

# *BRCA2* in Ovarian Development and Function

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Rare variants in FANCA induce premature ovarian insufficiency. <i>Human Genetics</i> , 2019, 138, 1227-1236.	1.8	56
2	BRCA2 deficiency is a potential driver for human primary ovarian insufficiency. <i>Cell Death and Disease</i> , 2019, 10, 474.	2.7	28
3	A novel EIF4ENIF1 mutation associated with a diminished ovarian reserve and premature ovarian insufficiency identified by whole-exome sequencing. <i>Journal of Ovarian Research</i> , 2019, 12, 119.	1.3	27
4	A TOP6BL mutation abolishes meiotic DNA double-strand break formation and causes human infertility. <i>Science Bulletin</i> , 2020, 65, 2120-2129.	4.3	18
5	Variants in Homologous Recombination Genes <i>EXO1</i> and <i>RAD51</i> Related with Premature Ovarian Insufficiency. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2020, 105, e3566-e3574.	1.8	21
6	MicroRNA-127-5p impairs function of granulosa cells via <i>HMGB2</i> gene in premature ovarian insufficiency. <i>Journal of Cellular Physiology</i> , 2020, 235, 8826-8838.	2.0	24
7	Novel pathogenic mutations in minichromosome maintenance complex component 9 (MCM9) responsible for premature ovarian insufficiency. <i>Fertility and Sterility</i> , 2020, 113, 845-852.	0.5	24
8	Big data-driven precision medicine: Starting the custom-made era of iatrogeny. <i>Biomedicine and Pharmacotherapy</i> , 2020, 129, 110445.	2.5	19
9	<i>FANCL</i> gene mutations in premature ovarian insufficiency. <i>Human Mutation</i> , 2020, 41, 1033-1041.	1.1	32
10	Genetics of Primary Ovarian Insufficiency in the Next-Generation Sequencing Era. <i>Journal of the Endocrine Society</i> , 2020, 4, bvz037.	0.1	45
11	NOTCH2 variant D1853H is mutated in two non-syndromic premature ovarian insufficiency patients from a Chinese pedigree. <i>Journal of Ovarian Research</i> , 2020, 13, 41.	1.3	7
12	Homozygous hypomorphic <i>BRCA2</i> variant in primary ovarian insufficiency without cancer or Fanconi anaemia trait. <i>Journal of Medical Genetics</i> , 2021, 58, 125-134.	1.5	24
13	Concerns regarding the potentially causal role of FANCA heterozygous variants in human primary ovarian insufficiency. <i>Human Genetics</i> , 2021, 140, 691-694.	1.8	2
14	Homozygous mutations in <i>C14orf39/SIX6OS1</i> cause non-obstructive azoospermia and premature ovarian insufficiency in humans. <i>American Journal of Human Genetics</i> , 2021, 108, 324-336.	2.6	50
15	Genetic etiologic analysis in 74 Chinese Han women with idiopathic premature ovarian insufficiency by combined molecular genetic testing. <i>Journal of Assisted Reproduction and Genetics</i> , 2021, 38, 965-978.	1.2	9
16	Meiotic Recombination Defects and Premature Ovarian Insufficiency. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 652407.	1.8	25
17	Haploinsufficiency in non-homologous end joining factor 1 induces ovarian dysfunction in humans and mice. <i>Journal of Medical Genetics</i> , 2022, 59, 579-588.	1.5	3
18	Fanconi Anemia Gene Variants in Patients with Gonadal Dysfunction. <i>Reproductive Sciences</i> , 2022, 29, 1408-1413.	1.1	6

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19	A Preliminary Study on the Mechanism of Premature Ovarian Failure. <i>Proceedings of Anticancer Research</i> , 2021, 5, 34-39.	0.0	0
20	OB-Folds and Genome Maintenance: Targeting Protein-DNA Interactions for Cancer Therapy. <i>Cancers</i> , 2021, 13, 3346.	1.7	6
22	Should FANCL heterozygous pathogenic variants be considered as potentially causative of primary ovarian insufficiency?. <i>Human Mutation</i> , 2020, 41, 1697-1699.	1.1	2
23	lncRNA DDGC participates in premature ovarian insufficiency through regulating RAD51 and WT1. <i>Molecular Therapy - Nucleic Acids</i> , 2021, 26, 1092-1106.	2.3	12
24	Genetics of ovarian insufficiency and defects of folliculogenesis. <i>Best Practice and Research in Clinical Endocrinology and Metabolism</i> , 2022, 36, 101594.	2.2	36
25	Effects of BRCA gene mutation on female reproductive potential: A systematic review. <i>Maturitas</i> , 2020, 137, 11-17.	1.0	5
26	Genetics of Primary Ovarian Insufficiency. <i>Clinical Obstetrics and Gynecology</i> , 2020, 63, 687-705.	0.6	4
27	Impairment of Pol $\beta$ -related DNA base-excision repair leads to ovarian aging in mice. <i>Aging</i> , 2020, 12, 25207-25228.	1.4	7
28	Pathogenic Variations of Homologous Recombination Gene HSF2BP Identified in Sporadic Patients With Premature Ovarian Insufficiency. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 768123.	1.8	7
29	Biallelic <i>SPATA22</i> variants cause premature ovarian insufficiency and nonobstructive azoospermia due to meiotic arrest. <i>Clinical Genetics</i> , 2022, , .	1.0	7
30	Homozygous Variant in <i>KASH5</i> Causes Premature Ovarian Insufficiency by Disordered Meiotic Homologous Pairing. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2022, 107, 2589-2597.	1.8	7
31	Progress in modern reproductive biology research in China. <i>Biology of Reproduction</i> , 0, , .	1.2	0
32	A novel cancer risk prediction score for the natural course of FA patients with biallelic <i>BRCA2/FANCD1</i> mutations. <i>Human Molecular Genetics</i> , 2023, 32, 1836-1849.	1.4	1
33	Landscape of pathogenic mutations in premature ovarian insufficiency. <i>Nature Medicine</i> , 2023, 29, 483-492.	15.2	34
34	Single-cell transcriptome analysis of the mouse and primate ovaries reveals oocyte-specific expression patterns of risk genes in ovarian aging. <i>MedComm</i> , 2023, 4, .	3.1	0