Kenia B El-Jaick

List of Publications by Year
in descending order

19	825	840119 11	839053 18
papers	citations	h-index	g-index

19	19	19	1106
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Homozygotes NAT2*5B slow acetylators are highly associated with hepatotoxicity induced by anti-tuberculosis drugs. Memorias Do Instituto Oswaldo Cruz, 2022, 117, e210328.	0.8	0
2	PGAM1 and TP53 mRNA levels in canine mammary carcinomas â€“ Short communication. Acta Veterinaria Hungarica, 2021, 69, 50-54.	0.2	4
3	Could polymorphisms in ABCB1 gene represent a genetic risk factor for the development of mammary tumors in dogs?. Veterinary Journal, 2019, 248, 58-63.	0.6	1
4	TP53 gene expression levels and tumor aggressiveness in canine mammary carcinomas. Journal of Veterinary Diagnostic Investigation, 2017, 29, 865-868.	0.5	9
5	Molecular analysis of holoprosencephaly in South America. Genetics and Molecular Biology, 2014, 37, 250-262.	0.6	8
6	The role of cigarette smoking and liver enzymes polymorphisms in anti-tuberculosis drug-induced hepatotoxicity in Brazilian patients. Tuberculosis, 2014, 94, 299-305.	0.8	30
7	Unique CYP3A4 genetic variant in Brazilian tuberculosis patients with/without HIV. Molecular Medicine Reports, 2012, 5, 153-61.	1.1	3
8	The roles ofGSTM1andGSTT1null genotypes and other predictors in anti-tuberculosis drug-induced liver injury. Journal of Clinical Pharmacy and Therapeutics, 2012, 37, 712-718.	0.7	27
9	TGIF Mutations in Human Holoprosencephaly: Correlation between Genotype and Phenotype. Molecular Syndromology, 2010, 1, 211-222.	0.3	19
10	Clinical spectrum of SIX3-associated mutations in holoprosencephaly: correlation between genotype, phenotype and function. Journal of Medical Genetics, 2009, 46, 389-398.	1.5	88
11	The mutational spectrum of holoprosencephaly-associated changes within theSHHgene in humans predicts loss-of-function through either key structural alterations of the ligand or its altered synthesis. Human Mutation, 2009, 30, E921-E935.	1.1	77
12	Detection of mutations inGATA1gene using automated denaturing high-performance liquid chromatography and direct sequencing in children with Down syndrome. Leukemia and Lymphoma, 2009, 50, 834-840.	0.6	2
13	Regulation of a remote Shh forebrain enhancer by the Six3 homeoprotein. Nature Genetics, 2008, 40, 1348-1353.	9.4	182
14	Mutations in the human SIX3 gene in holoprosencephaly are loss of function. Human Molecular Genetics, 2008, 17, 3919-3928.	1.4	56
15	No association of the polyhistidine tract polymorphism of the ZIC2 gene with neural tube defects in a South American (ECLAMC) population. Molecular Medicine Reports, 2008, , .	1.1	1
16	Functional analysis of mutations in TGIF associated with holoprosencephaly. Molecular Genetics and Metabolism, 2007, 90, 97-111.	0.5	63
17	Single median maxillary central incisor: New data and mutation review. Birth Defects Research Part A: Clinical and Molecular Teratology, 2007, 79, 573-580.	1.6	32
18	A functional screen for sonic hedgehog regulatory elements across a 1 Mb interval identifies long-range ventral forebrain enhancers. Development (Cambridge), 2006, 133, 761-772.	1.2	198

#	Article	IF	Citations
19	SIX3 mutations with holoprosencephaly. American Journal of Medical Genetics, Part A, 2006, 140A, 2577-2583.	0.7	25