

Epigenetics of Cardiovascular Disease: A New 'Beat' i

Medical Epigenetics

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Mechanisms Mediating Environmental Chemical-Induced Endocrine Disruption in the Adrenal Gland. <i>Frontiers in Endocrinology</i> , 2015, 6, 29.	1.5	46
2	Cell-free DNA for diagnosing myocardial infarction: not ready for prime time. <i>Clinical Chemistry and Laboratory Medicine</i> , 2015, 53, 1895-901.	1.4	12
3	Epigenetic Regulation of Angiogenesis by JARID1B-Induced Repression of HOXA5. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2015, 35, 1645-1652.	1.1	33
5	Regulation of Nox enzymes expression in vascular pathophysiology: Focusing on transcription factors and epigenetic mechanisms. <i>Redox Biology</i> , 2015, 5, 358-366.	3.9	96
6	Endothelial Epigenetics in Biomechanical Stress. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2015, 35, 1317-1326.	1.1	57
7	Epigenetics in the Vascular Endothelium. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2015, 35, 2297-2306.	1.1	48
8	Association of interleukin-6 methylation in leukocyte DNA with serum level and the risk of ischemic heart disease. <i>Scandinavian Journal of Clinical and Laboratory Investigation</i> , 2016, 76, 291-295.	0.6	11
9	Circulating long noncoding RNAs as novel biomarkers of human diseases. <i>Biomarkers in Medicine</i> , 2016, 10, 757-769.	0.6	31
10	Association between global leukocyte DNA methylation and cardiovascular risk in postmenopausal women. <i>BMC Medical Genetics</i> , 2016, 17, 71.	2.1	26
11	Epigenetic associations in relation to cardiovascular prevention and therapeutics. <i>Clinical Epigenetics</i> , 2016, 8, 4.	1.8	62
12	Gene variants at FTO, 9p21, and 2q36.3 are age-independently associated with myocardial infarction in Czech men. <i>Clinica Chimica Acta</i> , 2016, 454, 119-123.	0.5	15
13	Long noncoding RNA variations in cardiometabolic diseases. <i>Journal of Human Genetics</i> , 2017, 62, 97-104.	1.1	40
14	Epigenetics in Cardiovascular Disease. , 2017, , 135-157.		3
15	Epigenetics-by-Sex Interaction for Coronary Artery Disease Risk Conferred by the Cystathionine β -Lyase Gene Promoter Methylation. <i>OMICS A Journal of Integrative Biology</i> , 2017, 21, 741-748.	1.0	19
16	Genome-wide DNA methylome alterations in acute coronary syndrome. <i>International Journal of Molecular Medicine</i> , 2017, 41, 220-232.	1.8	11
17	Exercise Training and Epigenetic Regulation: Multilevel Modification and Regulation of Gene Expression. <i>Advances in Experimental Medicine and Biology</i> , 2017, 1000, 281-322.	0.8	29
18	Biofluids, cell mechanics and epigenetics: Flow-induced epigenetic mechanisms of endothelial gene expression. <i>Journal of Biomechanics</i> , 2017, 50, 3-10.	0.9	12
19	miRNAs may change rapidly with thoughts: The Relaxation Response after myocardial infarction. <i>European Journal of Integrative Medicine</i> , 2018, 20, 63-72.	0.8	10

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20	Epigenetic Regulation of Endothelial Function: With Focus on MicroRNAs. , 2018, , 171-187.		0
21	Short-Term Exposure to High Sucrose Levels near Weaning Has a Similar Long-Lasting Effect on Hypertension as a Long-Term Exposure in Rats. <i>Nutrients</i> , 2018, 10, 728.	1.7	13
22	Genetic Markers for Coronary Artery Disease. <i>Medicina (Lithuania)</i> , 2018, 54, 36.	0.8	9
23	Integrating Genes Affecting Coronary Artery Disease in Functional Networks by Multi-OMICs Approach. <i>Frontiers in Cardiovascular Medicine</i> , 2018, 5, 89.	1.1	23
24	Epigenetic Determinants of Flow-Mediated Vascular Endothelial Gene Expression. <i>Hypertension</i> , 2019, 74, 467-476.	1.3	19
25	DNA methylation of antisense noncoding RNA in the INK locus (ANRIL) is associated with coronary artery disease in a Chinese population. <i>Scientific Reports</i> , 2019, 9, 15340.	1.6	8
26	Effect of Sucrose Ingestion at the End of a Critical Window that Increases Hypertension Susceptibility on Peripheral Mechanisms Regulating Blood Pressure in Rats. Role of Sirtuins 1 and 3. <i>Nutrients</i> , 2019, 11, 309.	1.7	8
27	TETs Regulate Proepicardial Cell Migration through Extracellular Matrix Organization during Zebrafish Cardiogenesis. <i>Cell Reports</i> , 2019, 26, 720-732.e4.	2.9	22
28	Genome-wide DNA Methylation Profiling of Blood from Monozygotic Twins Discordant for Myocardial Infarction. <i>In Vivo</i> , 2020, 34, 361-367.	0.6	8
29	Epigenetic-sensitive pathways in personalized therapy of major cardiovascular diseases. , 2020, 210, 107514.		87
30	LncRNA 0003250 accelerates heart autophagy and binds to miRâ€17â€5p as a competitive endogenous RNA in chicken induced by selenium deficiency. <i>Journal of Cellular Physiology</i> , 2021, 236, 157-177.	2.0	34
31	The Histone Demethylase PHF8 Is Essential for Endothelial Cell Migration. <i>PLoS ONE</i> , 2016, 11, e0146645.	1.1	27
32	Genomics era and complex disorders. <i>Journal of Postgraduate Medicine</i> , 2016, 62, 188-198.	0.2	14
33	Study of the expression of genes associated with post-translational changes in histones in the internal thoracic artery and the saphenous vein grafts used in coronary artery bypass grafting procedure. <i>Medical Journal of Cell Biology (discontinued)</i> , 2020, 8, 183-189.	0.2	1
34	Dissecting the functional pleiotropism of lysine demethylase 5B in physiology and pathology. <i>Journal of Cancer Research and Practice</i> , 2020, 7, 49.	0.2	0
35	Polygenic risk for coronary artery disease in the Scottish and English population. <i>BMC Cardiovascular Disorders</i> , 2021, 21, 586.	0.7	6
36	Epigenetics and Gut Microbiota Crosstalk: A potential Factor in Pathogenesis of Cardiovascular Disorders. <i>Bioengineering</i> , 2022, 9, 798.	1.6	1