

# Valerie Turcot

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5757758/publications.pdf>

Version: 2024-02-01

15  
papers

1,342  
citations

758635

12  
h-index

996533

15  
g-index

16  
all docs

16  
docs citations

16  
times ranked

4863  
citing authors

#	ARTICLE	IF	CITATIONS
1	Meta-analysis of up to 622,409 individuals identifies 40 novel smoking behaviour associated genetic loci. <i>Molecular Psychiatry</i> , 2020, 25, 2392-2409.	4.1	83
2	Protein-coding variants implicate novel genes related to lipid homeostasis contributing to body-fat distribution. <i>Nature Genetics</i> , 2019, 51, 452-469.	9.4	89
3	Exome Chip Meta-analysis Fine Maps Causal Variants and Elucidates the Genetic Architecture of Rare Coding Variants in Smoking and Alcohol Use. <i>Biological Psychiatry</i> , 2019, 85, 946-955.	0.7	69
4	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. <i>Nature Genetics</i> , 2018, 50, 26-41.	9.4	286
5	Rare and low-frequency coding variants alter human adult height. <i>Nature</i> , 2017, 542, 186-190.	13.7	544
6	Lower Methylation of the ANGPTL2 Gene in Leukocytes from Post-Acute Coronary Syndrome Patients. <i>PLoS ONE</i> , 2016, 11, e0153920.	1.1	18
7	Validation of fatty acid intakes estimated by a food frequency questionnaire using erythrocyte fatty acid profiling in the Montreal Heart Institute Biobank. <i>Journal of Human Nutrition and Dietetics</i> , 2015, 28, 646-658.	1.3	18
8	Rare and low-frequency coding variants in CXCR2 and other genes are associated with hematological traits. <i>Nature Genetics</i> , 2014, 46, 629-634.	9.4	113
9	Postnatal Growth and DNA Methylation Are Associated With Differential Gene Expression of the TACSTD2 Gene and Childhood Fat Mass. <i>Diabetes</i> , 2012, 61, 391-400.	0.3	55
10	Influence of Shielding TPN From Photooxidation on the Number of Early Blood Transfusions in ELBW Premature Neonates. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2012, 55, 398-402.	0.9	4
11	Genetic contribution to C-reactive protein levels in severe obesity. <i>Molecular Genetics and Metabolism</i> , 2012, 105, 494-501.	0.5	20
12	Bioinformatic selection of putative epigenetically regulated loci associated with obesity using gene expression data. <i>Gene</i> , 2012, 499, 99-107.	1.0	9
13	A polymorphism of the interferon-gamma-inducible protein 30 gene is associated with hyperglycemia in severely obese individuals. <i>Human Genetics</i> , 2012, 131, 57-66.	1.8	13
14	Cardiovascular risk factors in severe obesity are associated with the inflammatory and immune-response IFI30 and TSLP genes. <i>Canadian Journal of Diabetes</i> , 2011, 35, 195.	0.4	0
15	Long-term impact of an antioxidant-deficient neonatal diet on lipid and glucose metabolism. <i>Free Radical Biology and Medicine</i> , 2009, 47, 275-282.	1.3	14