

# Yohann Ndlec

## List of Publications by Year in Descending Order

**Source:** <https://exaly.com/author-pdf/11014538/yohann-nedelec-publications-by-year.pdf>

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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.

10  
papers

647  
citations

8  
h-index

10  
g-index

10  
ext. papers

874  
ext. citations

14.9  
avg, IF

2.99  
L-index

#	Paper	IF	Citations
10	Genetic regulatory effects modified by immune activation contribute to autoimmune disease associations. <i>Nature Communications</i> , <b>2017</b> , 8, 266	17.4	93
9	Deciphering the genetic control of gene expression following <i>Mycobacterium leprae</i> antigen stimulation. <i>PLoS Genetics</i> , <b>2017</b> , 13, e1006952	6	17
8	Genetic Ancestry and Natural Selection Drive Population Differences in Immune Responses to Pathogens. <i>Cell</i> , <b>2016</b> , 167, 657-669.e21	56.2	264
7	Widespread Shortening of 3' Untranslated Regions and Increased Exon Inclusion Are Evolutionarily Conserved Features of Innate Immune Responses to Infection. <i>PLoS Genetics</i> , <b>2016</b> , 12, e1006338	6	53
6	Adaptively introgressed Neandertal haplotype at the OAS locus functionally impacts innate immune responses in humans. <i>Genome Biology</i> , <b>2016</b> , 17, 246	18.3	70
5	A genomic portrait of the genetic architecture and regulatory impact of microRNA expression in response to infection. <i>Genome Research</i> , <b>2014</b> , 24, 850-9	9.7	49
4	When genetics meets epigenetics: deciphering the mechanisms controlling inter-individual variation in immune responses to infection. <i>Current Opinion in Immunology</i> , <b>2014</b> , 29, 119-26	7.8	6
3	Adaptive, convergent origins of the pygmy phenotype in African rainforest hunter-gatherers. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2014</b> , 111, E3596-603	11.5	70
2	Gene set signature of reversal reaction type I in leprosy patients. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003624	6	23
1	Widespread shortening of 3' untranslated regions and increased exon inclusion are evolutionarily conserved features of innate immune responses to infection		2