

# Lee E Edsall

## 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.

16  
papers

3,020  
citations

12  
h-index

22  
g-index

22  
ext. papers

3,594  
ext. citations

15.1  
avg, IF

3.41  
L-index

#	Paper	IF	Citations
16	A comparative encyclopedia of DNA elements in the mouse genome. <i>Nature</i> , <b>2014</b> , 515, 355-64	50.4	1026
15	Distinct epigenomic landscapes of pluripotent and lineage-committed human cells. <i>Cell Stem Cell</i> , <b>2010</b> , 6, 479-91	18	630
14	Global DNA hypomethylation coupled to repressive chromatin domain formation and gene silencing in breast cancer. <i>Genome Research</i> , <b>2012</b> , 22, 246-58	9.7	385
13	An encyclopedia of mouse DNA elements (Mouse ENCODE). <i>Genome Biology</i> , <b>2012</b> , 13, 418	18.3	340
12	5mC oxidation by Tet2 modulates enhancer activity and timing of transcriptome reprogramming during differentiation. <i>Molecular Cell</i> , <b>2014</b> , 56, 286-297	17.6	226
11	Integrative analysis of haplotype-resolved epigenomes across human tissues. <i>Nature</i> , <b>2015</b> , 518, 350-354	50.4	155
10	Dynamic chromatin states in human ES cells reveal potential regulatory sequences and genes involved in pluripotency. <i>Cell Research</i> , <b>2011</b> , 21, 1393-409	24.7	80
9	Construction of a medicinal leech transcriptome database and its application to the identification of leech homologs of neural and innate immune genes. <i>BMC Genomics</i> , <b>2010</b> , 11, 407	4.5	44
8	Exome analysis identified a novel mutation in the RBP4 gene in a consanguineous pedigree with retinal dystrophy and developmental abnormalities. <i>PLoS ONE</i> , <b>2012</b> , 7, e50205	3.7	42
7	DNA binding by FOXP3 domain-swapped dimer suggests mechanisms of long-range chromosomal interactions. <i>Nucleic Acids Research</i> , <b>2015</b> , 43, 1268-82	20.1	35
6	Systematic detection of putative tumor suppressor genes through the combined use of exome and transcriptome sequencing. <i>Genome Biology</i> , <b>2010</b> , 11, R114	18.3	32
5	Incomplete MyoD-induced transdifferentiation is associated with chromatin remodeling deficiencies. <i>Nucleic Acids Research</i> , <b>2017</b> , 45, 11684-11699	20.1	17
4	Evaluating Chromatin Accessibility Differences Across Multiple Primate Species Using a Joint Modeling Approach. <i>Genome Biology and Evolution</i> , <b>2019</b> , 11, 3035-3053	3.9	3
3	Epstein-Barr virus nuclear antigen 2 extensively rewires the human chromatin landscape at autoimmune risk loci. <i>Genome Research</i> , <b>2021</b> ,	9.7	3
2	Epigenetic Analysis of the Chromatin Landscape Identifies a Repertoire of Murine Eosinophil-Specific PU.1-Bound Enhancers. <i>Journal of Immunology</i> , <b>2021</b> , 207, 1044-1054	5.3	2
1	Epigenetic and transcriptional dysregulation in CD4+ T cells in patients with atopic dermatitis.. <i>PLoS Genetics</i> , <b>2022</b> , 18, e1009973	6	0