

# Nicolas Frankel

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1066550/publications.pdf>

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11  
papers

1,352  
citations

1163117

8  
h-index

1281871

11  
g-index

11  
all docs

11  
docs citations

11  
times ranked

2143  
citing authors

#	ARTICLE	IF	CITATIONS
1	Pleiotropic Enhancers are Ubiquitous Regulatory Elements in the Human Genome. <i>Genome Biology and Evolution</i> , 2022, 14, .	2.5	5
2	<i>Cis</i> -regulatory variation in the <i>shavenbaby</i> gene underlies intraspecific phenotypic variation, mirroring interspecific divergence in the same trait. <i>Evolution; International Journal of Organic Evolution</i> , 2021, 75, 427-436.	2.3	7
3	A complex gene regulatory architecture underlies the development and evolution of cuticle morphology in <i>Drosophila</i> . <i>Current Opinion in Genetics and Development</i> , 2021, 69, 21-27.	3.3	11
4	Actors with Multiple Roles: Pleiotropic Enhancers and the Paradigm of Enhancer Modularity. <i>Trends in Genetics</i> , 2019, 35, 423-433.	6.7	61
5	Comprehensive Analysis of a cis-Regulatory Region Reveals Pleiotropy in Enhancer Function. <i>Cell Reports</i> , 2018, 22, 3021-3031.	6.4	81
6	Gene regulatory network architecture in different developmental contexts influences the genetic basis of morphological evolution. <i>PLoS Genetics</i> , 2018, 14, e1007375.	3.5	38
7	Low Affinity Binding Site Clusters Confer Hox Specificity and Regulatory Robustness. <i>Cell</i> , 2015, 160, 191-203.	28.9	322
8	The structure and evolution of <i>cis</i> -regulatory regions: the <i>shavenbaby</i> story. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2013, 368, 20130028.	4.0	71
9	Conserved regulatory architecture underlies parallel genetic changes and convergent phenotypic evolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 20975-20979.	7.1	82
10	Morphological evolution caused by many subtle-effect substitutions in regulatory DNA. <i>Nature</i> , 2011, 474, 598-603.	27.8	199
11	Phenotypic robustness conferred by apparently redundant transcriptional enhancers. <i>Nature</i> , 2010, 466, 490-493.	27.8	475