## Ioanna Ntalla

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/679100/publications.pdf

Version: 2024-02-01

|          |                 | 759233       | 996975         |
|----------|-----------------|--------------|----------------|
| 14       | 1,974 citations | 12           | 15             |
| papers   | citations       | h-index      | g-index        |
|          |                 |              |                |
|          |                 |              |                |
|          |                 |              | 4              |
| 19       | 19              | 19           | 6519           |
| all docs | docs citations  | times ranked | citing authors |
|          |                 |              |                |

| #  | Article   | IF   | Citations |
|----|---|------|-----------|
| 1  | Shared genetic risk between eating disorder†and substance†use†related phenotypes: Evidence from genome†wide association studies. Addiction Biology, 2021, 26, e12880.   | 2.6  | 28        |
| 2  | Cohort Profile: Extended Cohort for E-health, Environment and DNA (EXCEED). International Journal of Epidemiology, 2019, 48, 678-679j.  | 1.9  | 9         |
| 3  | Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. Nature Genetics, 2019, 51, 1207-1214.  | 21.4 | 641       |
| 4  | Exome-wide analysis of rare coding variation identifies novel associations with COPD and airflow limitation in <i>MOCS3 </i> , <i>IFIT3 </i> and <i>SERPINA12 </i> . Thorax, 2016, 71, 501-509.                             | 5.6  | 22        |
| 5  | Genome-wide association analysis identifies three new susceptibility loci for childhood body mass index. Human Molecular Genetics, 2016, 25, 389-403.   | 2.9  | 275       |
| 6  | Sixteen new lung function signals identified through 1000 Genomes Project reference panel imputation. Nature Communications, 2015, 6, 8658.   | 12.8 | 108       |
| 7  | Low-frequency and rare exome chip variants associate with fasting glucose and type 2 diabetes susceptibility. Nature Communications, 2015, 6, 5897.   | 12.8 | 173       |
| 8  | Dietary Intake, <i>FTO</i> Genetic Variants, and Adiposity: A Combined Analysis of Over 16,000 Children and Adolescents. Diabetes, 2015, 64, 2467-2476.   | 0.6  | 74        |
| 9  | Novel insights into the genetics of smoking behaviour, lung function, and chronic obstructive pulmonary disease (UK BiLEVE): a genetic association study in UK Biobank. Lancet Respiratory Medicine, the, 2015, 3, 769-781. | 10.7 | 346       |
| 10 | A novel common variant in DCST2 is associated with length in early life and height in adulthood. Human Molecular Genetics, 2015, 24, 1155-1168.   | 2.9  | 109       |
| 11 | Genetic characterization of Greek population isolates reveals strong genetic drift at missense and trait-associated variants. Nature Communications, 2014, 5, 5345.   | 12.8 | 60        |
| 12 | Body composition and eating behaviours in relation to dieting involvement in a sample of urban Greek adolescents from the TEENAGE (TEENs of Attica: Genes & Environment) study. Public Health Nutrition, 2014, 17, 561-568. | 2.2  | 12        |
| 13 | Genome-wide association study of sexual maturation in males and females highlights a role for body mass and menarche loci in male puberty. Human Molecular Genetics, 2014, 23, 4452-4464.                                   | 2.9  | 82        |
| 14 | Replication of Established Common Genetic Variants for Adult BMI and Childhood Obesity in Greek Adolescents: The TEENAGE Study. Annals of Human Genetics, 2013, 77, 268-274.  | 0.8  | 29        |