Romain Barres

List of Publications by Citations

Source: https://exaly.com/author-pdf/2470493/romain-barres-publications-by-citations.pdf

Version: 2024-04-27

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.

75
papers

5,323
citations

49
h-index
g-index

88
ext. papers

6,532
ext. citations

9
avg, IF
L-index

#	Paper	IF	Citations
75	Chronic high-fat diet in fathers programs Etell dysfunction in female rat offspring. <i>Nature</i> , 2010 , 467, 963-6	50.4	1043
74	Acute exercise remodels promoter methylation in human skeletal muscle. <i>Cell Metabolism</i> , 2012 , 15, 405-11	24.6	586
73	Non-CpG methylation of the PGC-1alpha promoter through DNMT3B controls mitochondrial density. <i>Cell Metabolism</i> , 2009 , 10, 189-98	24.6	463
72	Skeletal muscle PGC-1¶ modulates kynurenine metabolism and mediates resilience to stress-induced depression. <i>Cell</i> , 2014 , 159, 33-45	56.2	453
71	Obesity and Bariatric Surgery Drive Epigenetic Variation of Spermatozoa in Humans. <i>Cell Metabolism</i> , 2016 , 23, 369-78	24.6	320
70	High-fat diet reprograms the epigenome of rat spermatozoa and transgenerationally affects metabolism of the offspring. <i>Molecular Metabolism</i> , 2016 , 5, 184-197	8.8	217
69	Weight loss after gastric bypass surgery in human obesity remodels promoter methylation. <i>Cell Reports</i> , 2013 , 3, 1020-7	10.6	192
68	Sperm epigenetics and influence of environmental factors. <i>Molecular Metabolism</i> , 2018 , 14, 1-11	8.8	134
67	In Situ Fixation Redefines Quiescence and Early Activation of Skeletal Muscle Stem Cells. <i>Cell Reports</i> , 2017 , 21, 1982-1993	10.6	133
66	ZBED6, a novel transcription factor derived from a domesticated DNA transposon regulates IGF2 expression and muscle growth. <i>PLoS Biology</i> , 2009 , 7, e1000256	9.7	124
65	The role of diet and exercise in the transgenerational epigenetic landscape of T2DM. <i>Nature Reviews Endocrinology</i> , 2016 , 12, 441-51	15.2	119
64	DNA methylation in metabolic disorders. American Journal of Clinical Nutrition, 2011, 93, 897S-900	7	111
63	Time of Exercise Specifies the Impact on Muscle Metabolic Pathways and Systemic Energy Homeostasis. <i>Cell Metabolism</i> , 2019 , 30, 92-110.e4	24.6	88
62	Evidence Suggesting Absence of Mitochondrial DNA Methylation. Frontiers in Genetics, 2017, 8, 166	4.5	85
61	Altered DNA methylation of glycolytic and lipogenic genes in liver from obese and type 2 diabetic patients. <i>Molecular Metabolism</i> , 2016 , 5, 171-183	8.8	74
60	The Microbiota and Epigenetic Regulation of T Helper 17/Regulatory T Cells: In Search of a Balanced Immune System. <i>Frontiers in Immunology</i> , 2017 , 8, 417	8.4	72
59	DNA methylation is altered in B and NK lymphocytes in obese and type 2 diabetic human. <i>Metabolism: Clinical and Experimental</i> , 2014 , 63, 1188-97	12.7	70

(2006-2016)

58	The Emerging Role of Epigenetics in Inflammation and Immunometabolism. <i>Trends in Endocrinology and Metabolism</i> , 2016 , 27, 782-795	8.8	66	
57	Fatty acid-induced insulin resistance: role of insulin receptor substrate 1 serine phosphorylation in the retroregulation of insulin signalling. <i>Biochemical Society Transactions</i> , 2003 , 31, 1152-6	5.1	62	
56	Hyperosmotic stress inhibits insulin receptor substrate-1 function by distinct mechanisms in 3T3-L1 adipocytes. <i>Journal of Biological Chemistry</i> , 2003 , 278, 26550-7	5.4	58	
55	Endurance training remodels sperm-borne small RNA expression and methylation at neurological gene hotspots. <i>Clinical Epigenetics</i> , 2018 , 10, 12	7.7	55	
54	Nicotinamide riboside does not alter mitochondrial respiration, content or morphology in skeletal muscle from obese and insulin-resistant men. <i>Journal of Physiology</i> , 2020 , 598, 731-754	3.9	53	
53	Altered promoter methylation of PDK4, IL1 B, IL6, and TNF after Roux-en Y gastric bypass. <i>Surgery for Obesity and Related Diseases</i> , 2014 , 10, 671-8	3	52	
52	Epigenetic changes in healthy human skeletal muscle following exercise- a systematic review. <i>Epigenetics</i> , 2019 , 14, 633-648	5.7	51	
51	Evidence for non-CpG methylation in mammals. Experimental Cell Research, 2011, 317, 2555-61	4.2	46	
50	Exercise training alters the genomic response to acute exercise in human adipose tissue. <i>Epigenomics</i> , 2018 , 10, 1033-1050	4.4	41	
49	Alteration in insulin action: role of IRS-1 serine phosphorylation in the retroregulation of insulin signalling. <i>Annales DrEndocrinologie</i> , 2004 , 65, 43-8	1.7	38	
48	Transcriptomic and epigenetic responses to short-term nutrient-exercise stress in humans. <i>Scientific Reports</i> , 2017 , 7, 15134	4.9	33	
47	Role of adenosine 5Whonophosphate-activated protein kinase in interleukin-6 release from isolated mouse skeletal muscle. <i>Endocrinology</i> , 2009 , 150, 600-6	4.8	31	
46	Altered expression and insulin-induced trafficking of Na+-K+-ATPase in rat skeletal muscle: effects of high-fat diet and exercise. <i>American Journal of Physiology - Endocrinology and Metabolism</i> , 2009 , 297, E38-49	6	28	
45	C-peptide increases Na,K-ATPase expression via PKC- and MAP kinase-dependent activation of transcription factor ZEB in human renal tubular cells. <i>PLoS ONE</i> , 2011 , 6, e28294	3.7	27	
44	Altered Methylation Profile of Lymphocytes Is Concordant with Perturbation of Lipids Metabolism and Inflammatory Response in Obesity. <i>Journal of Diabetes Research</i> , 2016 , 2016, 8539057	3.9	26	
43	Dynamic epigenetic responses to muscle contraction. <i>Drug Discovery Today</i> , 2014 , 19, 1010-4	8.8	25	
42	Ionizing Radiation Potentiates High-Fat Diet-Induced Insulin Resistance and Reprograms Skeletal Muscle and Adipose Progenitor Cells. <i>Diabetes</i> , 2016 , 65, 3573-3584	0.9	25	
41	Enigma interacts with adaptor protein with PH and SH2 domains to control insulin-induced actin cytoskeleton remodeling and glucose transporter 4 translocation. <i>Molecular Endocrinology</i> , 2006 , 20, 2864-75		24	

40	Time-restricted feeding alters lipid and amino acid metabolite rhythmicity without perturbing clock gene expression. <i>Nature Communications</i> , 2020 , 11, 4643	17.4	22
39	Muscle Contraction Induces Acute Hydroxymethylation of the Exercise-Responsive Gene. <i>Frontiers in Endocrinology</i> , 2016 , 7, 165	5.7	22
38	Preadipocytes from obese humans with type 2 diabetes are epigenetically reprogrammed at genes controlling adipose tissue function. <i>International Journal of Obesity</i> , 2019 , 43, 306-318	5.5	22
37	The interaction between the adaptor protein APS and Enigma is involved in actin organisation. <i>Experimental Cell Research</i> , 2005 , 308, 334-44	4.2	21
36	T cell epigenetic remodeling and accelerated epigenetic aging are linked to long-term immune alterations in childhood cancer survivors. <i>Clinical Epigenetics</i> , 2018 , 10, 138	7.7	19
35	Chronic erythropoietin treatment improves diet-induced glucose intolerance in rats. <i>Journal of Endocrinology</i> , 2015 , 225, 77-88	4.7	14
34	Skeletal muscle enhancer interactions identify genes controlling whole-body metabolism. <i>Nature Communications</i> , 2020 , 11, 2695	17.4	14
33	Atlas of exercise metabolism reveals time-dependent signatures of metabolic homeostasis <i>Cell Metabolism</i> , 2022 ,	24.6	14
32	Strenuous physical exercise adversely affects monocyte chemotaxis. <i>Thrombosis and Haemostasis</i> , 2011 , 105, 122-30	7	13
31	Epigenetic Reprogramming of Immune Cells in Women With PCOS Impact Genes Controlling Reproductive Function. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2019 , 104, 6155-6170	5.6	12
30	Paternal high-fat diet transgenerationally impacts hepatic immunometabolism. <i>FASEB Journal</i> , 2019 , 33, 6269-6280	0.9	10
29	Transcriptomic and epigenomics atlas of myotubes reveals insight into the circadian control of metabolism and development. <i>Epigenomics</i> , 2020 , 12, 701-713	4.4	10
28	Grandpaternal-induced transgenerational dietary reprogramming of the unfolded protein response in skeletal muscle. <i>Molecular Metabolism</i> , 2017 , 6, 621-630	8.8	10
27	Identification of two microRNA nodes as potential cooperative modulators of liver metabolism. <i>Hepatology Research</i> , 2019 , 49, 1451-1465	5.1	8
26	Afadin is a scaffold protein repressing insulin action via HDAC6 in adipose tissue. <i>EMBO Reports</i> , 2019 , 20, e48216	6.5	8
25	Nampt controls skeletal muscle development by maintaining Ca homeostasis and mitochondrial integrity. <i>Molecular Metabolism</i> , 2021 , 53, 101271	8.8	7
24	Genes controlling the activation of natural killer lymphocytes are epigenetically remodeled in intestinal cells from germ-free mice. <i>FASEB Journal</i> , 2019 , 33, 2719-2731	0.9	6
23	Methodology for Accurate Detection of Mitochondrial DNA Methylation. <i>Journal of Visualized Experiments</i> , 2018 ,	1.6	6

(2021-2020)

22	OBEDIS Core Variables Project: European Expert Guidelines on a Minimal Core Set of Variables to Include in Randomized, Controlled Clinical Trials of Obesity Interventions. <i>Obesity Facts</i> , 2020 , 13, 1-28	5.1	5
21	Nutritional status affects the epigenomic profile of peripheral blood cells. <i>Epigenomics</i> , 2011 , 3, 259-60	4.4	5
20	Contraction influences Per2 gene expression in skeletal muscle through a calcium-dependent pathway. <i>Journal of Physiology</i> , 2020 , 598, 5739-5752	3.9	5
19	Environmental factors influence the epigenetic signature of newborns from mothers with gestational diabetes. <i>Epigenomics</i> , 2019 , 11, 861-873	4.4	4
18	Epigenetic and Transcriptomic Characterization of Pure Adipocyte Fractions From Obese Pigs Identifies Candidate Pathways Controlling Metabolism. <i>Frontiers in Genetics</i> , 2019 , 10, 1268	4.5	4
17	Ablation of DNA-methyltransferase 3A in skeletal muscle does not affect energy metabolism or exercise capacity. <i>PLoS Genetics</i> , 2021 , 17, e1009325	6	4
16	Insulin-induced serine 22 phosphorylation of retinoid X receptor alpha is dispensable for adipogenesis in brown adipocytes. <i>Adipocyte</i> , 2020 , 9, 142-152	3.2	3
15	Disrupted circadian oscillations in type 2 diabetes are linked to altered rhythmic mitochondrial metabolism in skeletal muscle. <i>Science Advances</i> , 2021 , 7, eabi9654	14.3	3
14	GREM1 is epigenetically reprogrammed in muscle cells after exercise training and controls myogenesis and metabolism		3
13	Perinatal exposure to nicotine alters spermatozoal DNA methylation near genes controlling nicotine action. <i>FASEB Journal</i> , 2021 , 35, e21702	0.9	3
12	Exercise during pregnancy mitigates negative effects of parental obesity on metabolic function in adult mouse offspring. <i>Journal of Applied Physiology</i> , 2021 , 130, 605-616	3.7	3
11	Structured supervised exercise training or motivational counselling during pregnancy on physical activity level and health of mother and offspring: FitMum study protocol. <i>BMJ Open</i> , 2021 , 11, e043671	3	3
10	Comparative analysis of oral and intraperitoneal glucose tolerance tests in mice <i>Molecular Metabolism</i> , 2022 , 57, 101440	8.8	2
9	Endurance exercise training-responsive miR-19b-3p improves skeletal muscle glucose metabolism. <i>Nature Communications</i> , 2021 , 12, 5948	17.4	2
8	White adipose remodeling during browning in mice involves YBX1 to drive thermogenic commitment. <i>Molecular Metabolism</i> , 2021 , 44, 101137	8.8	2
7	Cold-induction of afadin in brown fat supports its thermogenic capacity. Scientific Reports, 2021, 11, 979	94 .9	1
6	Circadian Transcriptomic and Epigenomic Remodeling in Response to Lipid Overload and Human Obesit	Ξy	1
5	Epigenetic rewiring of skeletal muscle enhancers after exercise training supports a role in whole-body function and human health. <i>Molecular Metabolism</i> , 2021 , 53, 101290	8.8	1

4	Translational Medicine, 2021 , 13, eabc7367	17.5	1
3	Non-cell autonomous mechanisms control mitochondrial gene dysregulation in polycystic ovary syndrome. <i>Journal of Molecular Endocrinology</i> , 2021 , 68, 63-76	4.5	0
2	Time trends in epigenetic signatures and population health risks 2021 , 285-298		
1	DNA Methylation and Gene Expression in Blood and Adipose Tissue of Adult Offspring of Women with Diabetes in Pregnancy Validation Study of DNA Methylation Changes Identified in Adolescent Offspring. <i>Biomedicines.</i> 2022 . 10, 1244	4.8	