA sequences. <i>Nucleic Acids Research</i> , 2008 , 36, 6907-17	20.1	12
31	Conserved cysteine residues of Oct-2 POU domain confer sensitivity to oxidation but are dispensable for sequence-specific DNA binding. <i>Biochimica Et Biophysica Acta Gene Regulatory Mechanisms</i> , 1993 , 1173, 141-6		12
30	Long-range activation of transcription by SV40 enhancer is affected by "inhibitory" or "permissive" DNA sequences between enhancer and promoter. <i>Somatic Cell and Molecular Genetics</i> , 1989 , 15, 591-60)3	12
29	Dumpy-30 family members as determinants of male fertility and interaction partners of metal-responsive transcription factor 1 (MTF-1) in Drosophila. <i>BMC Developmental Biology</i> , 2008 , 8, 68	3.1	11
28	Characterization of metal-responsive transcription factor (MTF-1) from the giant rodent capybara reveals features in common with human as well as with small rodents (mouse, rat). Short communication. <i>Chemistry and Biodiversity</i> , 2008 , 5, 1485-94	2.5	10
27	Strong transcriptional activators isolated from viral DNA by the Pactivator trapRa novel selection system in mammalian cells. <i>Nucleic Acids Research</i> , 1994 , 22, 4031-8	20.1	10
26	In vitro transcription complementation assay with miniextracts of transiently transfected COS-1 cells. <i>Nucleic Acids Research</i> , 1992 , 20, 5855-6	20.1	10
25	Immediate early protein of pseudorabies virus is a general transactivator but stimulates only suboptimally utilized promoters. A clue to specificity?. <i>Journal of Molecular Biology</i> , 1990 , 215, 301-11	6.5	10
24	In vivo construction of transgenes in Drosophila. <i>Genetics</i> , 2007 , 175, 2019-28	4	9
23	Periodicity of eight nucleotides in purine distribution around human genomic CpG dinucleotides. <i>Somatic Cell and Molecular Genetics</i> , 1995 , 21, 91-8		8
22	The leucine zipper of c-Jun binds to ribosomal protein L18a: a role in Jun protein regulation?. <i>Biological Chemistry Hoppe-Seyler</i> , 1995 , 376, 321-5		8
21	Cloning and characterization of the murine B-cell specific transcriptional coactivator Bob1. <i>Biological Chemistry Hoppe-Seyler</i> , 1996 , 377, 139-45		8
20	Tissue-specific gene expression. <i>Trends in Neurosciences</i> , 1985 , 8, 100-104	13.3	8
19	Improved "activator trap" method for the isolation of transcriptional activation domains from random DNA fragments. <i>BioTechniques</i> , 1996 , 21, 848-54	2.5	7
18	A transcriptional terminator between enhancer and promoter does not affect remote transcriptional control. <i>Somatic Cell and Molecular Genetics</i> , 1990 , 16, 351-60		7

LIST OF PUBLICATIONS

17	Transcriptional Regulation by Heavy Metals, Exemplified at the Metallothionein Genes 1995 , 206-240		7
16	Transcriptional activation by recombinant GAL4-VP16 in the Xenopus oocyte. <i>Nucleic Acids Research</i> , 1993 , 21, 2775	20.1	6
15	POU-specific domain of Oct-2 factor confers RoctamerRmotif DNA binding specificity on heterologous Antennapedia homeodomain. <i>FEBS Letters</i> , 1992 , 314, 361-5	3.8	6
14	Fine mapping of protein interaction surfaces with a PCR-based mutagenesis screen in yeast. <i>Trends in Genetics</i> , 1996 , 12, 393-4	8.5	5
13	A factor known to bind to endogenous Ig heavy chain enhancer only in lymphocytes is a ubiquitously active transcription factor. <i>FEBS Journal</i> , 1990 , 187, 507-13		4
12	Transcription enhancers as major determinants of SV40 polyomavirus growth efficiency and host cell tropism. <i>Journal of General Virology</i> , 2016 , 97, 1597-1603	4.9	4
11	The legless lizard Anguis fragilis (slow worm) has a potent metal-responsive transcription factor 1 (MTF-1). <i>Biological Chemistry</i> , 2014 , 395, 425-31	4.5	2
10	Simian virus 40 strains with novel properties generated by replacing the viral enhancer with synthetic oligonucleotides. <i>Journal of Virology</i> , 2012 , 86, 3135-42	6.6	2
9	Short-lived mammals (shrew, mouse) have a less robust metal-responsive transcription factor than humans and bats. <i>BioMetals</i> , 2016 , 29, 423-32	3.4	2
8	Liver degeneration and embryonic lethality in mouse null mutants for the metal-responsive transcriptional activator MTF-1 1999 , 223-226		1
7	Transient Expression of Cloned Genes in Mammalian Cells 1983 , 19-32		1
6	2 Regulation of Metallothionein Gene Expression 2015 , 31-50		
5	BZLF1 (ZEBRA, Zta) Protein of Epstein-Barr Virus Selected in a Yeast One-Hybrid System by Binding to a Consensus Site in the IgH Intronic Enhancer: A Role in Immunoglobulin Expression?. <i>Biological Chemistry Hoppe-Seyler</i> , 1996 , 377, 669-674		
4	Wie werden unsere Gene ein- und ausgeschaltet? 2000 , 7-14		
3	Inducible and constitutive sequence elements in the enhancer of the mouse metallothionein-I gene. <i>Exs</i> , 1987 , 52, 415-22		
2	Cloning of Sequence-Specific DNA-Binding Proteins by Screening &DNA Expression Libraries with Radiolabelled Binding-Site Probes 1991 , 233-244		
1	Embryonic Liver Degeneration and Increased Sensitivity Towards Heavy Metal and H2O2 in Mice Lacking the Metal-Responsive Transcription Factor MTF-1 1999 , 339-352		