

Andreas Laner

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

407
citations

1163117

8
h-index

1199594

12
g-index

12
all docs

12
docs citations

12
times ranked

971
citing authors

#	ARTICLE	IF	CITATIONS
1	Exome Sequencing Identifies Biallelic MSH3 Germline Mutations as a Recessive Subtype of Colorectal Adenomatous Polyposis. American Journal of Human Genetics, 2016, 99, 337-351.	6.2	198
2	Biallelic MUTYH mutations can mimic Lynch syndrome. European Journal of Human Genetics, 2014, 22, 1334-1337.	2.8	87
3	Solving patients with rare diseases through programmatic reanalysis of genome-phenome data. European Journal of Human Genetics, 2021, 29, 1337-1347.	2.8	34
4	Comprehensive analysis of the <i>MLH1</i> promoter region in 480 patients with colorectal cancer and 1150 controls reveals new variants including one with a heritable constitutional <i>MLH1</i> epimutation. Journal of Medical Genetics, 2018, 55, 240-248.	3.2	23
5	Full-length transcript amplification and sequencing as universal method to test mRNA integrity and biallelic expression in mismatch repair genes. European Journal of Human Genetics, 2019, 27, 1808-1820.	2.8	16
6	Prevalence of CNV-neutral structural genomic rearrangements in MLH1, MSH2, and PMS2 not detectable in routine NGS diagnostics. Familial Cancer, 2020, 19, 161-167.	1.9	11
7	Targeted deep-intronic sequencing in a cohort of unexplained cases of suspected Lynch syndrome. European Journal of Human Genetics, 2020, 28, 597-608.	2.8	10
8	A mosaic PIK3CA variant in a young adult with diffuse gastric cancer: case report. European Journal of Human Genetics, 2021, 29, 1354-1358.	2.8	9
9	Splicing analyses for variants in MMR genes: best practice recommendations from the European Mismatch Repair Working Group. European Journal of Human Genetics, 2022, 30, 1051-1059.	2.8	7
10	Analysis of 3297 individuals suggests that the pathogenic germline 5'UTR variant BRCA1 c.-107A>&T is not common in south-east Germany. Familial Cancer, 2020, 19, 211-213.	1.9	6
11	Long-term chemoprevention in patients with adenomatous polyposis coli: an observational study. Familial Cancer, 2022, 21, 463-472.	1.9	4
12	Solving the genetic aetiology of hereditary gastrointestinal tumour syndromes – a collaborative multicentre endeavour within the project Solve-RD. European Journal of Medical Genetics, 2022, 65, 104475.	1.3	2