

Mutations in the coding regions of the hepatocyte nuclear factor ϵ alpha in Iranian families with maturity onset diabetes of the young

Taghavi, S.M.^a, Fatemi, S.S.^b, Rafatpanah, H.^b, Ganjali, R.^b, Tavakolafshari, J.^b, Valizadeh, N.^c

^a Internal Medicine Department, Ghaem Hospital and Endocrine Research Center, **Mashhad University of Medical Sciences**, Parastar st, Ahmad abad blvd, **Mashhad**, Iran

^b Immunogenetics Department, Immunology Research Center, Bu-Ali Research Center, **Mashhad University of Medical Sciences**, Bu Ali Square, Ferdowsi Square, **Mashhad**, Iran

^c Microbiology and Virology Department, Bu-Ali Research Center, **Mashhad University of Medical Sciences**, Bu Ali Square, Ferdowsi Square, **Mashhad**, Iran

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Abstract

Background: Hepatocyte nuclear factor ϵ (HNF ϵ) is a nuclear receptor involved in glucose homeostasis and is required for normal β cell function. Mutations in the HNF ϵ gene are associated with maturity onset diabetes of the young type 1 (MODY1). The aim of the present study was to determine the prevalence and nature of mutations in HNF ϵ gene in Iranian patients with a clinical diagnosis of MODY and their family members. Twelve families including 20 patients with clinically MODY diagnosis and 11 members of their family were examined using PCR-RFLP method and in case of mutation confirmed by sequencing techniques. Fifty age and sex matched subjects with normal fasting blood sugar (FBS) and Glucose tolerance test (GTT) were constituted the control group and investigated in the similar pattern. Single mutation of V100M in the HNF ϵ gene was detected. This known mutation was found in 1 of 20 patients and 2 of 11 individuals in relatives. Fifty healthy control subjects did not show any mutation. Here, it is indicated that the prevalence of HNF ϵ mutation among Iranian patients with clinical MODY is considerable. This mutation was present in 26.7% of our patients, but nothing was found in control group. In the family members, 2 subjects with the age of ≤ 20 years old carried this mutation. Therefore, holding this mutation in this range of age could be a predisposing factor for developing diabetes in future. © 2019 Taghavi et al; licensee BioMed Central Ltd.

Reaxys Database Information

Indexed Keywords

EMTREE drug terms: hepatocyte nuclear factor ϵ ; HNF ϵ A protein, human; glucose; hepatocyte nuclear factor ϵ alpha

EMTREE medical terms: adolescent; adult; article; Asian; blood; case control study; ethnology; female; genetic predisposition; genetics; glucose blood level; glucose tolerance test; human; Iran; male; mutation; non insulin dependent diabetes mellitus; nucleotide sequence; onset age; open reading frame; pedigree; phenotype; polymerase chain reaction; risk factor; clinical article; controlled study; family; gene mutation; gene sequence; maturity onset diabetes mellitus; pathogenesis; prevalence; restriction fragment length polymorphism

MeSH: Adolescent; Adult; Age of Onset; Asian Continental Ancestry Group; Blood Glucose; Case-Control Studies; Diabetes Mellitus, Type 1; DNA Mutational Analysis; Female; Genetic Predisposition to Disease; Glucose Tolerance Test; Hepatocyte Nuclear Factor ϵ ; Humans; Iran; Male; Mutation; Open Reading Frames; Pedigree; Phenotype; Polymerase Chain Reaction; Risk Factors; Young Adult

Medline is the source for the MeSH terms of this document.

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