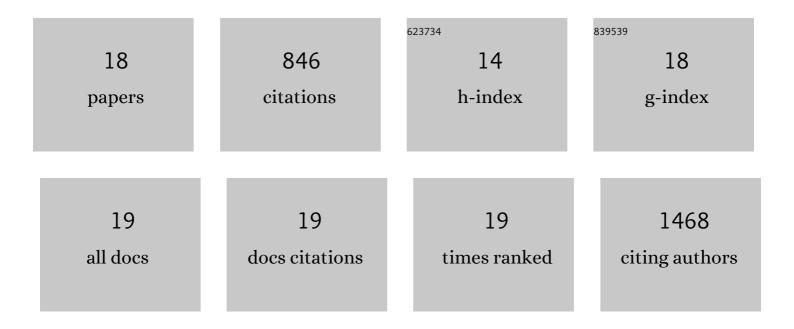
Claire Vandiedonck

List of Publications by Year in descending order

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#	Article	lF	CITATIONS
1	Regulation of inflammation in diabetes: From genetics to epigenomics evidence. Molecular Metabolism, 2020, 41, 101041.	6.5	23
2	Genetic association of molecular traits: A help to identify causative variants in complex diseases. Clinical Genetics, 2018, 93, 520-532.	2.0	45
3	Network-based analysis of omics data: the LEAN method. Bioinformatics, 2017, 33, 701-709.	4.1	29
4	Juvenile-Onset Diabetes and Congenital Cataract: "Double-Gene―Mutations Mimicking a Syndromic Diabetes Presentation. Genes, 2017, 8, 309.	2.4	8
5	A functional AT/G polymorphism in the 5′-untranslated region of SETDB2 in the IgE locus on human chromosome 13q14. Genes and Immunity, 2015, 16, 488-494.	4.1	6
6	Fine mapping genetic determinants of the highly variably expressed MHC gene ZFP57. European Journal of Human Genetics, 2014, 22, 568-571.	2.8	16
7	Allele-specific transcription of the asthma-associated PHD finger protein 11 gene (PHF11) modulated by octamer-binding transcription factor 1 (Oct-1). Journal of Allergy and Clinical Immunology, 2011, 127, 1054-1062.e2.	2.9	15
8	Pervasive haplotypic variation in the spliceo-transcriptome of the human major histocompatibility complex. Genome Research, 2011, 21, 1042-1054.	5.5	63
9	Association of HLA-A in autoimmune myasthenia gravis with thymoma. Journal of Neuroimmunology, 2009, 210, 120-123.	2.3	28
10	The human Major Histocompatibility Complex as a paradigm in genomics research. Briefings in Functional Genomics & Proteomics, 2009, 8, 379-394.	3.8	85
11	<i>Genetic Factors in Autoimmune Myasthenia Gravis</i> . Annals of the New York Academy of Sciences, 2008, 1132, 180-192.	3.8	79
12	Chromatin profiling across the human tumour necrosis factor gene locus reveals a complex, cell type-specific landscape with novel regulatory elements. Nucleic Acids Research, 2008, 36, 4845-4862.	14.5	23
13	Validating Discovered Cis-Acting Regulatory Genetic Variants: Application of an Allele Specific Expression Approach to HapMap Populations. PLoS ONE, 2008, 3, e4105.	2.5	22
14	An IRF8-binding promoter variant and AIRE control CHRNA1 promiscuous expression in thymus. Nature, 2007, 448, 934-937.	27.8	167
15	Association of the PTPN22*R620W polymorphism with autoimmune myasthenia gravis. Annals of Neurology, 2006, 59, 404-407.	5.3	103
16	Genetics of autoimmune myasthenia gravis: The multifaceted contribution of the HLA complex. Journal of Autoimmunity, 2005, 25, 6-11.	6.5	33
17	Pleiotropic effects of the 8.1 HLA haplotype in patients with autoimmune myasthenia gravis and thymus hyperplasia. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 15464-15469.	7.1	81
18	Genetic linkage of progressive pseudorheumatoid dysplasia to a 3-cM interval of chromosome 6q22. Human Genetics, 1998, 103, 60-64.	3.8	20