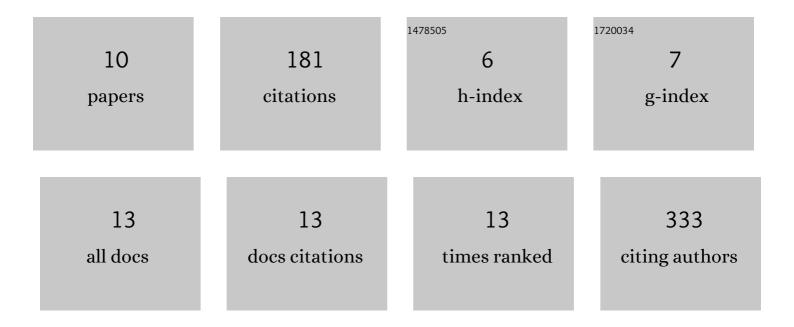
## Amna Khamis

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7999149/publications.pdf Version: 2024-02-01



ΔΜΝΙΛ ΚΗΛΜΙς

#	Article	IF	CITATIONS
1	Persistent or Transient Human $\hat{l}^2$ Cell Dysfunction Induced by Metabolic Stress: Specific Signatures and Shared Gene Expression with Type 2 Diabetes. Cell Reports, 2020, 33, 108466.	6.4	65
2	The expression of genes in top obesity-associated loci is enriched in insula and substantia nigra brain regions involved in addiction and reward. International Journal of Obesity, 2020, 44, 539-543.	3.4	38
3	Laser capture microdissection of human pancreatic islets reveals novel eQTLs associated with type 2 diabetes. Molecular Metabolism, 2019, 24, 98-107.	6.5	26
4	Maternal Glycemic Dysregulation During Pregnancy and Neonatal Blood DNA Methylation: Meta-analyses of Epigenome-Wide Association Studies. Diabetes Care, 2022, 45, 614-623.	8.6	19
5	Epigenome-Wide Association Study Reveals Methylation Loci Associated With Offspring Gestational Diabetes Mellitus Exposure and Maternal Methylome. Diabetes Care, 2021, 44, 1992-1999.	8.6	17
6	Histone deacetylase 9 promoter hypomethylation associated with adipocyte dysfunction is a statin-related metabolic effect. Clinical Epigenetics, 2020, 12, 68.	4.1	10
7	Chromatin 3D interaction analysis of the STARD10 locus unveils FCHSD2 as a regulator of insulin secretion. Cell Reports, 2021, 34, 108703.	6.4	4
8	Chromatin 3D Interaction Analysis of the <i>STARD10</i> Locus Unveils <i>FCHSD2</i> as a New Regulator of Insulin Secretion. SSRN Electronic Journal, 0, , .	0.4	0
9	Persistent or Transient Human β-Cell Dysfunction Induced by Metabolic Stress Associates with Specific Signatures and Shared Gene Expression of Type 2 Diabetes. SSRN Electronic Journal, 0, , .	0.4	0
10	Epigenetic changes associated with hyperglycaemia exposure in the longitudinal D.E.S.I.R. cohort. Diabetes and Metabolism, 2022, 48, 101347.	2.9	0