

Multiplex ARMS analysis to detect 13 common mutations

Clinical Genetics

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Citation Report

#	ARTICLE	IF	CITATIONS
1	PCSK9: un exemple de recherche translationnelle. <i>Medecine Des Maladies Metaboliques</i> , 2008, 2, 10-14.	0.1	0
2	Reductions in all-cause, cancer, and coronary mortality in statin-treated patients with heterozygous familial hypercholesterolaemia: a prospective registry study. <i>European Heart Journal</i> , 2008, 29, 2625-2633.	2.2	391
3	Identifying patients with familial hypercholesterolaemia in primary care. <i>Heart</i> , 2008, 94, 695-696.	2.9	8
4	What is the clinical utility of DNA testing in patients with familial hypercholesterolaemia?. <i>Current Opinion in Lipidology</i> , 2008, 19, 362-368.	2.7	62
5	Identification of loci conferring risk for premature CAD and heterozygous familial hyperlipidemia in the LDLR, APOB and PCSK9 genes. <i>International Journal of Diabetes Mellitus</i> , 2009, 1, 16-21.	0.6	5
6	Evaluation of high-resolution melting analysis for screening the LDL receptor gene. <i>Clinical Biochemistry</i> , 2009, 42, 528-535.	1.9	18
7	Multiplex ligation-dependent probe amplification analysis to screen for deletions and duplications of the LDLR gene in patients with familial hypercholesterolaemia. <i>Clinical Genetics</i> , 2009, 76, 69-75.	2.0	29
8	Mutation screening in patients for familial hypercholesterolaemia (ADH). <i>Clinical Genetics</i> , 2010, 77, 97-99.	2.0	18
9	Mutation detection rate and spectrum in familial hypercholesterolaemia patients in the UK pilot cascade project. <i>Clinical Genetics</i> , 2010, 77, 572-580.	2.0	155
10	Genetic screening for homozygous and heterozygous familial hypercholesterolemia. <i>The Application of Clinical Genetics</i> , 2010, 3, 147.	3.0	5
11	A double heterozygote for familial hypercholesterolaemia and familial defective apolipoprotein B-100. <i>Annals of Clinical Biochemistry</i> , 2010, 47, 487-490.	1.6	16
12	Development of a high-resolution melting method for mutation detection in familial hypercholesterolaemia patients. <i>Annals of Clinical Biochemistry</i> , 2010, 47, 44-55.	1.6	38
13	Familial hypercholesterolemia: epidemiology, Neolithic origins and modern geographic distribution. <i>Critical Reviews in Clinical Laboratory Sciences</i> , 2011, 48, 1-18.	6.1	57
14	Familial hypercholesterolaemia: A model of care for Australasia. <i>Atherosclerosis Supplements</i> , 2011, 12, 221-263.	1.2	181
15	Mechanisms and genetic determinants regulating sterol absorption, circulating LDL levels, and sterol elimination: implications for classification and disease risk. <i>Journal of Lipid Research</i> , 2011, 52, 1885-1926.	4.2	76
16	A DNA Microarray for the Detection of Point Mutations and Copy Number Variation Causing Familial Hypercholesterolemia in Europe. <i>Journal of Molecular Diagnostics</i> , 2013, 15, 362-372.	2.8	17
17	The use of next-generation sequencing in clinical diagnosis of familial hypercholesterolemia. <i>Genetics in Medicine</i> , 2013, 15, 948-957.	2.4	69
18	Multiplex ARMS PCR to Detect 8 Common Mutations of ATP7B Gene in Patients With Wilson Disease. <i>Hepatitis Monthly</i> , 2013, 13, e8375.	0.2	6

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19	Molecular characterization of a Chinese woman homozygous for the familial hypercholesterolemiaLDLRc.1474G>A (p.Asp492Asn) mutation. <i>Clinical Lipidology</i> , 2014, 9, 163-170.	0.4	1
20	Screening for Familial Hypercholesterolaemia: Universal or Cascade? A Critique of Current FH Recognition Strategies. <i>Current Cardiovascular Risk Reports</i> , 2015, 9, 1.	2.0	3
21	Mutation p.L799R in the LDLR, which affects the transmembrane domain of the LDLR, prevents membrane insertion and causes secretion of the mutant LDLR. <i>Human Molecular Genetics</i> , 2015, 24, 5836-5844.	2.9	15
22	Childâ€Parent Familial Hypercholesterolemia Screening in Primary Care. <i>New England Journal of Medicine</i> , 2016, 375, 1628-1637.	27.0	250
23	PCSK9 Variants in Familial Hypercholesterolemia: A Comprehensive Synopsis. <i>Frontiers in Genetics</i> , 2020, 11, 1020.	2.3	29
24	Genetics of Cardiovascular Diseases. , 2009, , 281-293.		2
26	Elucigene FH20 and LIPOchip for the diagnosis of familial hypercholesterolaemia: a systematic review and economic evaluation.. <i>Health Technology Assessment</i> , 2012, 16, 1-266.	2.8	15
27	Prioritization of Candidate Nonsynonymous Single Nucleotide Polymorphisms via Sequence Conservation Features. <i>International Journal of Engineering and Manufacturing</i> , 2011, 1, 66-72.	0.7	1
28	Genetic testing for familial hypercholesterolemiaâ€past, present, and future. <i>Journal of Lipid Research</i> , 2021, 62, 100139.	4.2	20
29	cSNP Identification and Genotyping from C4B and BAT2 Assigned to the SLA Class III Region. <i>Journal of Animal Science and Technology</i> , 2007, 49, 549-558.	2.5	0
30	Cloning, cSNP Identification, and Genotyping of Pig Complement Factor B(CFB) Gene Located on the SLA Class III Region. <i>Journal of Animal Science and Technology</i> , 2008, 50, 753-762.	2.5	0
31	Extraction of Sequence Conservation Features for the Prioritization of Candidate Single Amino Acid Polymorphisms. <i>International Journal of Information Engineering and Electronic Business</i> , 2011, 3, 1-10.	1.2	3
32	A 72-Year-Old Patient with Longstanding, Untreated Familial Hypercholesterolemia but no Coronary Artery Calcification: A Case Report. <i>Cureus</i> , 2018, 10, e2452.	0.5	1