

# THE ATP-BINDING CASSETTE TRANSPORTER A1 GENE POLYMORPHISMS AND TYPE 2 DIABETES MELLITUS

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Received on : 07-01-2022

Accepted on : 11-05-2022

## ABSTRACT

Insulin resistance (IR), secretion of insulin, and abnormalities of lipid metabolism are all markers of type 2 diabetes (T2DM), which is a progressive and complex metabolic disorder. Major risk factors for the development of T2DM were identified as genetic, environmental, and lifestyle factors. Several studies found that many genes contribute to T2DM susceptibility after glucose tolerance. Adenosine Binding Cassette Transporter Proteins 1 is a member of the ABC gene superfamily that is involved in cholesterol transport and HDL cholesterol (HDL-C) biosynthesis. Abnormal cholesterol metabolism, particularly high-density lipoprotein, has been related to genetic variations in the ABCA1 gene (HDL-C). Previous research suggested that ABCA1 gene polymorphisms may a hereditary risk factor for type 2 diabetes, along with lower HDL levels in various populations.

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**KEYWORDS:** T2DM, ABCA1 gene, Genetic variations, SNPs

## DIABETES MELLITUS

Diabetes mellitus is among the most prevalent chronic diseases worldwide. Hyperglycemia characterizes this metabolic disorder due to insulin insensitivity or insufficient levels. Long-term complications caused by Diabetic Mellitus are death and damage to different organs (1, 2). Diabetes currently affects 422 million people worldwide. This number is expected to increase to million by 2040 (<http://www.diabetesatlas.org>) (3).

Type two diabetes (T2DM) is a type of diabetes caused by insulin resistance or insufficiency. Monozygotic twins are more likely to be identical than dizygotic twins. Genetic and environmental conditions that affect beta-cell function and structure, resulting in altered tissue sensitivity to insulin, were found to play a significant role in Type 2 Diabetes Mellitus (T2DM). Cigarette smoking, lack of activity, low physical activity, poor diet, being overweight, and eating sugary foods all play a role in the progression of that disorder (4-6).

## ATP-BINDING CASSETTE TRANSPORTER A1 (ABCA1) TRANSPORTER PROTEIN

The ABCA1 genes have been identified in 1994 as a novel subclass A member of the ABC superfamily. ABCA1 is a 149-kb gene on chromosome 9 (q31) that

encodes a 240-kDa integral membrane protein with 2,261 amino acids (7). This protein forms an ATPase-dependent channel that transports lipids across the membrane (8). Cholesterol, phospholipids, and other lipophilic molecules are carried across the cell membrane through this channel.

The ATP-Binding Cassette Transporter A1 (ABCA1) is a symmetrical carrier with two transmembrane domains and two nucleotide binding domains (9, 10). Tangier disease is caused by mutations in the ABCA1 gene and has a genetic pattern characterized by autosomal dominance. Low HDL cholesterol levels in the blood and an accumulation of cholesteryl esters in the tissues are the characteristics of this disease (11). ABCA1 maintains intracellular cholesterol levels by removing excess cholesterol from cells (12). This lipid transporter is overexpressed in hepatocytes and non-hepatocytes, in accordance with its physiological function in these cells. To generate pre-HDL, ABCA1 transports all phospholipids and cholesterol from the cell to a lipid-free or lipid-poor apoA-I cell (13-17). ABCA1 is required for insulin secretion in beta cells; genetic changes in the ABCA1 gene disrupt high-density lipoprotein (HDL) metabolism, resulting in HDL deficiency (18). Cholesterol toxicity impairs beta cells' ability to secrete plasma and insulin, leading to

an increased risk of cardiovascular disorder and type-2 diabetes (19).

### **ABCA1 VARIANTS ARE ASSOCIATED WITH AN INCREASED RISK OF TYPE 2 DIABETES MELLITUS**

The Gene for the ATP-Binding Cassette Transporter A1 (ABCA1) contains at least 50 exons with lengths ranging from 33 to 245 bp (20). The ABCA1 gene has over 70 rare mutations, mostly in the apoA-I binding site. These uncommon and common variants may contribute to susceptibility to type 2 diabetes. They prevent the transporter from maturing and the protein from binding to ApoA-I. Individuals with ABCA1 SNPs may have lower high-density lipoprotein (HDL) or ApoA-I levels in their plasma (21).

Several researches have established the effect of these gene changes located in the exons, promoter, introns, and 3' untranslated area on transporter function and structure, thereby contributing to the emergence of Type-2 diabetes and its complications (ABCA1 Genetic variants and Type 2 Diabetes Mellitus and Its Health problems) (18).

### **THE PROMOTER REGION OF RS1800977 (C69T) GENETIC POLYMORPHISM**

Gene polymorphisms near the regulatory components of the ATP-Binding Cassette Transporter A1 Gene (ABCA1) proximal promoter area is related with an increased risk of developing more extreme hypertension (22). In this way, this variant of ABCA1 may play a central role in regulating the transcription of cholesterol trafficking, alone or in conjunction with several other genetic variations within the same gene or even other associated genes (23). A possible association between rs1800977 C69T variant and susceptibility to type 2 diabetes developments has also been described in various populations (24-27). The rs1800977 T allele and TT genotype, each thought to play a protective effect against T2DM, has been observed to be more prevalent in the subject of Turkey (25). A parallel result for the T allele has been demonstrated in a Saudi residents (26). As well as in the Malaysian residents (27). In individuals with and without prior myocardial injury, there is a significant association between the TT genotype and incidence of myocardial injury (24).

In a meta-analysis of the data from three eligible intervention studies, this polymorphism was suggested not to significantly impact the development of T2DM (25-27).

Cardiovascular disease account for the great majority of deaths among diabetics. High cholesterol and blood sugar levels in T2DM patients can result in blood

vessel obstruction, atherosclerosis, and cerebral infarction (28-30). HDL receptor ABCA1 is essential in the metabolism of HDL. Decreased plasma HDL-C levels have been connected to, and the ABCA1 transporter in the surface of blood vessel cells, where HDL helps extracts cholesterol particles from the extracellular space, can contribute to severe atherosclerosis. Thus, this transporter plays a protective role in aortic atherosclerosis (31-33).

The patient with diabetes has a more risk of atherosclerosis, cardiovascular disease, since their lowered HDL concentrations. Patients suffering from Type 2 diabetes have lipid profiles that are not normal, making them more prone to heart disease and cerebral infarction (34). The results of a national study in Japan that included 3432 diabetes and non-diabetic cerebrovascular disease, as well as 2070 healthy controls (adjusted for gender, hypertension, diabetes, and hypercholesterolemia) indicate that the T allele variant of the rs1800977 gene is a potential risk for atherothrombotic cerebral infarction (35).

The role of the rs1800977 is not to act alone. It must work with additional SNPs in the ABCA1 or with other genes to do its job. The population subgroup with the HMGCR rs3931914 profile and the MTHFR rs1800977 and ApoE rs2422493 SNPs was identified as one of the potential risks for Alzheimer's disease in Spain (36). In accordance with the findings of a study conducted in non-diabetic Dutch people with CHD, the rs1800977 polymorphism cooperates with the rs2422493, rs1800976, and rs2740483 polymorphisms that cause the ABCA1 gene (37). In addition, Rs1800977 interacts with 3-hydroxy-3-methylglutaryl-CoA gene polymorphisms (HMGCR). This enzyme converts HMG-CoA to mevalonic acid, which is important in cholesterol synthesis. Then, the cholesterol is secreted by the ABCA1 transporter. A combination of poor ABCA1 expression and over-expression of HMGCR causes an increase in cholesterol levels in the plasma and brain, resulting in atherosclerosis and Alzheimer's disease (38, 39).

In the atherogenesis as a consequence of type 2 diabetes, the role of rs1800977 in the ABCA1 promoter or other areas of the ABCA1 gene remains unknown. Many studies have attempted to correlate rs1800977 to lipid profiles (40). The greatest significant effects were observed in models that included age, the interaction of rs1800977 and rs708272, and the isomer of cholesteryl ester transfer protein (CETP). Cholesteryl esters are transported from HDL to other lipoproteins by this protein, which is essential for this process to take place (41).

### THE EXONIC REGION OF RS9282541 (R230C) GENETIC POLYMORPHISM

Rs9282541 is a rare non-synonymous SNP identified in the ABCA1 gene. As a result, the ABCA1 structure and function may be affected by the mutation. The substitution of C for T (CGT for TGT) results in the conversion of arginine to cysteine at the ABCA1 transporter's ED1 (23, 26). ApoA-I is triggered on the cell surface of the endoplasmic reticulum by directly interacting with the putative binding site in the ED-1 and ED-4 to stimulate ATP hydrolysis and subsequent cholesterol efflux from the ER (42). Clues that mutations in the ABCA1 transporter and ABCA1 stability sites are possible can lead to cellular lipid translocation deficits (43). ABCA1 is the gene that determines HDL levels. The rs9282541 gene, on the other hand, is associated with vulnerability to T2DM or high HDL levels in Mexican patients, but not in Colombians (18, 27, 44-47). The Mexican CT carriers were more likely to get T2DM early (44).

Other studies on the rs9282541 polymorphism have found no significant association with type 2 diabetes (27, 44, 46, 48). The rs9282541 is attributed to low HDL and high triglyceride levels and also with obesity risk. The CT/TT carriers had higher values of the HDL-3a/HDL-3b ratio as compared to the CC carriers (49-51). Flores-Dorantes et al. showed that this variation decreases HDL levels as a result of the gender-BMI interaction. Also, the examination of gender and genotype showed that men and CT carriers had the highest effect sizes (49).

Acua-Alonzo et al. discovered that Native Americans have a greater prevalence of the allele TT at the rs9282541 locus, indicating a potential selective advantage for higher HDL levels and other metabolic features. In this population, decreased HDL-C levels were found to be associated with a higher BMI. In vitro study revealed that the A allele resulted in a 27% decrease in cholesterol efflux. This finding shows how this protein shows functionality in this study (52). As a result, the rs9282541 T allele could be a key factor in spreading this ethnic group's resistance to metabolic diseases (52). However, another study did not support the use of traditional markers in admixture mapping. Evaluation of genome regions in the subject showed no deviation from the Native American genome. The researchers propose that socioeconomic status is a useful indicator of T2DM within that ethnicity (47).

Previous research has shown a correlation between rs9282541 and lifestyle factors. The association of rs9282541 with dietary carbohydrate and fat intake, gamma-glutamyl transpeptidase, alkaline phosphatase, adiponectin, and insulin resistance may

lower HDL levels in Mexicans (10, 53). Individuals with CT genotypes reacted better to dietary pattern treatment with soy protein, nopal, oat, and chia seed than those with TT genotypes. There was an even stronger therapeutic impact seen with more metabolic syndrome individuals (54, 55).

### THE EXONIC REGION OF RS2230806 (R219K) GENETIC POLYMORPHISM

There is a mutation in the ABCA1 gene called rs2230806. This mutation is in exon 7. There are two types of polymorphisms in this gene: one that changes the amino acid arginine to lysine (R219K) and one that changes a G nucleotide to an A nucleotide. It has a beneficial effect on the structure and stability of proteins (12, 23). The rs2230806 is associated with high high-density lipoprotein levels related to type 2 diabetes (56, 57). HDL-C, which is in our bloodstream, directly regulates the plasma's glucose concentration by stimulating insulin secretion by our pancreas and altering glucose uptake in our skeletal muscles (58).

Researchers found a possible correlation between a particular risk allele on an AA genotype and higher high-density lipoprotein levels in Asians (57). A study in Malaysia has shown that A allele and AA genotype carriers are more suitable for T2DM than alternative alleles or genotypes. This finding has been supported by Haghvirdizadeh et al (27). The low incidence of A allele was also found in diabetes patients with previous coronary heart disease (24).

In a study of Type 2 diabetes patients, the gene rs2230806 was also found to increase the chances of responding well to a drug that lowers blood sugar levels. Specifically, in Type 2 diabetes patients, the rs2230806 was associated with the antidiabetic drug response, rosiglitazone (59). ABCA1 is a highly expressed protein in the mouse pancreatic beta cells. Deficiency in this specific transporter results in cellular cholesterol accumulation and reduces insulin secretion. This results in a progressive impairment in glycaemic control. Rosiglitazone increased insulin sensitivity in the mice with T2DM by enhancing the expression of ABCA1 (60). In Chinese ethnicities, carriers of the r2 allele of rs2230806 have a stronger response to rosiglitazone than carriers of the A allele (61).

Other researchers showed that the A allele variant plays a protective effect against atherosclerosis (61), Coronary heart disease (CHD) (62), and ischemic stroke (63), but doesn't in Alzheimer's disease (AD) (64, 65).

Studies show that high HDL levels decrease atherosclerosis chances by increasing the endothelium's oxidation and restoration and by



exerting anti-inflammation and anti-apoptotic effects (65). Other studies showed that low levels of High-Density Lipoprotein (HDL) are closely associated with disease progression of Type 2 Diabetes Mellitus (T2DM) (66). They may accelerate and promote atherosclerosis, myocardial infarction, and Coronary heart disease (CHD). The rs2230806 is strongly correlated with serum high-density lipoprotein (HDL) levels and response to antidiabetic drugs in T2DM (67). The ABCA1 is an efflux transporter that is critical for regulating HDL function and cholesterol homeostasis in the body; identifying the ABCA1 gene and its monogenic mutations has become a new strategy for earlier diagnosis and prognosis of 2 diabetes(68).

## CONCLUSION

A number of studies have shown that having the ABCA1 gene polymorphism may increase one's threat of developing type 2 diabetes and its complications, and also conditions that contribute to the development of type 2 diabetes, such as lower serum high-density lipoprotein (HDL) levels type 2 diabetes and/or heart disease (CHD) (69). To recognize and better understand the role of these SNPs, we summarized the existing evidence by focusing more on rs2230806 (R219K), rs1800977 (C69T), and rs9282541 (R230C) variants. Specific variants are associated with reduced cholesterol efflux, This raises the risk for type 2 diabetes and its consequences and the medication's responsiveness in the patients.

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