Heikki Allan Koistinen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1374772/publications.pdf

Version: 2024-02-01

36 papers 4,408 citations

236925 25 h-index 315739 38 g-index

41 all docs

41 docs citations

41 times ranked

9562 citing authors

#	Article	IF	CITATIONS
1	Circulating levels of urocortin neuropeptides are impaired in children with overweight. Obesity, 2022, 30, 472-481.	3.0	3
2	Rare coding variants in 35 genes associate with circulating lipid levelsâ€"A multi-ancestry analysis of 170,000 exomes. American Journal of Human Genetics, 2022, 109, 81-96.	6.2	24
3	ACE2 expression in adipose tissue is associated with cardio-metabolic risk factors and cell type compositionâ€"implications for COVID-19. International Journal of Obesity, 2022, 46, 1478-1486.	3.4	18
4	Sex-dimorphic genetic effects and novel loci for fasting glucose and insulin variability. Nature Communications, 2021, 12, 24.	12.8	87
5	The trans-ancestral genomic architecture of glycemic traits. Nature Genetics, 2021, 53, 840-860.	21.4	341
6	Determinants of penetrance and variable expressivity in monogenic metabolic conditions across 77,184 exomes. Nature Communications, 2021, 12, 3505.	12.8	49
7	Skeletal muscle proteomes reveal downregulation of mitochondrial proteins in transition from prediabetes into type 2 diabetes. IScience, 2021, 24, 102712.	4.1	20
8	Urocortin 3 overexpression reduces ER stress and heat shock response in 3T3-L1 adipocytes. Scientific Reports, 2021, 11, 15666.	3.3	8
9	The power of genetic diversity in genome-wide association studies of lipids. Nature, 2021, 600, 675-679.	27.8	353
10	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. Nature Genetics, 2020, 52, 1314-1332.	21.4	91
11	Simvastatin profoundly impairs energy metabolism in primary human muscle cells. Endocrine Connections, 2020, 9, 1103-1113.	1.9	0
12	Simvastatin profoundly impairs energy metabolism in primary human muscle cells. Endocrine Connections, 2020, 9, 1103-1113.	1.9	5
13	Multiancestry Genome-Wide Association Study of Lipid Levels Incorporating Gene-Alcohol Interactions. American Journal of Epidemiology, 2019, 188, 1033-1054.	3.4	85
14	Exome sequencing of 20,791Âcases of type 2 diabetes and 24,440Âcontrols. Nature, 2019, 570, 71-76.	27.8	248
15	Integrative analysis of gene expression, DNA methylation, physiological traits, and genetic variation in human skeletal muscle. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 10883-10888.	7.1	114
16	A multi-ancestry genome-wide study incorporating gene–smoking interactions identifies multiple new loci for pulse pressure and mean arterial pressure. Human Molecular Genetics, 2019, 28, 2615-2633.	2.9	31
17	A Large-Scale Multi-ancestry Genome-wide Study Accounting for Smoking Behavior Identifies Multiple Significant Loci for Blood Pressure. American Journal of Human Genetics, 2018, 102, 375-400.	6.2	123
18	A Strategy for Discovery of Endocrine Interactions with Application to Whole-Body Metabolism. Cell Metabolism, 2018, 27, 1138-1155.e6.	16.2	58

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19	A Partial Loss-of-Function Variant in <i>AKT2</i> Is Associated With Reduced Insulin-Mediated Glucose Uptake in Multiple Insulin-Sensitive Tissues: A Genotype-Based Callback Positron Emission Tomography Study. Diabetes, 2018, 67, 334-342.	0.6	37
20	Interactions between genetic variation and cellular environment in skeletal muscle gene expression. PLoS ONE, 2018, 13, e0195788.	2.5	18
21	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. Nature Communications, 2017, 8, 14977.	12.8	169
22	Exome-wide association study of plasma lipids in >300,000 individuals. Nature Genetics, 2017, 49, 1758-1766.	21.4	470
23	Large meta-analysis of genome-wide association studies identifies five loci for lean body mass. Nature Communications, 2017, 8, 80.	12.8	147
24	Sequence data and association statistics from 12,940 type 2 diabetes cases and controls. Scientific Data, 2017, 4, 170179.	5.3	31
25	The genetic architecture of type 2 diabetes. Nature, 2016, 536, 41-47.	27.8	952
26	The genetic regulatory signature of type 2 diabetes in human skeletal muscle. Nature Communications, 2016, 7, 11764.	12.8	114
27	Genome-Wide Association Study of the Modified Stumvoll Insulin Sensitivity Index Identifies <i>BCL2</i> and <i>FAM19A2</i> as Novel Insulin Sensitivity Loci. Diabetes, 2016, 65, 3200-3211.	0.6	67
28	The transcription factor Prox1 is essential for satellite cell differentiation and muscle fibre-type regulation. Nature Communications, 2016, 7, 13124.	12.8	62
29	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. PLoS Genetics, 2015, 11, e1005378.	3.5	331
30	The effects of acute hyperinsulinemia on bone metabolism. Endocrine Connections, 2015, 4, 155-162.	1.9	32
31	Dyslipidemia and a reversible decrease in insulin sensitivity induced by therapy with 13-cis-retinoic acid. Diabetes/Metabolism Research and Reviews, 2001, 17, 391-395.	4.0	46
32	Subcutaneous adipose tissue expression of plasminogen activator inhibitor-1 (PAI-1) in nondiabetic and Type 2 diabetic subjects. Diabetes/Metabolism Research and Reviews, 2000, 16, 364-369.	4.0	20
33	Fatty acid transport protein-1 mRNA expression in skeletal muscle and in adipose tissue in humans. American Journal of Physiology - Endocrinology and Metabolism, 2000, 279, E1072-E1079.	3.5	81
34	Changes in Leptin Concentration during the Early Postnatal Period: Adjustment to Extrauterine Life?. Pediatric Research, 1999, 45, 197-201.	2.3	36
35	Is brain uptake of leptin in vivo saturable and reduced by fasting?. European Journal of Nuclear Medicine and Molecular Imaging, 1998, 25, 607-612.	6.4	27
36	Insulin-independent glucose transport regulates insulin sensitivity. FEBS Letters, 1998, 436, 301-303.	2.8	99

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