

# Halit Ongen

## List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

15  
papers

6,203  
citations

12  
h-index

21  
g-index

21  
ext. papers

7,949  
ext. citations

19.4  
avg, IF

4.37  
L-index

#	Paper	IF	Citations
15	Human genomics. The Genotype-Tissue Expression (GTEx) pilot analysis: multitissue gene regulation in humans. <i>Science</i> , <b>2015</b> , 348, 648-60	33.3	3242
14	Genetic variants in novel pathways influence blood pressure and cardiovascular disease risk. <i>Nature</i> , <b>2011</b> , 478, 103-9	50.4	1564
13	Passive and active DNA methylation and the interplay with genetic variation in gene regulation. <i>ELife</i> , <b>2013</b> , 2, e00523	8.9	295
12	Fast and efficient QTL mapper for thousands of molecular phenotypes. <i>Bioinformatics</i> , <b>2016</b> , 32, 1479-85	7.2	241
11	Using an atlas of gene regulation across 44 human tissues to inform complex disease- and trait-associated variation. <i>Nature Genetics</i> , <b>2018</b> , 50, 956-967	36.3	239
10	Tissue-specific effects of genetic and epigenetic variation on gene regulation and splicing. <i>PLoS Genetics</i> , <b>2015</b> , 11, e1004958	6	140
9	Putative cis-regulatory drivers in colorectal cancer. <i>Nature</i> , <b>2014</b> , 512, 87-90	50.4	115
8	Estimating the causal tissues for complex traits and diseases. <i>Nature Genetics</i> , <b>2017</b> , 49, 1676-1683	36.3	106
7	A complete tool set for molecular QTL discovery and analysis. <i>Nature Communications</i> , <b>2017</b> , 8, 15452	17.4	103
6	Molecular-Subtype-Specific Biomarkers Improve Prediction of Prognosis in Colorectal Cancer. <i>Cell Reports</i> , <b>2017</b> , 19, 1268-1280	10.6	55
5	Combined genetic and transcriptome analysis of patients with SLE: distinct, targetable signatures for susceptibility and severity. <i>Annals of the Rheumatic Diseases</i> , <b>2019</b> , 78, 1079-1089	2.4	45
4	Alternative Splicing QTLs in European and African Populations. <i>American Journal of Human Genetics</i> , <b>2015</b> , 97, 567-75	11	39
3	Contribution of allelic imbalance to colorectal cancer. <i>Nature Communications</i> , <b>2018</b> , 9, 3664	17.4	11
2	Comprehensive evaluation of coding region point mutations in microsatellite-unstable colorectal cancer. <i>EMBO Molecular Medicine</i> , <b>2018</b> , 10,	12	6
1	MethCORR modelling of methylomes from formalin-fixed paraffin-embedded tissue enables characterization and prognostication of colorectal cancer. <i>Nature Communications</i> , <b>2020</b> , 11, 2025	17.4	2