

Tbx1 haploinsufficiency in the DiGeorge syndrome region mice

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Isolation and Characterization of a Novel Gene from the DiGeorge Chromosomal Region That Encodes for a Mediator Subunit. <i>Genomics</i> , 2001, 74, 320-332.	1.3	33
2	Introducing Defined Chromosomal Rearrangements into the Mouse Genome. <i>Methods</i> , 2001, 24, 81-94.	1.9	30
3	A common cis-acting sequence in the DiGeorge critical region regulates bi-directional transcription of UFD1L and CDC45L. <i>Mechanisms of Development</i> , 2001, 108, 81-92.	1.7	9
4	Cloning and characterization of the gene encoding human NPL4, a protein interacting with the ubiquitin fusion-degradation protein (UFD1L). <i>Gene</i> , 2001, 275, 39-46.	1.0	11
6	Subnuclear localization and mitotic phosphorylation of HIRA, the human homologue of <i>Saccharomyces cerevisiae</i> transcriptional regulators Hir1p/Hir2p. <i>Biochemical Journal</i> , 2001, 358, 447.	1.7	14
7	Subnuclear localization and mitotic phosphorylation of HIRA, the human homologue of <i>Saccharomyces cerevisiae</i> transcriptional regulators Hir1p/Hir2p. <i>Biochemical Journal</i> , 2001, 358, 447-455.	1.7	18
8	Benign idiopathic partial seizures in the velocardiofacial syndrome: Report of two cases. <i>American Journal of Medical Genetics Part A</i> , 2001, 103, 172-175.	2.4	13
9	The T-box transcription factor gene TBX22 is mutated in X-linked cleft palate and ankyloglossia. <i>Nature Genetics</i> , 2001, 29, 179-183.	9.4	245
10	Getting the T-box dose right. <i>Nature Medicine</i> , 2001, 7, 1185-1186.	15.2	20
11	Genomic organisation of the ≈ 1.5 Mb Smith-Magenis syndrome critical interval: Transcription map, genomic contig, and candidate gene analysis. <i>European Journal of Human Genetics</i> , 2001, 9, 892-902.	1.4	30
13	Genetics of craniofacial development and malformation. <i>Nature Reviews Genetics</i> , 2001, 2, 458-468.	7.7	380
14	Engineering chromosomal rearrangements in mice. <i>Nature Reviews Genetics</i> , 2001, 2, 780-790.	7.7	229
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16	From mouse to man: generating megabase chromosome rearrangements. <i>Trends in Genetics</i> , 2001, 17, 331-339.	2.9	88
17	Causes of the phenotype-genotype dissociation in DiGeorge syndrome: clues from mouse models. <i>Trends in Genetics</i> , 2001, 17, 551-554.	2.9	8
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19	A region of homozygosity within 22q11.2 associated with congenital heart disease: recessive DiGeorge/velocardiofacial syndrome?. <i>Journal of Medical Genetics</i> , 2001, 38, 533-536.	1.5	8
20	High frequency of the ApoB-100 R3500Q mutation in Bulgarian hypercholesterolaemic subjects. <i>Journal of Medical Genetics</i> , 2001, 38, 536-540.	1.5	12

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21	Mutation analysis of TBX1 in non-deleted patients with features of DGS/VCFS or isolated cardiovascular defects. <i>Journal of Medical Genetics</i> , 2001, 38, 45e-45.	1.5	135
22	Mice deleted for the DiGeorge/velocardiofacial syndrome region show abnormal sensorimotor gating and learning and memory impairments. <i>Human Molecular Genetics</i> , 2001, 10, 2645-2650.	1.4	140
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