

## Original Article

# Genetic polymorphisms of *ADIPOQ* and osteonecrosis of the femoral head risk in Chinese population

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Received January 4, 2016; Accepted March 18, 2016; Epub August 1, 2016; Published August 15, 2016

**Abstract:** Aims: The objective of this study was to investigate the effect of adiponectin gene (*ADIPOQ*) polymorphisms on osteonecrosis of the femoral head (ONFH) susceptibility. Methods: Polymerase chain reaction (PCR) was used to detect the genotypes of *ADIPOQ* polymorphisms in 104 patients with ONFH and 131 controls matched by age and gender with the former.  $\chi^2$  test were conducted to compare the frequency difference of genotypes and alleles in *ADIPOQ* polymorphisms between the above two groups and detect the Hardy-Weinberg equilibrium (HWE). Odds ratio (OR) with 95% confidence interval (95% CI) was calculated to represent the association intensity of gene polymorphisms with disease. The linkage disequilibrium (LD) and haplotype of these two polymorphisms were analyzed by Haploview. Results: In *ADIPOQ* polymorphisms, neither genotype nor allele of rs3774261 was found the significant frequency difference between the case and control groups ( $P>0.05$ ). However, TT genotype of rs1063537 was correlated to the increased risk of ONFH development (OR=2.66, 95% CI=1.04-6.78), T allele had also a 1.57 times risk to suffer from ONFH, compared with C allele (OR=1.57, 95% CI=1.04-2.36). There was strong LD between these two polymorphisms. Conclusion: *ADIPOQ* rs1063537 polymorphism is associated with the onset of ONFH, but not rs3774261 in Han population based on our study. A-T haplotype might be a risk factor in ONFH.

**Keywords:** *ADIPOQ*, osteonecrosis of the femoral head risk, polymorphism, haplotype

## Introduction

Osteonecrosis of the femoral head (ONFH) is a devastating bone disease in femoral head that always induces arthropathy of the hip joint [1]. This disease is characterized by the interruption of local blood supply and the death of osteocytes and bone marrow cells, usually occurring in the age of 20-50 years old [2, 3]. In clinical, ONFH is divided into traumatic- and non-traumatic ONFH and the latter has been studied widely in pathogenesis, prevention and treatment [4-6]. Non-traumatic ONFH is a complex multifactorial disease and the known risk factors involve corticosteroid misuse, alcohol abuse, thrombotic interference and idiopathic origins [7, 8]. Among of these causes, genetic factors play a vital role in ONFH development, especially the single nucleotide polymorphism (SNP) of gene [9-11].

Adiponectin is firstly found by Scherer et al. in 1995 [12] and then is proved to be secreted by adipose tissues. As a kind of hormone, it is encoded by *ADIPOQ* gene located on chromo-

some 3q27 [13]. It can regulate a number of metabolic processes, such as glucose regulation and fatty acid breakdown. In previous studies, *ADIPOQ* has been reported to participate in the generation of various diseases, type 2 diabetes mellitus (T2DM), cardiovascular disease (CVD), insulin resistance and obesity are the common diseases in clinical [14-16]. Recently, adiponectin is also found to be associated with the function of endothelial cells and monocyte-macrophages and may be play an important role in restraining the occurrence of atherosclerosis [17]. In bone metabolism, the formation of osteoblasts and osteoclasts is considered to be accelerated by adiponectin which can promote bone matrix mineralization, too. In addition, adiponectin is also involved in lipid metabolism and dyslipidemia is a leading cause of ONFH development. What's more, the studies show that the genetic variant of *ADIPOQ* influences the level of adiponectin and lipoproteins [18].

But the publications about the role of *ADIPOQ* polymorphism with ONFH are few nowadays. Therefore, in this research, the association

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**Table 1.** The primer sequences of *ADIPOQ* polymorphisms

SNP	Primer sequences	Location	Length
rs3774261	Fro. 5'-GCTCAGTCCTGCCTTTGG-3'	Intron 2	239 bp
	Rev. 5'-CCTCCTCTATTCTGCCTAC-3'		
rs1063537	Fro. 5'-GTTGAGGTAGTTGATGGTGGTA-3'	3'UTR	270 bp
	Rev. 5'-CCTGGCAACAGAGGGAAA-3'		

Note: SNP: Single nucleotide polymorphism; 3'UTR: 3'untranslated region.

**Table 2.** The relationship between *ADIPOQ* gene polymorphisms and ONFH susceptibility

SNP	Genotype/allele	Case/control, n	OR (95%CI)	P	P <sub>HWE</sub>
rs3774261	AA	36/41	1.00 (Ref.)	-	0.12
	AG	56/72	0.89 (0.50-1.56)	0.68	
	GG	12/18	0.76 (0.32-1.79)	0.53	
	A	128/154	1.00 (Ref.)	-	
	G	80/108	0.89 (0.61-1.29)	0.54	
rs1063537	CC	52/79	1.00 (Ref.)	-	0.58
	CT	38/44	1.31 (0.75-2.29)	0.34	
	TT	14/8	2.66 (1.04-6.78)	0.04	
	C	142/202	1.00 (Ref.)	-	
	T	66/60	1.57 (1.04-2.36)	0.03	

Note: HWE: Hardy-Weinberg equilibrium.

between *ADIPOQ* polymorphisms and ONFH was explored and two SNPs were selected in *ADIPOQ* gene, that is, rs3774261, rs1063537. In addition, the linkage disequilibrium was also paid attention between these two SNPs.

### Materials and methods

#### Subjects

104 patients with ONFH and 131 healthy people were as the case and control groups respectively to conduct this case-control study. The cases were all non-traumatic ONFH diagnosed by pathobiology in orthopedics department of Shandong Provincial Western Hospital from 3, 2012 to 7, 2013, consisted of 63 males and 41 females with the mean age of 46.76±10.49. All patients didn't accept the surgery before collecting the blood sample. The controls were also from the hospital similar to the cases at the same time, including 72 males and 59 females. They were confirmed to be healthy by the physical examination center of the hospital and the age range was from 34 to 72 years old.

All subjects were Chinese Han population from Shandong province and had no relationship by

blood each other. The study was accepted by the Research Ethics Committee of Shandong Provincial Western Hospital and participants were informed the study process and the detailed study content. Written consents must be obtained from all subjects before starting the whole study.

#### DNA extraction

2 ml peripheral venous blood from every subject was collected and put into corresponding tubes for blood specimen collection, and stored at -80°C. Genomic DNA was extracted from peripheral blood using an AxyPrep Blood Genomic DNA Miniprep Kit (Axygen Biosciences, Union, CA, USA).

#### Genotyping

In this study, the genotyping of *ADIPOQ* gene polymorphisms was performed by polymerase chain reaction (PCR) and sequencing. The polymorphisms of rs3774261, rs1063537 were selected in *ADIPOQ* and the corresponding PCR primers were designed using Primer Premier 5.0 software. The primer sequences and their detailed information were listed in **Table 1**. 20.0 µl PCR system was used, including 1.0 µl DNA template, each 0.5 µl of forward and reverse primers, 10.0 µl PCR Mix and 8.0 µl sterile ddH<sub>2</sub>O. The PCR program was as follows: initial denaturation at 95°C for 2 min, 32 cycles with denaturation at 95°C for 30 s, annealing at 58°C for 45 s and extension at 72°C for 45 s, final extension at 72°C for 7 min. And then, PCR products were detected the quality and concentration.

The eligible PCR products were sequenced to determine the genotype of every individual in Shanghai Sangon Biotech Co., Ltd.

#### Statistical analysis

The relative data were processed in PASW Statistics 18.0 software. Hardy-Weinberg disequilibrium (HWE) in controls was evaluated based on *ADIPOQ* polymorphisms. Age distribution between the cases and controls were

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**Table 3.** The detected results of linkage disequilibrium and haplotype between two polymorphisms of ADIPOQ

Haplotype SNP1-SNP2	Case (%)	Control (%)	OR (95% CI)	P
G-C	80 (38.46)	108 (41.22)	1.00 (Ref.)	-
A-C	62 (29.81)	94 (35.88)	0.89 (0.58-1.37)	0.60
A-T	66 (31.73)	60 (22.90)	1.49 (0.94-2.34)	0.09

Note: SNP1: rs3774261; SNP2: rs1063537.

assessed by t-test and the  $\chi^2$  test was used to measure the differences of genotype, allele, haplotype and the other indexes between the above two groups. The odds ratio (OR) and 95% confidence interval (CI) were calculated to evaluate the association strength between gene polymorphisms and disease. The statistical significant difference was set at  $P < 0.05$ . The linkage disequilibrium (LD) and haplotype were analyzed referring to Haploview software.

### Results

#### *Characteristics of patients and controls*

A total of 235 subjects were enrolled and contained 104 ONFH patients and 131 healthy persons. The average age in the control group was  $46.76 \pm 10.49$ , which showed no significant difference with the average age of  $47.52 \pm 11.27$  in the case group ( $P = 0.64$ ). In the case group, nearly 61% subjects were males and the percentage of males in the control group was about 55%. So there was no obvious difference between two groups in gender ( $P = 0.39$ ).

#### *HWE test and the association analysis of ADIPOQ polymorphisms with ONFH*

The genotype distributions of ADIPOQ polymorphisms in the control group were detected to conform to HWE ( $P > 0.05$ ) in **Table 2**, the results showed that our study population was consistent with the standard of the selection and possessed the representativeness of group.

In views of the association analysis between ADIPOQ polymorphisms with ONFH susceptibility, the results were listed in **Table 2**. The genotype distribution of rs3774261 had no significant difference between two groups and this polymorphism might be not directly associated with ONFH risk ( $P = 0.68, 0.53$  respectively). The effect of rs3774261 allele on ONFH was not significant ( $P = 0.54$ ). Referring to rs1063537, differing from rs3774261, TT genotype fre-

quency in cases was significantly different from the controls ( $P = 0.04$ ) and the carriage of TT genotype was a risk factor for ONFH occurrence (OR=2.66, 95% CI=1.04-6.78). T allele frequency was also higher in cases than that of in controls ( $P = 0.03$ ) and it was directly correlated to the development of ONFH (OR=1.57, 95% CI=1.04-2.36).

#### *Association between haplotypes and ONFH susceptibility*

We found that there was the strong LD between rs3774261 and rs1063537 polymorphisms from Haploview software ( $D' = 1.0, r^2 = 0.256$ ). Three haplotypes were detected in our population, namely G-C, A-C, A-T and their frequencies in the case and control groups were 38.46%, 29.81%, 31.73% and 41.22%, 35.88%, 22.90%, respectively. The detailed information of haplotypes was showed in **Table 3**. In these haplotypes, their frequency differences between two study groups didn't reached the significant level of  $P < 0.05$ , but A-T haplotype might be a risk factor for ONFH (OR=1.49, 95% CI=0.94-2.34).

### Discussion

Recently, OPG-RANKL-RANK system is found to play a fatal role in bone destruction [19, 20]. Receptor activator of NF- $\kappa$ B ligand (RANKL) mainly promotes the differentiation, maturation, activation of osteoclasts and meanwhile inhibits their apoptosis through combining with receptor activator of NF- $\kappa$ B (RANK) in the surface of preosteoclasts [21]. Osteoprotegerin (OPG) is a member of tumor necrosis factor superfamily and it can effectively prevent bone loss mediated by RANKL in the case of integrating RANKL. Bone resorption is also modulated by the balance of OPG and RANKL. Studies show that a number of hormones, cytokines affect bone metabolism directly or indirectly through regulating the OPG-RANKL-RANK pathway, which leads to bone metabolism diseases, such as ONFH.

Adiponectin is a kind of hormone secreted by adipocytes, consisted of 244 amino acids. In the primary sequence of adiponectin, four functional domains are revealed, namely a signal peptide containing 18 amino acids, a non-helical domain of N-terminal, a collagen repetitive

sequence containing 22 amino acids and a globular domain in C-terminal. It is persisted in tripolymer, hexamer and multimer status within plasma [22]. Adiponectin regulates bone metabolism by the way of affecting the expression of OPG, RANKL, RANK mainly. On the one hand, it can promote the proliferation, differentiation of osteoblasts by the adiponectin receptor 1 (AdipoR1)/JNK, AdipoR1/p38 (mitogen-activated protein kinase, MAPK) pathways [23]. On the other hand, recombinant adiponectin accelerates the expression of RANKL and inhibit OPG expression through AdipoR1/p38 MAPK pathway in osteoblasts [24]. Therefore, adiponectin plays an important role in the formation of osteoclasts and osteoblasts and it realizes the regulation for osteoclasts on the basis of osteoblasts. In addition, adiponectin regulates the blood glucose, blood fat and insulin-resistance by the way of PPAR (peroxisome proliferator activated receptor) and AMPK (AMP-activated protein kinase) signal transduction pathway, so it also plays a role in lipid metabolism.

ONFH occurrence is closely associated with lipid metabolism disorders. In the study of Cui et al., the genetic polymorphisms of apolipoprotein A5 (ApoA5) which may result in lipid metabolism disorders are associated with the occurrence of ONFH in Chinese population [25]. In order to determine the association of hypercholesterolemia with idiopathic ONFH risk, Moskal et al. detect cholesterol levels in serum of patients with ONFH, which suggests that hypercholesterolemia may have a systemic influence on the pathophysiology of ONFH [26]. In the meanwhile, the role of OPG-RANKL-RANK pathway in ONFH development has been proved in previous study. Jiang et al. have verified this conclusion in rats, they found that *Achyranthes bidentata* extract (ABE) can inhibit steroid-induced ONFH and remiss the bone deterioration induced by steroid through down-regulating OPG and up-regulating RANK and RANKL in OPG-RANKL-RANK system [27]. So adiponectin which can regulate lipid metabolism and OPG-RANKL-RANK pathway may be closely correlated to the generation and development of ONFH.

In present study, the genetic polymorphisms of *ADIPOQ* were analyzed the association with ONFH susceptibility. Two common SNPs were selected, namely rs3774261, rs1063537 and

the former is located in intron 2, the latter is in 3'-UTR. In the genotypes and alleles analyses of rs3774261, no obvious difference was found between the case and control groups, which indicated that this polymorphism was not directly associated with the onset of ONFH in our study population. Differently, the genotype distribution of rs1063537 had a significant difference between the two groups, that is, TT genotype carriers has a larger risk to suffer from ONFH than people carrying CC genotype. T allele of rs1063537 was also a risk factor in this study population. The strong LD was found between these two polymorphisms of *ADIPOQ* and three haplotypes were analyzed the effect on ONFH, but all haplotype distribution differences between the case and control groups didn't reached a statistical significance and maybe A-T haplotype is a risk factor in the development of ONFH based on a larger study population.

All in all, *ADIPOQ* polymorphisms affect the level of adiponectin in humans and them result in a series of biological processes disorders, which causes various diseases. In our study, *ADIPOQ* rs1063537 is associated with the onset of ONFH, but not rs3774261 in Chinese Han population based on our study. But the results need to be verified due to the small sample size, ignoring the environmental factors and region difference in gene polymorphism. In the future, studies with well design, large samples are conducted to deeply reveal the effect of *ADIPOQ* polymorphisms on ONFH pathobiology.

### Disclosure of conflict of interest

None.

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