

Netta MÄäkinen

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

1,994
citations

471509

17
h-index

713466

21
g-index

22
all docs

22
docs citations

22
times ranked

2155
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>MED12</i> , the <i>Mediator Complex Subunit 12</i> Gene, Is Mutated at High Frequency in Uterine Leiomyomas. <i>Science</i> , 2011, 334, 252-255.	12.6	547
2	Characterization of Uterine Leiomyomas by Whole-Genome Sequencing. <i>New England Journal of Medicine</i> , 2013, 369, 43-53.	27.0	280
3	Integrated data analysis reveals uterine leiomyoma subtypes with distinct driver pathways and biomarkers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 1315-1320.	7.1	166
4	Genomics of uterine leiomyomas: Insights from high-throughput sequencing. <i>Fertility and Sterility</i> , 2014, 102, 621-629.	1.0	164
5	Uterine Leiomyoma-Linked <i>MED12</i> Mutations Disrupt Mediator-Associated CDK Activity. <i>Cell Reports</i> , 2014, 7, 654-660.	6.4	125
6	<i>MED12</i> exon 2 mutations are common in uterine leiomyomas from South African patients. <i>Oncotarget</i> , 2011, 2, 966-969.	1.8	95
7	Exome Sequencing of Uterine Leiomyosarcomas Identifies Frequent Mutations in TP53, ATRX, and <i>MED12</i> . <i>PLoS Genetics</i> , 2016, 12, e1005850.	3.5	94
8	Characterization of <i>MED12</i> , <i>HMGA2</i> , and <i>FH</i> alterations reveals molecular variability in uterine smooth muscle tumors. <i>Molecular Cancer</i> , 2017, 16, 101.	19.2	74
9	<i>MED12</i> exon 2 mutations in histopathological uterine leiomyoma variants. <i>European Journal of Human Genetics</i> , 2013, 21, 1300-1303.	2.8	66
10	<i>MED12</i> mutation frequency in unselected sporadic uterine leiomyomas. <i>Fertility and Sterility</i> , 2014, 102, 1137-1142.	1.0	62
11	Exome-wide somatic mutation characterization of small bowel adenocarcinoma. <i>PLoS Genetics</i> , 2018, 14, e1007200.	3.5	62
12	Deficient H2A.Z deposition is associated with genesis of uterine leiomyoma. <i>Nature</i> , 2021, 596, 398-403.	27.8	53
13	<i>MED12</i> mutations and <i>FH</i> inactivation are mutually exclusive in uterine leiomyomas. <i>British Journal of Cancer</i> , 2016, 114, 1405-1411.	6.4	43
14	Exomic landscape of <i>MED12</i> mutation-negative and -positive uterine leiomyomas. <i>International Journal of Cancer</i> , 2014, 134, 1008-1012.	5.1	36
15	Somatic <i>MED12</i> mutations in prostate cancer and uterine leiomyomas promote tumorigenesis through distinct mechanisms. <i>Prostate</i> , 2016, 76, 22-31.	2.3	33
16	Global metabolomic profiling of uterine leiomyomas. <i>British Journal of Cancer</i> , 2017, 117, 1855-1864.	6.4	29
17	Somatic <i>MED12</i> Nonsense Mutation Escapes mRNA Decay and Reveals a Motif Required for Nuclear Entry. <i>Human Mutation</i> , 2017, 38, 269-274.	2.5	20
18	Clonally related uterine leiomyomas are common and display branched tumor evolution. <i>Human Molecular Genetics</i> , 2015, 24, 4407-4416.	2.9	19

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19	Patterns of chromosome 18 loss of heterozygosity in multifocal ileal neuroendocrine tumors. <i>Genes Chromosomes and Cancer</i> , 2020, 59, 535-539.	2.8	16
20	Comparison of 2SC, AKR1B10, and FH Antibodies as Potential Biomarkers for FH-deficient Uterine Leiomyomas. <i>American Journal of Surgical Pathology</i> , 2022, 46, 537-546.	3.7	8
21	Parity associates with chromosomal damage in uterine leiomyomas. <i>Nature Communications</i> , 2021, 12, 5448.	12.8	2
22	Genomic Evolution in a Patient With Lung Adenocarcinoma With a Germline EGFR T790M Mutation. <i>JTO Clinical and Research Reports</i> , 2021, 2, 100146.	1.1	0