

# Claudia Ha Ting Tam

## List of Publications by Year in descending order

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Version: 2024-02-01

28  
papers

2,925  
citations

516561

16  
h-index

501076

28  
g-index

29  
all docs

29  
docs citations

29  
times ranked

7093  
citing authors

#	ARTICLE	IF	CITATIONS
1	Risk Associations of Glycemic Burden and Obesity With Liver Cancer—A 10-Year Analysis of 15,280 Patients With Type 2 Diabetes. <i>Hepatology Communications</i> , 2022, 6, 1350-1360.	2.0	13
2	Shortened Leukocyte Telomere Length Is Associated With Glycemic Progression in Type 2 Diabetes: A Prospective and Mendelian Randomization Analysis. <i>Diabetes Care</i> , 2022, 45, 701-709.	4.3	37
3	Clinical Predictors and Long-term Impact of Acute Kidney Injury on Progression of Diabetic Kidney Disease in Chinese Patients With Type 2 Diabetes. <i>Diabetes</i> , 2022, 71, 520-529.	0.3	6
4	Associations of the HOMA2-%B and HOMA2-%R with progression to diabetes and glycaemic deterioration in young and middle-aged Chinese. <i>Diabetes/Metabolism Research and Reviews</i> , 2022, 38, e3525.	1.7	12
5	Relative leucocyte telomere length is associated with incident end-stage kidney disease and rapid decline of kidney function in type 2 diabetes: analysis from the Hong Kong Diabetes Register. <i>Diabetologia</i> , 2022, 65, 375-386.	2.9	11
6	Multi-ancestry genetic study of type 2 diabetes highlights the power of diverse populations for discovery and translation. <i>Nature Genetics</i> , 2022, 54, 560-572.	9.4	250
7	Vitamin D Levels During Pregnancy Are Associated With Offspring Telomere Length: A Longitudinal Mother-Child Study. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2022, 107, e3901-e3909.	1.8	1
8	Development of genome-wide polygenic risk scores for lipid traits and clinical applications for dyslipidemia, subclinical atherosclerosis, and diabetes cardiovascular complications among East Asians. <i>Genome Medicine</i> , 2021, 13, 29.	3.6	18
9	Shortened relative leukocyte telomere length is associated with all-cause mortality in type 2 diabetes—analysis from the Hong Kong Diabetes Register. <i>Diabetes Research and Clinical Practice</i> , 2021, 173, 108649.	1.1	10
10	Determinants of penetrance and variable expressivity in monogenic metabolic conditions across 77,184 exomes. <i>Nature Communications</i> , 2021, 12, 3505.	5.8	49
11	Association between FGF19, FGF21 and lipocalin-2, and diabetes progression in PCOS. <i>Endocrine Connections</i> , 2021, 10, 1243-1252.	0.8	6
12	Long-term maternal cardiometabolic outcomes 22 years after gestational diabetes mellitus. <i>Journal of Diabetes Investigation</i> , 2020, 11, 985-993.	1.1	6
13	Shortened Relative Leukocyte Telomere Length Is Associated With Prevalent and Incident Cardiovascular Complications in Type 2 Diabetes: Analysis From the Hong Kong Diabetes Register. <i>Diabetes Care</i> , 2020, 43, 2257-2265.	4.3	31
14	Obesity, clinical, and genetic predictors for glycemic progression in Chinese patients with type 2 diabetes: A cohort study using the Hong Kong Diabetes Register and Hong Kong Diabetes Biobank. <i>PLoS Medicine</i> , 2020, 17, e1003209.	3.9	31
15	Identification of type 2 diabetes loci in 433,540 East Asian individuals. <i>Nature</i> , 2020, 582, 240-245.	13.7	282
16	SNPs in PRKCA-HIF1A-GLUT1 are associated with diabetic kidney disease in a Chinese Han population with type 2 diabetes. <i>European Journal of Clinical Investigation</i> , 2020, 50, e13264.	1.7	9
17	Circulating branched-chain amino acids and incident heart failure in type 2 diabetes: The Hong Kong Diabetes Register. <i>Diabetes/Metabolism Research and Reviews</i> , 2020, 36, e3253.	1.7	20
18	Progression of glucose intolerance and cardiometabolic risk factors over a decade in Chinese women with polycystic ovary syndrome: A case-control study. <i>PLoS Medicine</i> , 2019, 16, e1002953.	3.9	38

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19	Exome sequencing of 20,791 cases of type 2 diabetes and 24,440 controls. <i>Nature</i> , 2019, 570, 71-76.	13.7	248
20	Progression of diabetic kidney disease and trajectory of kidney function decline in Chinese patients with Type 2 diabetes. <i>Kidney International</i> , 2019, 95, 178-187.	2.6	105
21	IFITM3, TLR3, and CD55 Gene SNPs and Cumulative Genetic Risks for Severe Outcomes in Chinese Patients With H7N9/H1N1pdm09 Influenza. <i>Journal of Infectious Diseases</i> , 2017, 216, 97-104.	1.9	54
22	The genetic architecture of type 2 diabetes. <i>Nature</i> , 2016, 536, 41-47.	13.7	952
23	Genome-wide associations for birth weight and correlations with adult disease. <i>Nature</i> , 2016, 538, 248-252.	13.7	406
24	Genome-wide association studies in the Japanese population identify seven novel loci for type 2 diabetes. <i>Nature Communications</i> , 2016, 7, 10531.	5.8	149
25	Genetic and clinical variables identify predictors for chronic kidney disease in type 2 diabetes. <i>Kidney International</i> , 2016, 89, 411-420.	2.6	22
26	Familial Young-Onset Diabetes, Pre-Diabetes and Cardiovascular Disease Are Associated with Genetic Variants of DACH1 in Chinese. <i>PLoS ONE</i> , 2014, 9, e84770.	1.1	16
27	Genetic Variants of the Protein Kinase C- $\beta$ 1 Gene and Development of End-Stage Renal Disease in Patients With Type 2 Diabetes. <i>JAMA - Journal of the American Medical Association</i> , 2010, 304, 881.	3.8	58
28	Common Polymorphisms in MTNR1B, G6PC2 and GCK Are Associated with Increased Fasting Plasma Glucose and Impaired Beta-Cell Function in Chinese Subjects. <i>PLoS ONE</i> , 2010, 5, e11428.	1.1	65