

# Suzanne B R Jacobs

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6273069/publications.pdf>

Version: 2024-02-01

12  
papers

1,789  
citations

933264

10  
h-index

1199470

12  
g-index

13  
all docs

13  
docs citations

13  
times ranked

5334  
citing authors

| #  | ARTICLE  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Sequence variants in SLC16A11 are a common risk factor for type 2 diabetes in Mexico. <i>Nature</i> , 2014, 506, 97-101.   | 13.7 | 439       |
| 2  | Loss-of-function mutations in SLC30A8 protect against type 2 diabetes. <i>Nature Genetics</i> , 2014, 46, 357-363.   | 9.4  | 428       |
| 3  | Genome-wide meta-analysis identifies new susceptibility loci for migraine. <i>Nature Genetics</i> , 2013, 45, 912-917.   | 9.4  | 338       |
| 4  | Association of a Low-Frequency Variant in <i>HNF1A</i> With Type 2 Diabetes in a Latino Population. <i>JAMA - Journal of the American Medical Association</i> , 2014, 311, 2305.               | 3.8  | 230       |
| 5  | Type 2 Diabetes Variants Disrupt Function of SLC16A11 through Two Distinct Mechanisms. <i>Cell</i> , 2017, 170, 199-212.e20.   | 13.5 | 121       |
| 6  | Polyunsaturated Fatty Acid Desaturation Is a Mechanism for Glycolytic NAD <sup>+</sup> Recycling. <i>Cell Metabolism</i> , 2019, 29, 856-870.e7.   | 7.2  | 87        |
| 7  | A Loss-of-Function Splice Acceptor Variant in <i>IGF2</i> Is Protective for Type 2 Diabetes. <i>Diabetes</i> , 2017, 66, 2903-2914.  | 0.3  | 52        |
| 8  | A Low-Frequency Inactivating <i>AKT2</i> Variant Enriched in the Finnish Population Is Associated With Fasting Insulin Levels and Type 2 Diabetes Risk. <i>Diabetes</i> , 2017, 66, 2019-2032. | 0.3  | 47        |
| 9  | The SLC16A11 risk haplotype is associated with decreased insulin action, higher transaminases and large-size adipocytes. <i>European Journal of Endocrinology</i> , 2019, 180, 99-107.         | 1.9  | 19        |
| 10 | Interaction Between Type 2 Diabetes Prevention Strategies and Genetic Determinants of Coronary Artery Disease on Cardiometabolic Risk Factors. <i>Diabetes</i> , 2020, 69, 112-120.            | 0.3  | 13        |
| 11 | Analysis of Glucocorticoid-Related Genes Reveal <i>CCHCR1</i> as a New Candidate Gene for Type 2 Diabetes. <i>Journal of the Endocrine Society</i> , 2020, 4, bvaa121.                         | 0.1  | 8         |
| 12 | Gain-of-Function Claims for Type-2-Diabetes-Associated Coding Variants in SLC16A11 Are Not Supported by the Experimental Data. <i>Cell Reports</i> , 2019, 29, 778-780.                        | 2.9  | 6         |