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Methylation of Specific CpG Sites in Genes is Affected by Hyperglycaemia in Type 2 Diabetic Patients

DOI: 10.1080/08820139.2019.1656227 Immunological Investigations, 2020, 49, 287-298.

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9	The inflammatory effect of epigenetic factors and modifications in type 2 diabetes. <i>Inflammopharmacology</i> , 2020 , 28, 345-362	5.1	6
8	Silencing cyclophilin A improves insulin secretion, reduces cell apoptosis, and alleviates inflammation as well as oxidant stress in high glucose-induced pancreatic Eells via MAPK/NF-kb signaling pathway. <i>Bioengineered</i> , 2020 , 11, 1047-1057	5.7	11
7	DNA methylation alterations and their potential influence on macrophage in periodontitis. <i>Oral Diseases</i> , 2020 ,	3.5	4
6	Molecular prospect of type-2 diabetes: Nanotechnology based diagnostics and therapeutic intervention. <i>Reviews in Endocrine and Metabolic Disorders</i> , 2021 , 22, 421-451	10.5	8
5	Phytochemicals as Potential Epidrugs in Type 2 Diabetes Mellitus. <i>Frontiers in Endocrinology</i> , 2021 , 12, 656978	5.7	4
4	Screening key genes and signaling pathways in COVID-19 infection and its associated complications by integrated bioinformatics analysis.		
3	Expression of obesity- and type-2 diabetes-associated genes in omental adipose tissue of individuals with obesity <i>Gene</i> , 2022 , 815, 146181	3.8	O
2	Bioinformatics Analysis of next generation sequencing data for Risk Prediction in Patients with Type 1 diabetes mellitus.		
1	Epigenetic Reprogramming of the Inflammatory Response in Obesity and Type 2 Diabetes. <i>Biomolecules</i> , 2022 , 12, 982	5.9	