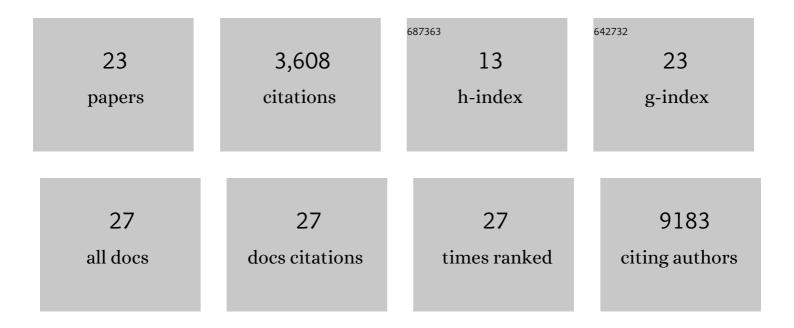
## Ilya E Vorontsov

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	ANANASTRA: annotation and enrichment analysis of allele-specific transcription factor binding at SNPs. Nucleic Acids Research, 2022, 50, W51-W56.	14.5	7
2	Landscape of allele-specific transcription factor binding in the human genome. Nature Communications, 2021, 12, 2751.	12.8	55
3	Enhanced C/EBP binding to G·T mismatches facilitates fixation of CpG mutations in cancer and adult stem cells. Cell Reports, 2021, 35, 109221.	6.4	5
4	Assessing Ribosome Distribution Along Transcripts with Polarity Scores and Regression Slope Estimates. Methods in Molecular Biology, 2021, 2252, 269-294.	0.9	1
5	A holistic view of mouse enhancer architectures reveals analogous pleiotropic effects and correlation with human disease. BMC Genomics, 2020, 21, 754.	2.8	3
6	Functional annotation of human long noncoding RNAs via molecular phenotyping. Genome Research, 2020, 30, 1060-1072.	5.5	109
7	Insights gained from a comprehensive all-against-all transcription factor binding motif benchmarking study. Genome Biology, 2020, 21, 114.	8.8	39
8	What Do Neighbors Tell About You: The Local Context of Cis-Regulatory Modules Complicates Prediction of Regulatory Variants. Frontiers in Genetics, 2019, 10, 1078.	2.3	3
9	Abstract 5206: Role of super-enhancers in HPV+ head and neck squamous cell carcinoma. , 2019, , .		0
10	HOCOMOCO: towards a complete collection of transcription factor binding models for human and mouse via large-scale ChIP-Seq analysis. Nucleic Acids Research, 2018, 46, D252-D259.	14.5	660
11	Genome-wide map of human and mouse transcription factor binding sites aggregated from ChIP-Seq data. BMC Research Notes, 2018, 11, 756.	1.4	19
12	Multiple single nucleotide polymorphisms in the first intron of the IL2RA gene affect transcription factor binding and enhancer activity. Gene, 2017, 602, 50-56.	2.2	19
13	The single nucleotide variant rs12722489 determines differential estrogen receptor binding and enhancer properties of an IL2RA intronic region. PLoS ONE, 2017, 12, e0172681.	2.5	10
14	Early B-cell factor 1 (EBF1) is critical for transcriptional control of SLAMF1 gene in human B cells. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2016, 1859, 1259-1268.	1.9	20
15	Negative selection maintains transcription factor binding motifs in human cancer. BMC Genomics, 2016, 17, 395.	2.8	16
16	HOCOMOCO: expansion and enhancement of the collection of transcription factor binding sites models. Nucleic Acids Research, 2016, 44, D116-D125.	14.5	215
17	EpiFactors: a comprehensive database of human epigenetic factors and complexes. Database: the Journal of Biological Databases and Curation, 2015, 2015, bav067.	3.0	226
18	PERFECTOS-APE - Predicting Regulatory Functional Effect of SNPs by Approximate P-value Estimation. , 2015, , .		28

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#	Article	IF	CITATIONS
19	A promoter-level mammalian expression atlas. Nature, 2014, 507, 462-470.	27.8	1,838
20	Jaccard index based similarity measure to compare transcription factor binding site models. Algorithms for Molecular Biology, 2013, 8, 23.	1.2	53
21	In silico motif analysis suggests an interplay of transcriptional and translational control in mTOR response. Translation, 2013, 1, 18-24.	2.9	9
22	HOCOMOCO: a comprehensive collection of human transcription factor binding sites models. Nucleic Acids Research, 2013, 41, D195-D202.	14.5	206
23	FROM BINDING MOTIFS IN CHIP-SEQ DATA TO IMPROVED MODELS OF TRANSCRIPTION FACTOR BINDING SITES. Journal of Bioinformatics and Computational Biology, 2013, 11, 1340004.	0.8	58