

# Michelle R Jones

## List of Publications by Year in descending order

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Version: 2024-02-01

21  
papers

3,851  
citations

567281

15  
h-index

677142

22  
g-index

26  
all docs

26  
docs citations

26  
times ranked

9093  
citing authors

#	ARTICLE	IF	CITATIONS
1	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 349-362.	2.8	23
2	A Simple Optimization Workflow to Enable Precise and Accurate Imputation of Missing Values in Proteomic Data Sets. <i>Journal of Proteome Research</i> , 2021, 20, 3214-3229.	3.7	14
3	Large-scale cross-cancer fine-mapping of the 5p15.33 region reveals multiple independent signals. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100041.	1.7	6
4	Ovarian Cancer Risk Variants Are Enriched in Histotype-Specific Enhancers and Disrupt Transcription Factor Binding Sites. <i>American Journal of Human Genetics</i> , 2020, 107, 622-635.	6.2	14
5	GENAVi: a shiny web application for gene expression normalization, analysis and visualization. <i>BMC Genomics</i> , 2019, 20, 745.	2.8	40
6	Large-scale genome-wide meta-analysis of polycystic ovary syndrome suggests shared genetic architecture for different diagnosis criteria. <i>PLoS Genetics</i> , 2018, 14, e1007813.	3.5	341
7	Genetic epidemiology of ovarian cancer and prospects for polygenic risk prediction. <i>Gynecologic Oncology</i> , 2017, 147, 705-713.	1.4	69
8	Genetic determinants of polycystic ovary syndrome: progress and future directions. <i>Fertility and Sterility</i> , 2016, 106, 25-32.	1.0	103
9	Association study of androgen signaling pathway genes in polycystic ovary syndrome. <i>Fertility and Sterility</i> , 2016, 105, 467-473.e4.	1.0	11
10	Comprehensive assessment of expression of insulin signaling pathway components in subcutaneous adipose tissue of women with and without polycystic ovary syndrome. <i>Journal of Clinical and Translational Endocrinology</i> , 2015, 2, 99-104.	1.4	9
11	Further Investigation in Europeans of Susceptibility Variants for Polycystic Ovary Syndrome Discovered in Genome-Wide Association Studies of Chinese Individuals. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2015, 100, E182-E186.	3.6	57
12	Systems Genetics Reveals the Functional Context of PCOS Loci and Identifies Genetic and Molecular Mechanisms of Disease Heterogeneity. <i>PLoS Genetics</i> , 2015, 11, e1005455.	3.5	84
13	Discovery and refinement of loci associated with lipid levels. <i>Nature Genetics</i> , 2013, 45, 1274-1283.	21.4	2,641
14	Steroidogenic Regulatory Factor <i>FOS</i> Is Underexpressed in Polycystic Ovary Syndrome (PCOS) Adipose Tissue and Genetically Associated with PCOS Susceptibility. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2012, 97, E1750-E1757.	3.6	22
15	Metabolic and cardiovascular genes in polycystic ovary syndrome: A candidate-wide association study (CWAS). <i>Steroids</i> , 2012, 77, 317-322.	1.8	23
16	Replication of association of <i>DENND1A</i> and <i>THADA</i> variants with polycystic ovary syndrome in European cohorts. <i>Journal of Medical Genetics</i> , 2012, 49, 90-95.	3.2	165
17	FTO and MC4R Gene Variants Are Associated with Obesity in Polycystic Ovary Syndrome. <i>PLoS ONE</i> , 2011, 6, e16390.	2.5	92
18	Replication of association of a novel insulin receptor gene polymorphism with polycystic ovary syndrome. <i>Fertility and Sterility</i> , 2011, 95, 1736-1741.e11.	1.0	55

#	ARTICLE	IF	CITATIONS
19	Type 2 diabetes susceptibility single-nucleotide polymorphisms are not associated with polycystic ovary syndrome. <i>Fertility and Sterility</i> , 2011, 95, 2538-2541.e6.	1.0	31
20	Harnessing Expression Data to Identify Novel Candidate Genes in Polycystic Ovary Syndrome. <i>PLoS ONE</i> , 2011, 6, e20120.	2.5	12
21	Independent Confirmation of Association between Metabolic Phenotypes of Polycystic Ovary Syndrome and Variation in the Type 6 17 $\beta$ -Hydroxysteroid Dehydrogenase Gene. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2009, 94, 5034-5038.	3.6	32