## Hiroaki Okae

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3132576/publications.pdf

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37 2,563 21 papers citations h-index

40 40 40 4659 all docs docs citations times ranked citing authors

34

g-index

#	Article	IF	CITATIONS
1	Derivation of Human Trophoblast Stem Cells. Cell Stem Cell, 2018, 22, 50-63.e6.	11.1	570
2	The International Human Epigenome Consortium: A Blueprint for Scientific Collaboration and Discovery. Cell, 2016, 167, 1145-1149.	28.9	404
3	Genome-Wide Analysis of DNA Methylation Dynamics during Early Human Development. PLoS Genetics, 2014, 10, e1004868.	3.5	216
4	Naive Human Embryonic Stem Cells Can Give Rise to Cells with a Trophoblast-like Transcriptome and Methylome. Stem Cell Reports, 2020, 15, 198-213.	4.8	129
5	Characterization of DNA methylation errors in patients with imprinting disorders conceived by assisted reproduction technologies. Human Reproduction, 2012, 27, 2541-2548.	0.9	122
6	Induction of Human Trophoblast Stem Cells from Somatic Cells and Pluripotent Stem Cells. Cell Reports, 2020, 33, 108419.	6.4	117
7	Association of four imprinting disorders and ART. Clinical Epigenetics, 2019, 11, 21.	4.1	115
8	Allele-Specific Methylome and Transcriptome Analysis Reveals Widespread Imprinting in the Human Placenta. American Journal of Human Genetics, 2016, 99, 1045-1058.	6.2	103
9	Re-investigation and RNA sequencing-based identification of genes with placenta-specific imprinted expression. Human Molecular Genetics, 2012, 21, 548-558.	2.9	102
10	Imprinting methylation errors in ART. Reproductive Medicine and Biology, 2014, 13, 193-202.	2.4	66
11	Assessing loss of imprint methylation in sperm from subfertile men using novel methylation polymerase chain reaction Luminex analysis. Fertility and Sterility, 2011, 95, 129-134.e4.	1.0	60
12	RNA sequencing-based identification of aberrant imprinting in cloned mice. Human Molecular Genetics, 2014, 23, 992-1001.	2.9	57
13	Neural tube defects and impaired neural progenitor cell proliferation in <i>Gβ1</i> €deficient mice. Developmental Dynamics, 2010, 239, 1089-1101.	1.8	55
14	ASCL2 reciprocally controls key trophoblast lineage decisions during hemochorial placenta development. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	53
15	Genome-wide microRNA expression profiling in placentae from frozen-thawed blastocyst transfer. Clinical Epigenetics, 2017, 9, 79.	4.1	51
16	Epigenetic alterations in sperm associated with male infertility. Congenital Anomalies (discontinued), 2015, 55, 133-144.	0.6	49
17	Factors associated with aberrant imprint methylation and oligozoospermia. Scientific Reports, 2017, 7, 42336.	3.3	37
18	Stability of genomic imprinting in human induced pluripotent stem cells. BMC Genetics, 2013, 14, 32.	2.7	31

#	Article	IF	CITATIONS
19	<scp>DNA</scp> methylation errors in imprinting disorders and assisted reproductive technology. Pediatrics International, 2013, 55, 542-549.	0.5	30
20	Histone deacetylase 1 and 2 drive differentiation and fusion of progenitor cells in human placental trophoblasts. Cell Death and Disease, 2020, 11, 311.	6.3	30
21	Loss of p57 <sup>KIP2</sup> expression confers resistance to contact inhibition in human androgenetic trophoblast stem cells. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 26606-26613.	7.1	24
22	Transcriptomic signature of the follicular somatic compartment surrounding an oocyte with high developmental competence. Scientific Reports, 2017, 7, 6815.	3.3	22
23	The microRNA cluster C19MC confers differentiation potential into trophoblast lineages upon human pluripotent stem cells. Nature Communications, 2022, 13, .	12.8	20
24	Intersection of regulatory pathways controlling hemostasis and hemochorial placentation. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	19
25	Cutting Edge: G Protein Subunit $\hat{l}^2$ 1 Negatively Regulates NLRP3 Inflammasome Activation. Journal of Immunology, 2019, 202, 1942-1947.	0.8	15
26	Lactate dehydrogenase C is required for the protein expression of a sperm-specific isoform of lactate dehydrogenase A. Journal of Biochemistry, 2019, 165, 323-334.	1.7	15
27	Atypical protein kinase C iota (PKCλ/ι) ensures mammalian development by establishing the maternal–fetal exchange interface. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 14280-14291.	7.1	14
28	Genome-Scale Assessment of Age-Related DNA Methylation Changes in Mouse Spermatozoa. PLoS ONE, 2016, 11, e0167127.	2.5	14
29	Unique features and emerging in vitro models of human placental development. Reproductive Medicine and Biology, 2020, 19, 301-313.	2.4	9
30	High-throughput detection of aberrant imprint methylation in the ovarian cancer by the bisulphite PCR-Luminex method. BMC Medical Genomics, 2012, 5, 8.	1.5	4
31	Abnormal cleavage is involved in the self-correction of bovine preimplantation embryos. Biochemical and Biophysical Research Communications, 2021, 562, 76-82.	2.1	4
32	DNA Methylation Dynamics During Early Human Development. Journal of Mammalian Ova Research, 2016, 33, 101-107.	0.1	3
33	Therapeutic Approaches to Imprinting Diseases. , 2018, , 861-875.		1
34	G protein subunit $\hat{I}^21$ is an important mediator of the late stage of endochondral ossification. Biochemical and Biophysical Research Communications, 2020, 533, 90-96.	2.1	1
35	SS6-4 IL-17A and IL-17F are important for the development of intestinal polyps in APCmin mice by accelerating blood vessel formation. Cytokine, 2010, 52, 46.	<b>3.</b> 2	0
36	Incomplete reprogramming of germline DNA methylation in the human placenta. Placenta, 2016, 45, 113.	1.5	0

# ARTICLE IF CITATIONS

37 Epigenetic Alterations in Human Sperm., 2019, , 1161-1176. 0