## Tiffany Hung

List of Publications by Year in descending order

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Version: 2024-02-01

933447 1281871 7,789 11 10 11 citations h-index g-index papers 13 13 13 11898 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Pre―and postâ€treatment bloodâ€based genomic landscape of patients with <i>ROS1</i> or <i>NTRK</i> fusionâ€positive solid tumours treated with entrectinib. Molecular Oncology, 2022, 16, 2000-2014.	4.6	10
2	Toward best practice in cancer mutation detection with whole-genome and whole-exome sequencing. Nature Biotechnology, 2021, 39, 1141-1150.	17.5	66
3	Establishing community reference samples, data and call sets for benchmarking cancer mutation detection using whole-genome sequencing. Nature Biotechnology, 2021, 39, 1151-1160.	17.5	39
4	Whole genome and exome sequencing reference datasets from a multi-center and cross-platform benchmark study. Scientific Data, 2021, 8, 296.	5.3	15
5	An inducible long noncoding RNA amplifies DNA damage signaling. Nature Genetics, 2016, 48, 1370-1376.	21.4	195
6	Extensive and coordinated transcription of noncoding RNAs within cell-cycle promoters. Nature Genetics, 2011, 43, 621-629.	21.4	1,080
7	Long non-coding RNA HOTAIR reprograms chromatin state to promote cancer metastasis. Nature, 2010, 464, 1071-1076.	27.8	4,648
8	Long noncoding RNA in genome regulation. RNA Biology, 2010, 7, 582-585.	3.1	523
9	ING4 Mediates Crosstalk between Histone H3 K4 Trimethylation and H3 Acetylation to Attenuate Cellular Transformation. Molecular Cell, 2009, 33, 248-256.	9.7	191
10	HBO1 HAT Complexes Target Chromatin throughout Gene Coding Regions via Multiple PHD Finger Interactions with Histone H3 Tail. Molecular Cell, 2009, 33, 257-265.	9.7	163
11	ING2 PHD domain links histone H3 lysine 4 methylation to active gene repression. Nature, 2006, 442, 96-99.	27.8	851