

Anna E Jonsson

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

2,165
citations

566801

15
h-index

752256

20
g-index

20
all docs

20
docs citations

20
times ranked

5345
citing authors

#	ARTICLE	IF	CITATIONS
1	Hyperglucagonemia in Pediatric Adiposity Associates With Cardiometabolic Risk Factors but Not Hyperglycemia. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2022, 107, 1569-1576.	1.8	7
2	The Arg82Cys Polymorphism of the Protein Nepmucin Implies a Role in HDL Metabolism. <i>Journal of the Endocrine Society</i> , 2022, 6, bvac034.	0.1	1
3	Genome-wide association study of circulating levels of glucagon during an oral glucose tolerance test. <i>BMC Medical Genomics</i> , 2021, 14, 3.	0.7	3
4	Fasting Plasma GLP-1 Is Associated With Overweight/Obesity and Cardiometabolic Risk Factors in Children and Adolescents. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021, 106, 1718-1727.	1.8	22
5	Associations of Mitochondrial and Nuclear Mitochondrial Variants and Genes with Seven Metabolic Traits. <i>American Journal of Human Genetics</i> , 2019, 104, 112-138.	2.6	106
6	Evidence of a liver- α cell axis in humans: hepatic insulin resistance attenuates relationship between fasting plasma glucagon and glucagonotropic amino acids. <i>Diabetologia</i> , 2018, 61, 671-680.	2.9	76
7	Metabolite ratios as potential biomarkers for type 2 diabetes: a DIRECT study. <i>Diabetologia</i> , 2018, 61, 117-129.	2.9	32
8	Common variants in the hERG (KCNH2) voltage-gated potassium channel are associated with altered fasting and glucose-stimulated plasma incretin and glucagon responses. <i>BMC Genetics</i> , 2018, 19, 15.	2.7	12
9	Genetic evidence of a causal effect of insulin resistance on branched-chain amino acid levels. <i>Diabetologia</i> , 2017, 60, 873-878.	2.9	119
10	Selection in Europeans on Fatty Acid Desaturases Associated with Dietary Changes. <i>Molecular Biology and Evolution</i> , 2017, 34, 1307-1318.	3.5	90
11	A Genome-Wide Association Study of IVGTT-Based Measures of First-Phase Insulin Secretion Refines the Underlying Physiology of Type 2 Diabetes Variants. <i>Diabetes</i> , 2017, 66, 2296-2309.	0.3	102
12	Heterogeneity in glucose response curves during an oral glucose tolerance test and associated cardiometabolic risk. <i>Endocrine</i> , 2017, 55, 427-434.	1.1	21
13	Insulin Resistance Is Accompanied by Increased Fasting Glucagon and Delayed Glucagon Suppression in Individuals With Normal and Impaired Glucose Regulation. <i>Diabetes</i> , 2016, 65, 3473-3481.	0.3	137
14	Genetic Correlation between Body Fat Percentage and Cardiorespiratory Fitness Suggests Common Genetic Etiology. <i>PLoS ONE</i> , 2016, 11, e0166738.	1.1	18
15	Gene-Environment Interactions of Circadian-Related Genes for Cardiometabolic Traits. <i>Diabetes Care</i> , 2015, 38, 1456-1466.	4.3	52
16	Genetic fine mapping and genomic annotation defines causal mechanisms at type 2 diabetes susceptibility loci. <i>Nature Genetics</i> , 2015, 47, 1415-1425.	9.4	365
17	Effects of Common Genetic Variants Associated With Type 2 Diabetes and Glycemic Traits on β - and β ² -Cell Function and Insulin Action in Humans. <i>Diabetes</i> , 2013, 62, 2978-2983.	0.3	85
18	Association between parental history of diabetes and type 2 diabetes genetic risk scores in the PPP-Botnia and Framingham Offspring Studies. <i>Diabetes Research and Clinical Practice</i> , 2011, 93, e76-e79.	1.1	19

#	ARTICLE	IF	CITATIONS
19	A Variant in the <i>KCNQ1</i> Gene Predicts Future Type 2 Diabetes and Mediates Impaired Insulin Secretion. <i>Diabetes</i> , 2009, 58, 2409-2413.	0.3	86
20	Clinical Risk Factors, DNA Variants, and the Development of Type 2 Diabetes. <i>New England Journal of Medicine</i> , 2008, 359, 2220-2232.	13.9	812