

# Dinh Diep

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/505942/publications.pdf>

Version: 2024-02-01

10  
papers

636  
citations

2258059

3  
h-index

1872680

6  
g-index

13  
all docs

13  
docs citations

13  
times ranked

1768  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of methylation haplotype blocks aids in deconvolution of heterogeneous tissue samples and tumor tissue-of-origin mapping from plasma DNA. <i>Nature Genetics</i> , 2017, 49, 635-642.	21.4	384
2	Library-free methylation sequencing with bisulfite padlock probes. <i>Nature Methods</i> , 2012, 9, 270-272.	19.0	92
3	DNA methylation identifies genetically and prognostically distinct subtypes of myelodysplastic syndromes. <i>Blood Advances</i> , 2019, 3, 2845-2858.	5.2	32
4	5-Azacytidine Transiently Restores Dysregulated Erythroid Differentiation Gene Expression in TET2-Deficient Erythroleukemia Cells. <i>Molecular Cancer Research</i> , 2021, 19, 451-464.	3.4	3
5	Large-Scale Targeted DNA Methylation Analysis Using Bisulfite Padlock Probes. <i>Methods in Molecular Biology</i> , 2018, 1708, 365-382.	0.9	1
6	Reply to "DNA methylation haplotypes as cancer markers". <i>Nature Genetics</i> , 2018, 50, 1063-1066.	21.4	1
7	Genome-wide mapping of the sixth base. <i>Genome Biology</i> , 2011, 12, 116.	9.6	0
8	Efficient and fast identification of differentially methylated regions using whole-genome bisulfite sequencing data. <i>Journal of Genetics and Genomics</i> , 2018, 45, 455-457.	3.9	0
9	Cytosine Methylation Patterns As Biomarkers in MDS. <i>Blood</i> , 2015, 126, 5226-5226.	1.4	0
10	Methylation Profiling Among Patients with TET2 Mutations in Myelodysplastic Syndromes Is Associated with Overall Survival. <i>Blood</i> , 2016, 128, 3158-3158.	1.4	0