## Luca Lambertini

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A Research Strategy to Discover the Environmental Causes of Autism and Neurodevelopmental Disabilities. Environmental Health Perspectives, 2012, 120, a258-60.	6.0	191
2	First Experimental Demonstration of the Multipotential Carcinogenic Effects of Aspartame Administered in the Feed to Sprague-Dawley Rats. Environmental Health Perspectives, 2006, 114, 379-385.	6.0	190
3	Differential expression of imprinted genes in normal and IUGR human placentas. Epigenetics, 2009, 4, 235-240.	2.7	176
4	Global Methylation in the Placenta and Umbilical Cord Blood From Pregnancies With Maternal Gestational Diabetes, Preeclampsia, and Obesity. Reproductive Sciences, 2014, 21, 131-137.	2.5	138
5	Effect of postnatal low-dose exposure to environmental chemicals on the gut microbiome in a rodent model. Microbiome, 2016, 4, 26.	11.1	122
6	Results of Longâ€Term Experimental Studies on the Carcinogenicity of Ethyleneâ€bisâ€Dithiocarbamate (Mancozeb) in Rats. Annals of the New York Academy of Sciences, 2002, 982, 123-136.	3.8	96
7	Exploring the associations between microRNA expression profiles and environmental pollutants in human placenta from the National Children's Study (NCS). Epigenetics, 2015, 10, 793-802.	2.7	91
8	Placental expression profile of imprinted genes impacts birth weight. Epigenetics, 2015, 10, 842-849.	2.7	79
9	Cadmium-Associated Differential Methylation throughout the Placental Genome: Epigenome-Wide Association Study of Two U.S. Birth Cohorts. Environmental Health Perspectives, 2018, 126, 017010.	6.0	69
10	Insights into beta cell regeneration for diabetes via integration of molecular landscapes in human insulinomas. Nature Communications, 2017, 8, 767.	12.8	67
11	Stable heteroplasmy at the single-cell level is facilitated by intercellular exchange of mtDNA. Nucleic Acids Research, 2015, 43, 2177-2187.	14.5	62
12	Expression of imprinted genes in placenta is associated with infant neurobehavioral development. Epigenetics, 2015, 10, 834-841.	2.7	59
13	Expression quantitative trait loci (eQTLs) in human placentas suggest developmental origins of complex diseases. Human Molecular Genetics, 2017, 26, 3432-3441.	2.9	58
14	A sensitive functional assay reveals frequent loss of genomic imprinting in human placenta. Epigenetics, 2008, 3, 261-269.	2.7	54
15	Whole-transcriptome analysis delineates the human placenta gene network and its associations with fetal growth. BMC Genomics, 2017, 18, 520.	2.8	53
16	Neurodevelopmental consequences in offspring of mothers with preeclampsia during pregnancy: underlying biological mechanism via imprinting genes. Archives of Gynecology and Obstetrics, 2017, 295, 1319-1329.	1.7	50
17	Imprinted gene expression in fetal growth and development. Placenta, 2012, 33, 480-486.	1.5	49
18	Intrauterine multi-metal exposure is associated with reduced fetal growth through modulation of the placental gene network. Environment International, 2018, 120, 373-381.	10.0	46

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19	<i>In utero</i> exposures to environmental organic pollutants disrupt epigenetic marks linked to fetoplacental development. Environmental Epigenetics, 2016, 2, dvv013.	1.8	44
20	Mitochondrial Epigenetics and Environmental Exposure. Current Environmental Health Reports, 2016, 3, 214-224.	6.7	42
21	Placental imprinting variation associated with assisted reproductive technologies and subfertility. Epigenetics, 2017, 12, 653-661.	2.7	42
22	Genomic loss of imprinting in first-trimester human placenta. American Journal of Obstetrics and Gynecology, 2010, 202, 391.e1-391.e8.	1.3	38
23	Differential Methylation of Imprinted Genes in Growth-Restricted Placentas. Reproductive Sciences, 2011, 18, 1111-1117.	2.5	38
24	Genetic regulation of the placental transcriptome underlies birth weight and risk of childhood obesity. PLoS Genetics, 2018, 14, e1007799.	3.5	38
25	Placenta-Imprinted Gene Expression Association of Infant Neurobehavior. Journal of Pediatrics, 2012, 160, 854-860.e2.	1.8	36
26	Epigenetics in Women's Health Care. Mount Sinai Journal of Medicine, 2010, 77, 225-235.	1.9	35
27	Intrauterine Reprogramming of the Polycystic Ovary Syndrome: Evidence from a Pilot Study of Cord Blood Global Methylation Analysis. Frontiers in Endocrinology, 2017, 8, 352.	3.5	35
28	Mitochondrial Gene Expression Profiles Are Associated with Maternal Psychosocial Stress in Pregnancy and Infant Temperament. PLoS ONE, 2015, 10, e0138929.	2.5	35
29	Environmental Influences on Genomic Imprinting. Current Environmental Health Reports, 2015, 2, 155-162.	6.7	33
30	Effect of maternal exposure to endocrine disrupting chemicals on reproduction and mammary gland development in female Sprague-Dawley rats. Reproductive Toxicology, 2015, 54, 110-119.	2.9	31
31	Maternal residential air pollution and placental imprinted gene expression. Environment International, 2017, 108, 204-211.	10.0	26
32	Changes in mammary histology and transcriptome profiles by low-dose exposure to environmental phenols at critical windows of development. Environmental Research, 2017, 152, 233-243.	7.5	26
33	Timing of prenatal exposure to trauma and altered placental expressions of hypothalamicâ€pituitaryâ€ødrenal axis genes and genes driving neurodevelopment. Journal of Neuroendocrinology, 2018, 30, e12581.	2.6	24
34	Placental Expression of Imprinted Genes, Overall and in Sex-Specific Patterns, Associated with Placental Cadmium Concentrations and Birth Size. Environmental Health Perspectives, 2019, 127, 57005.	6.0	24
35	Myc Is Required for Adaptive β-Cell Replication in Young Mice but Is Not Sufficient in One-Year-Old Mice Fed With a High-Fat Diet. Diabetes, 2019, 68, 1934-1949.	0.6	23
36	Influences of Maternal Stress during Pregnancy on the Epi/genome: Comparison of Placenta and Umbilical Cord Blood. Journal of Depression & Anxiety, 2014, 03, .	0.1	22

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37	The atpIBEXF operon coding for the F 0 sector of the ATP synthase from the purple nonsulfur photosynthetic bacterium Rhodobacter capsulatus. Archives of Microbiology, 1998, 170, 385-388.	2.2	21
38	Results of a Long-Term Carcinogenicity Bioassay on Sprague-Dawley Rats Exposed to Sodium Arsenite Administered in Drinking Water. Annals of the New York Academy of Sciences, 2006, 1076, 578-591.	3.8	20
39	Paired Serum and Urine Concentrations of Biomarkers of Diethyl Phthalate, Methyl Paraben, and Triclosan in Rats. Environmental Health Perspectives, 2016, 124, 39-45.	6.0	18
40	The many lives of Myc in the pancreatic $\hat{l}^2$ -cell. Journal of Biological Chemistry, 2021, 296, 100122.	3.4	16
41	Intrauterine Growth Restriction Is Associated with Unique Features of the Reproductive Microbiome. Reproductive Sciences, 2021, 28, 828-837.	2.5	16
42	Prenatal exposure to maternal depression and anxiety on imprinted gene expression in placenta and infant neurodevelopment and growth. Pediatric Research, 2018, 83, 1075-1083.	2.3	15
43	Disrupting the DREAM complex enables proliferation of adult human pancreatic $\hat{l}^2$ cells. Journal of Clinical Investigation, 2022, 132, .	8.2	14
44	Genomic imprinting. Current Opinion in Pediatrics, 2014, 26, 237-242.	2.0	13
45	Mitochondrial gene expression profiles are associated with intrahepatic cholestasis of pregnancy. Placenta, 2016, 45, 16-23.	1.5	11
46	In-depth characterization of the placental imprintome reveals novel differentially methylated regions across birth weight categories. Epigenetics, 2020, 15, 47-60.	2.7	11
47	Copper associates with differential methylation in placentae from two US birth cohorts. Epigenetics, 2020, 15, 215-230.	2.7	11
48	Analysis of p53 Tumor Suppressor Gene, H-ras Protooncogene and Proliferating Cell Nuclear Antigen (PCNA) in Squamous Cell Carcinomas of HRA/Skh Mice Following Exposure to 8-Methoxypsoralen (8-MOP) and UVA Radiation (PUVA Therapy). Toxicologic Pathology, 2005, 33, 292-299.	1.8	9
49	Placental imprinted gene expression mediates the effects of maternal psychosocial stress during pregnancy on fetal growth. Journal of Developmental Origins of Health and Disease, 2019, 10, 196-205.	1.4	9
50	Aberrant methylation underlies insulin gene expression in human insulinoma. Nature Communications, 2020, 11, 5210.	12.8	9
51	Mitochondrial and glycolysis-regulatory gene expression profiles are associated with intrauterine growth restriction. Journal of Maternal-Fetal and Neonatal Medicine, 2018, 33, 1-10.	1.5	8
52	Moderate prenatal stress may buffer the impact of Superstorm Sandy on placental genes: Stress in Pregnancy (SIP) Study. PLoS ONE, 2020, 15, e0226605.	2.5	7
53	Demonstration of all-or-none loss of imprinting in mRNA expression in single cells. Nucleic Acids Research, 2009, 37, 7039-7046.	14.5	6
54	Genomic Imprinting in Human Placenta. , 2012, , .		5

Genomic Imprinting in Human Placenta. , 2012, , . 54

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55	441: Differential mitochondrial DNA methylation in growth restricted placentas. American Journal of Obstetrics and Gynecology, 2013, 208, S192.	1.3	3
56	465: Placental mitochondrial DNA mutation rate: a new association with intrauterine growth restriction (IUGR). American Journal of Obstetrics and Gynecology, 2016, 214, S255-S256.	1.3	2
57	631: The impact of postpartum hemorrhage drill training at a single institution. American Journal of Obstetrics and Gynecology, 2012, 206, S283.	1.3	1
58	308: Global methylation in placenta and umbilical cord blood from pregnancies with metabolic syndromes and the effect on birth outcomes. American Journal of Obstetrics and Gynecology, 2014, 210, S161.	1.3	1
59	Identification of endocrine disrupting chemical doses in rats to reproduce human urinary metabolite concentrations. ISEE Conference Abstracts, 2013, 2013, 5107.	0.0	1
60	38: Loss of imprinting in first trimester human placentas. American Journal of Obstetrics and Gynecology, 2009, 201, S22-S23.	1.3	0
61	712: The effects of exposure to BPA and BBP on the DNA methylation profile of the IGF2/H19 imprinting control region in HTR-8 cells. American Journal of Obstetrics and Gynecology, 2011, 204, S281.	1.3	0
62	705: Polymorphisms of the glucocorticoid receptor gene NR3C1 and the association with birth weight and gestational age at delivery. American Journal of Obstetrics and Gynecology, 2012, 206, S314.	1.3	0
63	193: Mitochondrial gene expression in intrahepatic cholestasis of pregnancy. American Journal of Obstetrics and Gynecology, 2015, 212, S111.	1.3	0
64	Intrauterine programming of polycystic ovary syndrome: evidence from cord blood global methylation analysis. Fertility and Sterility, 2017, 108, e248.	1.0	0
65	Developing Functional Assays for Genomic Imprinting Profile in Human Placenta. Epidemiology, 2007, 18, S138-S139.	2.7	Ο
66	Correlations of Urinary and Amniotic Fluid Phthalate Metabolite Concentrations and Expression of Imprinted Genes in Human Placenta. Epidemiology, 2009, 20, S110-S111.	2.7	0
67	Prenatal Phenol Exposure and Expression of Imprinted Genes in Human Placenta. Epidemiology, 2009, 20, S107.	2.7	Ο
68	Abstract LB-089: Defining windows of susceptibility for low-dose exposure to endocrine disruptors in rat mammary development by microRNA profiling. , 2015, , .		0