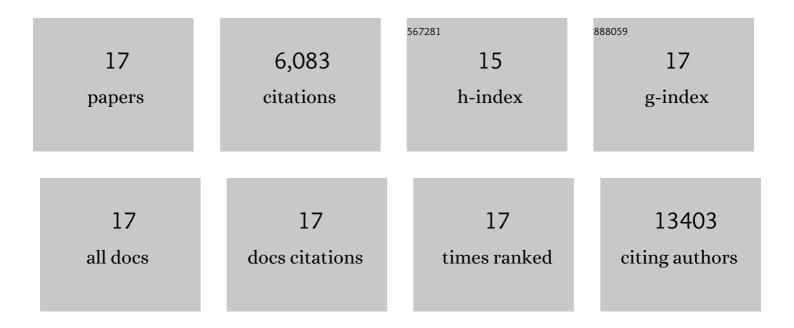
## Heidi Dvinge

List of Publications by Year in descending order

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HEIDI DVINCE

#	Article	IF	CITATIONS
1	Integrative Clinical Genomics of Advanced Prostate Cancer. Cell, 2015, 161, 1215-1228.	28.9	2,660
2	Systematic comparison of microarray profiling, real-time PCR, and next-generation sequencing technologies for measuring differential microRNA expression. Rna, 2010, 16, 991-1006.	3.5	588
3	RNA splicing factors as oncoproteins and tumour suppressors. Nature Reviews Cancer, 2016, 16, 413-430.	28.4	549
4	The shaping and functional consequences of the microRNA landscape in breast cancer. Nature, 2013, 497, 378-382.	27.8	370
5	Modulation of splicing catalysis for therapeutic targeting of leukemia with mutations in genes encoding spliceosomal proteins. Nature Medicine, 2016, 22, 672-678.	30.7	301
6	Widespread intron retention diversifies most cancer transcriptomes. Genome Medicine, 2015, 7, 45.	8.2	283
7	Modeling of C/EBPα Mutant Acute Myeloid Leukemia Reveals a Common Expression Signature of Committed Myeloid Leukemia-Initiating Cells. Cancer Cell, 2008, 13, 299-310.	16.8	225
8	NuRD-mediated deacetylation of H3K27 facilitates recruitment of Polycomb Repressive Complex 2 to direct gene repression. EMBO Journal, 2012, 31, 593-605.	7.8	224
9	PeakAnalyzer: Genome-wide annotation of chromatin binding and modification loci. BMC Bioinformatics, 2010, 11, 415.	2.6	210
10	<i>HTqPCR</i> : high-throughput analysis and visualization of quantitative real-time PCR data in R. Bioinformatics, 2009, 25, 3325-3326.	4.1	205
11	Coordinated alterations in RNA splicing and epigenetic regulation drive leukaemogenesis. Nature, 2019, 574, 273-277.	27.8	149
12	RNA components of the spliceosome regulate tissue- and cancer-specific alternative splicing. Genome Research, 2019, 29, 1591-1604.	5.5	96
13	Mapping Organelle Proteins and Protein Complexes in <i>Drosophila melanogaster</i> . Journal of Proteome Research, 2009, 8, 2667-2678.	3.7	71
14	Regulation of alternative <scp>mRNA</scp> splicing: old players and new perspectives. FEBS Letters, 2018, 592, 2987-3006.	2.8	71
15	Genome-wide characterization of Foxa2 targets reveals upregulation of floor plate genes and repression of ventrolateral genes in midbrain dopaminergic progenitors. Development (Cambridge), 2012, 139, 2625-2634.	2.5	55
16	Alternative Splicing and Cleavage of GLUT8. Molecular and Cellular Biology, 2021, 41, .	2.3	14
17	Human spliceosomal snRNA sequence variants generate variant spliceosomes. Rna, 2021, 27, 1186-1203.	3.5	12