

# Joshua F Robinson

## List of Publications by Year in descending order

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46  
papers

1,415  
citations

304743

22  
h-index

345221

36  
g-index

47  
all docs

47  
docs citations

47  
times ranked

1880  
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrated analysis of transcriptomic datasets to identify placental biomarkers of spontaneous preterm birth. <i>Placenta</i> , 2022, 122, 66-73.	1.5	0
2	Neutralizing antibody activity against SARS-CoV-2 variants in gestational age-matched mother-infant dyads after infection or vaccination. <i>JCI Insight</i> , 2022, 7, .	5.0	13
3	Organophosphate Flame Retardants, Highly Fluorinated Chemicals, and Biomarkers of Placental Development and Disease During Mid-Gestation. <i>Toxicological Sciences</i> , 2021, 181, 215-228.	3.1	22
4	Human placental cytotrophoblast epigenome dynamics over gestation and alterations in placental disease. <i>Developmental Cell</i> , 2021, 56, 1238-1252.e5.	7.0	29
5	Global proteomic analyses of human cytotrophoblast differentiation/invasion. <i>Development (Cambridge)</i> , 2021, 148, .	2.5	5
6	Cytotrophoblast extracellular vesicles enhance decidual cell secretion of immune modulators via TNF-alpha. <i>Development (Cambridge)</i> , 2020, 147, .	2.5	12
7	Differences in cytochrome p450 enzyme expression and activity in fetal and adult tissues. <i>Placenta</i> , 2020, 100, 35-44.	1.5	26
8	Racial/ethnic and geographic differences in polybrominated diphenyl ether (PBDE) levels across maternal, placental, and fetal tissues during mid-gestation. <i>Scientific Reports</i> , 2020, 10, 12247.	3.3	22
9	Up-regulated cytotrophoblast DOCK4 contributes to over-invasion in placenta accreta spectrum. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 15852-15861.	7.1	19
10	Association of polybrominated diphenyl ether (PBDE) levels with biomarkers of placental development and disease during mid-gestation. <i>Environmental Health</i> , 2020, 19, 61.	4.0	13
11	Retinoids and developmental neurotoxicity: Utilizing toxicogenomics to enhance adverse outcome pathways and testing strategies. <i>Reproductive Toxicology</i> , 2020, 96, 102-113.	2.9	8
12	Trisomy 21 is Associated with Caspase-2 Upregulation in Cytotrophoblasts at the Maternal-Fetal Interface. <i>Reproductive Sciences</i> , 2020, 27, 100-109.	2.5	0
13	Circulating Monocytes, Tissue Macrophages, and Malaria. <i>Journal of Tropical Medicine</i> , 2019, 2019, 1-9.	1.7	21
14	Using a Multi-Stage hESC Model to Characterize BDE-47 Toxicity During Neurogenesis. <i>Toxicological Sciences</i> , 2019, 171, 221-234.	3.1	20
15	Genomic Profiling of BDE-47 Effects on Human Placental Cytotrophoblasts. <i>Toxicological Sciences</i> , 2019, 167, 211-226.	3.1	32
16	Polybrominated diphenyl ethers (PBDEs) and hydroxylated PBDE metabolites (OH-PBDEs) in maternal and fetal tissues, and associations with fetal cytochrome P450 gene expression. <i>Environment International</i> , 2018, 112, 269-278.	10.0	66
17	Convergence of placenta biology and genetic risk for schizophrenia. <i>Nature Medicine</i> , 2018, 24, 792-801.	30.7	214
18	Transcriptional Dynamics of Cultured Human Villous Cytotrophoblasts. <i>Endocrinology</i> , 2017, 158, 1581-1594.	2.8	25

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19	A genomics-based framework for identifying biomarkers of human neurodevelopmental toxicity. <i>Reproductive Toxicology</i> , 2016, 60, 1-10.	2.9	8
20	Rbpj links uterine transformation and embryo orientation. <i>Cell Research</i> , 2014, 24, 1031-1032.	12.0	3
21	Valproic acid-induced gene expression responses in rat whole embryo culture and comparison across in vitro developmental and non-developmental models. <i>Reproductive Toxicology</i> , 2013, 41, 57-66.	2.9	13
22	Dose response analysis of monophthalates in the murine embryonic stem cell test assessed by cardiomyocyte differentiation and gene expression. <i>Reproductive Toxicology</i> , 2013, 35, 81-88.	2.9	27
23	Toxicogenomic Approaches in Developmental Toxicology Testing. <i>Methods in Molecular Biology</i> , 2013, 947, 451-473.	0.9	16
24	Complementary Detection of Embryotoxic Properties of Substances in the Neural and Cardiac Embryonic Stem Cell Tests. <i>Toxicological Sciences</i> , 2013, 132, 118-130.	3.1	37
25	Transcriptomic Analysis of Neurulation and Early Organogenesis in Rat Embryos: An In Vivo and Ex Vivo Comparison. <i>Toxicological Sciences</i> , 2012, 126, 255-266.	3.1	18
26	A Comparison of Gene Expression Responses in Rat Whole Embryo Culture and In Vivo: Time-Dependent Retinoic Acid-Induced Teratogenic Response. <i>Toxicological Sciences</i> , 2012, 126, 242-254.	3.1	34
27	Compound-specific effects of diverse neurodevelopmental toxicants on global gene expression in the neural embryonic stem cell test (ESTn). <i>Toxicology and Applied Pharmacology</i> , 2012, 262, 330-340.	2.8	38
28	Transcriptomic Concentration-Response Evaluation of Valproic Acid, Cyproconazole, and Hexaconazole in the Neural Embryonic Stem Cell Test (ESTn). <i>Toxicological Sciences</i> , 2012, 125, 430-438.	3.1	55
29	Dose-response analysis of phthalate effects on gene expression in rat whole embryo culture. <i>Toxicology and Applied Pharmacology</i> , 2012, 264, 32-41.	2.8	22
30	Triazole induced concentration-related gene signatures in rat whole embryo culture. <i>Reproductive Toxicology</i> , 2012, 34, 275-283.	2.9	47
31	A Review of Toxicogenomic Approaches in Developmental Toxicology. <i>Methods in Molecular Biology</i> , 2012, 889, 347-371.	0.9	18
32	Discriminating classes of developmental toxicants using gene expression profiling in the embryonic stem cell test. <i>Toxicology Letters</i> , 2011, 201, 143-151.	0.8	56
33	Arsenic- and cadmium-induced toxicogenomic response in mouse embryos undergoing neurulation. <i>Toxicology and Applied Pharmacology</i> , 2011, 250, 117-129.	2.8	45
34	Comparison of MeHg-induced toxicogenomic responses across in vivo and in vitro models used in developmental toxicology. <i>Reproductive Toxicology</i> , 2011, 32, 180-188.	2.9	35
35	Gene set assembly for quantitative prediction of developmental toxicity in the embryonic stem cell test. <i>Toxicology</i> , 2011, 284, 63-71.	4.2	33
36	Cadmium Induced p53-Dependent Activation of Stress Signaling, Accumulation of Ubiquitinated Proteins, and Apoptosis in Mouse Embryonic Fibroblast Cells. <i>Toxicological Sciences</i> , 2011, 120, 403-412.	3.1	32

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37	Time-Response Evaluation by Transcriptomics of Methylmercury Effects on Neural Differentiation of Murine Embryonic Stem Cells. <i>Toxicological Sciences</i> , 2011, 122, 437-447.	3.1	67
38	Embryonic toxicokinetic and dynamic differences underlying strain sensitivity to cadmium during neurulation. <i>Reproductive Toxicology</i> , 2010, 29, 279-285.	2.9	12
39	Methylmercury induced toxicogenomic response in C57 and SWV mouse embryos undergoing neural tube closure. <i>Reproductive Toxicology</i> , 2010, 30, 284-291.	2.9	30
40	Toxicogenomic profiling in maternal and fetal rodent brains following gestational exposure to chlorpyrifos. <i>Toxicology and Applied Pharmacology</i> , 2010, 245, 310-325.	2.8	40
41	Integrating genetic and toxicogenomic information for determining underlying susceptibility to developmental disorders. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2010, 88, 920-930.	1.6	10
42	A systems-based approach to investigate dose- and time-dependent methylmercury-induced gene expression response in C57BL/6 mouse embryos undergoing neurulation. <i>Birth Defects Research Part B: Developmental and Reproductive Toxicology</i> , 2010, 89, 188-200.	1.4	13
43	A System-Based Comparison of Gene Expression Reveals Alterations in Oxidative Stress, Disruption of Ubiquitin-Proteasome System and Altered Cell Cycle Regulation after Exposure to Cadmium and Methylmercury in Mouse Embryonic Fibroblast. <i>Toxicological Sciences</i> , 2010, 114, 356-377.	3.1	49
44	Embryotoxicant-Specific Transcriptomic Responses in Rat Postimplantation Whole-Embryo Culture. <i>Toxicological Sciences</i> , 2010, 118, 675-685.	3.1	38
45	Cadmium-Induced Differential Toxicogenomic Response in Resistant and Sensitive Mouse Strains Undergoing Neurulation. <i>Toxicological Sciences</i> , 2009, 107, 206-219.	3.1	44
46	Gene expression profiling analysis reveals arsenic-induced cell cycle arrest and apoptosis in p53-proficient and p53-deficient cells through differential gene pathways. <i>Toxicology and Applied Pharmacology</i> , 2008, 233, 389-403.	2.8	28