

# Jeffrey Conroy

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10472149/publications.pdf>

Version: 2024-02-01

12  
papers

1,477  
citations

840776

11  
h-index

1199594

12  
g-index

12  
all docs

12  
docs citations

12  
times ranked

1699  
citing authors

#	ARTICLE	IF	CITATIONS
1	Correlating array comparative genomic hybridization findings with histology and outcome in spitzoid melanocytic neoplasms. <i>International Journal of Clinical and Experimental Pathology</i> , 2010, 3, 593-9.	0.5	54
2	Duplication of the entire 22.9ÂMb human chromosome 21 syntenic region on mouse chromosome 16 causes cardiovascular and gastrointestinal abnormalities. <i>Human Molecular Genetics</i> , 2007, 16, 1359-1366.	2.9	165
3	Karyotypic changes detected by comparative genomic hybridization in a stillborn infant with chorioangioma and liver hemangioma. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2007, 79, 236-241.	1.6	12
4	Identification of large-scale human-specific copy number differences by inter-species array comparative genomic hybridization. <i>Human Genetics</i> , 2006, 119, 185-198.	3.8	35
5	Complex patterns of copy number variation at sites of segmental duplications: an important category of structural variation in the human genome. <i>Human Genetics</i> , 2006, 120, 270-284.	3.8	68
6	Array CGH analysis of pediatric medulloblastomas. <i>Genes Chromosomes and Cancer</i> , 2006, 45, 290-303.	2.8	59
7	Identification of consistent novel submegabase deletions in low-grade oligodendrogliomas using array-based comparative genomic hybridization. <i>Genes Chromosomes and Cancer</i> , 2005, 44, 85-96.	2.8	36
8	Novel karyotypic changes detected by comparative genomic hybridization in a case of congenital cervical immature teratoma. <i>Birth Defects Research Part A: Clinical and Molecular Teratology</i> , 2005, 73, 572-576.	1.6	7
9	Mapping segmental and sequence variations among laboratory mice using BAC array CGH. <i>Genome Research</i> , 2005, 15, 302-311.	5.5	64
10	Two Functional Coding Single Nucleotide Polymorphisms in STK15 (Aurora-A) Coordinately Increase Esophageal Cancer Risk. <i>Cancer Research</i> , 2005, 65, 3548-3554.	0.9	65
11	Application of bacterial artificial chromosome array-based comparative genomic hybridization and spectral karyotyping to the analysis of glioblastoma multiforme. <i>Cancer Genetics and Cytogenetics</i> , 2004, 151, 36-51.	1.0	58
12	Assembly of microarrays for genome-wide measurement of DNA copy number. <i>Nature Genetics</i> , 2001, 29, 263-264.	21.4	854