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Research Article

Clinical and Laboratory Characteristics of MODY (Maturity Onset Diabetes of Young) Cases, Genetic Mutation Spectrum and Phenotype-genotype Relationship

Özsu E et al. MODY in Childhood

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What is already known on this topic?

Maturity onset diabetes of the young (MODY) is the term used to describe a group of inherited, non-autoimmune forms of diabetes mellitus (DM). MODY diagnosis is often made at a young age (under 25 years of age), negative pancreatic autoantibodies, low insulin requirement, family history of autosomal dominant diabetes, no history of obesity and no history of diabetic ketoacidosis (DKA).

What this study adds?

This large-series national study showed that the diagnostic criteria in the prediagnostic process of MODY should be reconsidered in the presence of DKA, the presence of antibody positivity and the absence of family instory in patients with genetically diagnosed MODY.

Objective: M a t u r i t y - o n s e t d i a b e t e s o f th e yo u n g (MODY) occurs due to mutations in genes involved in pancreatic beta cell function and insulin secretion, has heterogeneous clinical and laboratory features, and account for 1-5% of all diabetes cases. The prevalence and distribution of MODY subtypes vary between countries. The aim of this study was to evaluate the clinical and laboratory characteristics, mutation distribution, and phenotype-genotype relationship in a large case series of pediatric Turkish patients genetically diagnosed with MODY.

Methods: MODY cases from 14 different pediatric endocrinology departments were included. Diagnosis, treatment, follow-up data, and results of genetic analysis were evaluated.

Results: A total of 224 patients were included, of whom 101 (45%) were female, and the mean age at diagnosis was 9.4 ± 4.1 years. Gene variant distribution was: 146 (65%) *GeK*; 43 (19%) *HNF1A*; 8 (3.6%) *HNF4A*, 8 (3.6%) *KLF11* and 7 (3.1%) *HNF1B*. The remaining 12 variant distribution was: 146 (65%) GCK, 42 (19%), *HNP1A*; 8 (5.6%) *HNP1A*, 8 (5.6%) *KLP11* and 7 (5.1%) *HNP1B*. The remaining 12 variants were: *PDX* (n=1), *NEUROD*: (n=3), *CLL* (n=1), *INS* (n=3), *ABCC8* (n=3) and *KJNC11* (n=1). Of the cases, 197 (87.9%) were diagnosed with incidental hyperplycentia, 6 with ketosis (7%) and 7 (3%) with diabetic ketoacidosis (DKA), while 30% presented with classical symptoms of diabetes. Two-hundred (89%) had a family history of diabetes. Anti-GAD antibody was detected in 13 cases, anti-islet antibody in eight and ant insulin antibody in four. Obesity was present in 16. Distribution of therapy was: 158 (71%) diet only; 23 (11%) intensive insulin treatment, 17 (7.6%) sulfonylureas; 10 (4.5%) metformin; and 6 (2.7%) insulin and oral antidiabetic treatment. **Conclusion:** This was the largest genetically diagnosed series from Turkey. The most common gene variants were *GCK* and *HNF1A* with much lower proportions for other MODY types. Hyperglycemia was the most common presenting symptom while 11% of patients had diabeter associated autoantibodies and 7% were obese. The majority of patients received dietary management only. Keywords: Childhood, MODY, Diagnosis

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Introduction

Maturity onset diabetes of the young (MODY) is the term used to describe a group of inherited, non-autoimmune forms of diabetes mellitus (DM). MODY was initially thought to be relatively rare but it has become apparent that it is more common than first thought. Despite increasing awareness of MODY, it has been estimated that up to 1-5 % of MODY cases remain undiagnosed. Since the clinical spectrum is variable due to the range of genetic defects, there may be diagnostic delays, and some cases may not be diagnosed until adulthood. Furthermore, cases may sometimes be misdiagnosed as Type 1 (T1DM) or Type 2 DM (T2DM) (1-3)

Given the natural history of MODY, that there is cost-effective treatment, and the potential impact on more than one family member, it is essential to diagnose these cases correctly. The presence of family history, clinical and physical examination findings incompatible with T1DM or T2DM, and negative diabetes autoantibodies, when present strengthen the diagnosis of MODY (3). In addition, urinary c peptide/creatinine ratio >0.2 nmol/mmol has high specificity and sensitivity in differentiating MODY from T1DM (4).

Genetic testing is important for definitive diagnosis and identification of other family members with the same variant. Mutation analyses classify the type of MODY and this in turn informs about the expected clinical courses and guides treatment planning. In patients with HNF4A and HNF1A mutations, forms of MODY which are responsive to sulphonylurea therapy, it is important to change treatment regimens from insulin treatment, as low-dose sulphonylurea treatment may provide good glycaemic control for years and, in patients with GCK mutations, will prevent unnecessary insulin treatment. Mutation screening is also crucial for asymptomatic relatives with a 50% risk of inheriting the mutation.

The distribution of MODY types differs between countries. Although HNF1A MODY is reported to be the most common type globally, there are countries where GCK MODY is observed more frequently (5). Therefore, knowing which of the MODY types is more common in one's own country is important. Determination of the MODY type will provide the most accurate approach for the pediatric endocrinologist to manage MODY patients. Moreover, managing clinicians should understand the phenotype-genotype relationship in MODY. The aim of this study was to investigate the distribution of MODY types in a pediatric population from Turkey and to investigate the phenotypegenotype relationship.

METHODS

The study included pediatric patients with genetically diagnosed MODY from 14 Pediatric Endocrinology and Diabetes Departments in Turkey. Ethical approval was obtained from Ankara University Faculty of Medicine with the number 12-112-20. The National Paediatric Endocrinology and Diabetes Association supported the study. An electronic case registration form was created in the Cedd-net data system of the association, and centers were asked to record their data using this system. The data entry took one year. Retrospective medical history (at leastthree generations of family history), clinical characteristics (age at diagnosis, year of diagnosis, duration of diabetes, presence of classical diabetes symptoms), laboratory results (blood glucose, insulin, C-peptide, glycated hemoglobin (HbA1c) levels, lipid profile, presence of acidosis at diagnosis, presence of autoantibodies at diagnosis, and reserve insulin values) and genetic data were transferred from electronic case record forms to Microsoft Excel (Microsoft Inc., Redmond, USA) and SPSS, version 20.0 (IBM Inc. Armonk, NY, USA) files for statistical evaluation.

Informed consent was obtained from the parents/guardians of the patient from all centers.

Patients with a positive family history in which at least two consecutive generations were affected, early-onset hyperglycemia, negative diabetes autoantibodies, and no classic type 1 or type 2 DM features (not insulin-dependent or not requiring insulin treatment even after three years of insulin treatment, or having insulin reserve with serum C-peptide level >0.60 ng/mL) were sough from the clinics. Genetic analysis was performed with a presumptive prediagnosis of MODY.Patients had already had genetic analyses and they were genetically diagnosed.

Body mass index (BMI) values of the cases were evaluated according to the Olcay Neyzi data adjusted for age and sex, and cases with $BMI > 95^{th}$ percentile were considered obese. Ketonemia was defined as a blood ketone level of 1 ng/dL, and diabetic ketoacidosis (DKA) was defined as a blood glucose level above 200 mg/dL with blood pH <7.3 or HCO₃ <15 mmol/L. cutoantibody status and reserve insulin values were recorded at diagnosis.

Mutation Interpretation

Genetic analyses of the cases were performed by next-generation sequencing and screening for 14 currently known mutations. Each patient was included in the study with the genetic diagnosis of MODY in the unit where they were followed.

MODY DNA sequence analysis panel was performed on each patient. All results were assessed by genetic specialist according to ACGM classification. Results, which were pathogenic or likely pathogenic were accepted as positive.

Statical Methods

Descriptive statistics were calculated for all 224 patients taking part in the study using SPSS, version 21.0 (IBM Corp., Armonk, NY, USA). Data were expressed as the mean ± SD, median [minimum; maximum]. The Kolmogorov-Smirnov test evaluated the normality of variables. Descriptive analyses were presented using means and standard deviations for normally distributed variables. A p value of less than 0,05 showed a statistically significant result in all analyses a statistically significant result in all analyses.

RESULTS

General Features

General Features A total of 224 cases were included in the study. Of these 101 (45%) were female, and the mean age at diagnosiswas 9.4 ± 4.1 and ranged widely from 0.2 to 17 years. On MODY genotyping, 146 (65%) cases were diagnosed as *GCK*, 43 (19%) cases were diagnosed as *HNF1A*, 8 (3.6%) cases were diagnosed as *HNF4A* and o turner eight with *KLF11* while 7 (3.125%) patients had an *HNF1B* variant. In the remaining twelve cases there were three (1.3%) each with *NEUROD1*, *INS* and *ABCC8* mutations while the last three had PDX (n=1), *CEL* (n=1) and *KJNC11* (n=1) mutations, respectively (Figure 1). Most of the cases (n=197, 87.9%) were diagn sed with incidental hyperglycemia, while 16 (7%) had ketosis and 7 (3%) had DKA; 30% had diabetes symptoms. There were 200 cases (29%) with a family history of DM; 93 (41%) had diabetes in two generations and 84 (37%) had diabetes in three generations. Relatives of 162 patients with diabetes were under the age of 40. In 24 patients, no family history was found. A proportion of the patients, where netically diagnosed MODY presented with type 1 DM-associated autoantihodies including 13

had diabetes in three generations. Relatives of 162 patients with diabetes were under the age of 40. In 24 patients, no family history was found. A proportion of the patients with genetically diagnosed MODY presented with type 1 DM-associated autoantibodies, including 13 (5.8%) with anti-GAD, 8 (3.6%) with anti-islet and 4 (1.8%) with anti-insulin antibodies. These caseswere in HNF4A, GCK, HNF1A, KLF11, KJNC11 MODY (Typ. 1,2,3, 1,2 MODY) types. Sixteen of the caseswere obese. Clinical and laboratory characteristics of MODY types are given in Table 1. While 158 (71%) of the patients were followed up with diet alone, 23 (11%) received intensive insulin therapy, 17 (7.6%) received sulfonylurea, 10 (4.5%), received a metformin, and 6 (2.7%) received intensive antibiabetic therapy (Table IIa-IIb). Seven patients

presented with diabete ketoacidosis or developed DKA during follow-up and these patients' genotypes were: HNF4A (n=2, %0.89), HNF1A (n=2, %0.89) HNF1B (n=1, %0.44), NEUROD (n=1, %0.44), and ABCC8 (n=1, %0.44) MODY types. The clinical and laboratory characteristics of the patients who presented with diabetic ketoacidosis are shown in Table III. An INS gene mutation was found in one patient who developed DKA during follow-up.

The characteristics of patients receiving intensive insulin treatment are shown. Insulin treatment was needed in 35 (15%) of the patients at diagnosis or follow-up. Of these patients, 13 (%5) had HNF1A and 6 (2%) had HNF1B MODY. The characteristics of MODY patients who used insulin treatment are given in Table IV.

Genotype-Phenotype Characteristics and Treatment according to MODY types

GCK MODY

GCK mutation was found in 146 (65%) of 224 patients. The mean±SD age was 8.8±3.9 years. Although all of the GCK MODYs were diagnosed with incidental hyperglycemia, 23 of them had typical symptoms of T1DM. In 50% of the cases, there was a family history of diabetes in at least two generations. At the time of diagnosis, the mean blood glucose level was 125±21 mg/dL, and c-peptide was 1.4±1.3 mg/dL. HbA1c ranged from 5.4-10.2% with a mean of 6.3±0.3%. Anti-GAD (n=6), anti-insulin (n=1) and islet cell antibodies (n=5) were positive. Comorbidities included arachnoid cyst, short stature, asthma, cystic fibrosis, precocious puberty, connective tissue diseases, dyslipidaemia and undescended testis. While 135 patients werefollowed up with the diet, six received oral anti-diabetic therapy, and five received intensive insulin treatment.

HNF1A-MODY

Nearly a fifth (19.2%, n=43) of the patients had this mutation. The mean±SD age at diagnosis was 10.6±4.3 years. Typical symptoms of T1DM were present in five patients, and 31 patients presented with incidental hyperglycemia. Thirty-five patients had a family history. At diagnosis, mean blood glucose was 209±162 mg/dL, and c-peptide was 1.5± 0.96 ng/mL. Mean HbA1c was 8.8±3.1%. Ten patients had autoantibody positivity. Comorbidities included exudative retinopathy (n=2), asthma (n=1), BHE (n=1), celiac disease (n=1), pelviectasis (n=1), hearing loss (n=2), and sensory neuropathy (n=1). Thirteen of the patients were on diet therapy, 15 were on oral anti-diabetic therapy and nine were on insulin therapy.

HNF4A MODY

This MODY subtype constituted 3.5% (n:8) of the cases. The mean age at diagnosis was 10.3±4.6years. There was one case with a birth weight of over 4000 g. Five cases had incidental hyperglycemia, and six cases had typical symptoms of diabetes. Acidosis was present in two cases, and autoantibody positivity was observed in one case. At the time of diagnosis, mean blood glucose was 306 ± 216 mg/dL, and c-peptide was 1.9 ± 1.2 ng/mL. while HbA1c range was 4.7-11.7%. One patient was observed to a diagnosis. The only comorbidity was spinocerebellar syndrome (n=3). While four of the patients were being managed with the diet, two were being treated with oral anti-diabetic therapyand two were being treated with insulin.

HNF1B-MODY

This mutation was detected in 3.1% (n=7) of the cases. The mean age at diagnosis was 12.4±3 years. Five patients had incidental hyperglycemia, and five patients had symptoms of diabetes. Two patients presented with diabetic ketoacidosis. Six patients had a family history. At the time of diagnosis, blood glucose was 308±177 mg/dL, and c-peptide was 1.18±1.0 ng/mL. HbA1c range was 5.4-14.5%. No autoantibody positivity was found in any of the patients. Additional anomalies were chronic renal failure and elevated transaminases in one, FSGS (Focal segmental glomerulosclerosis) in polycystic kidney in two, and renal cyst in one. Four of the patients were receiving insulin, one was on diet only and one was receiving oral anti-diabetic therapy and basal insulin.

KLF-MODY

KLF-MODY was present in 3.5% (n=8) of the patients. The mean age at diagnosis was 11.2±2.0 years. Three patients presented with incidental hyperglycemia, seven had symptoms of diabetes and three patients with diabetic ketosis. Family history was present in all cases. Atdiagnosis, blood glucose was 278±175 mg/dL, and c-peptide was 2.2±2.0 ng/mL. The range of HbA1c was 5.6-10.2%. Only one patient had autoantibodies, anti-GAD. One patient was on a diet, one on oral anti-diabetic therapy, two on oral anti-diabetic therapy and basal insulin, and four on intensive insulin treatment.

NEUROD1 MODY

NEUROD1 MODY was found in 1.3% (n=3) of the cases. Their age was 1, 12 and 15 years old. and one patient presented with DKA. No autoantibody positivity was detected. Additional anomalies included epilepsy in one patient and MMR (mental motor retarration) in one. At diagnosis, blood glucose was 164±100 mg/dL and c-peptide was 0.8±0.7 mg/dL. HbA1c was in the rang of 5.5 %. Family history was not found in any patient. Two patients were on a diet, and one received intensive insulin treatment.

INS-MODY

This MODY subtype was found in 1.3% (n=3) cases. Their age was 7.5, 7.8 and 11 years old. Two patients had no symptoms of diabetes, and one patient presented with DKA. All three patients had a familyhistory. Acidosis developed in one patient during follow-up. No autoantibody positivity was detected. At the time of diagnosis, blood glucose was 200 ± 64 mg tL, and c-peptide was 8.5 ± 4.3 mg/dL. HbA1c ranged widely from 6.2 to 14%. MMR was associated with one patient. One patient was on a diet, and two were receiving intensive insulin.

ABCC8-MODY

This subtype of MODY was found in 1.3% (n=3) of the cases. Their age was 8.5.11 and 12 years and diabetes symptoms were absent in two. Family history was present in all patients, and one patient was positive for autoantroodies. At diagnosis, blood glucose was 140 ± 26 mg/dL, and c-peptide was 2.4 ± 0.58 mg/dL. Two of the patients were taking oral ant-diabetic therapy and the other was managed with diet. PDX-MODY

A 3.9-year-old patient presented with DKA. Family history was positive. No autoantioodies were detected. Admission HbA1c was 5.2%. Blood glucose was 135 mg/dL, and c-peptide was 3.4 mg/dL at the time of diagnosis. He was being managed with diet only. **CEL-MODY**

CEL variant associated MODY was detected in one patient, 6 years old who presented with incidental hyperglycemia. There was no family history. Insulin was not started during the follow-up. Autoantibodic were negative and HbA1c was 5.6%. At diagnosis, blood glucose was 137 mg/dL, and c-peptide was 0.7 mg/dL. He v as managed with diet only.

KJNC11-MODY

This form of MODY was detected in one case 8.5 years old. Symptoms of diabetes and autoantibody positivity were present at presentation. There was a family history of DM. HbA1c was 7.8% and insulin was not started during follow-up. At the time of diagnosis, blood glucose was 191 mg/dL, and c-peptide was 2.1 mg/dL. The patients were followed up with oral anti-diabetic therapy. Of the 224 patients evaluated, eight patients present ed with acidosis. These cases were limited to the MODY types 1, 3, 5, and 6 and their detailed characteristics are shown in Table 3. DKA developed in one patient with *INS* mutation during follow-up. Table 5 shows the characteristics of the patients with positive theory heading.

characteristics of the patients with positive autoantibodies.

DISCUSSION

In this study, which is the largest generically diagnosed series from Turkey, the most common form of MODY was GCK-MODY(65%) followed by HNF1A-MODY (1990), while he other types were much rarer. Although the majority of cases were diagnosed because of

nonved by HNFTA-MODT (196), while do the types were much farer. Atthough the majority of cases were diagnosed because of incidental hyperglycemia, the e were a so cases with DKA and ketosis without a family history. Obesity was rarely associated with MODY, while comorbidities were seer in some MODY types. MODY is reported with different frequencies depending on the ethnicity, selected patient population, number of genes analyzed and technical characteristics. The distribution of subgroups may also differ according to the method, environment, and technique. In the present study, which presents a fair representation of centers from across Turkey, MODY2 (GCK-MODY) was the most common diagnosis, as reported pretored by (6,7). While *GCK* mutations were dual area in a sequence diagnosis, as expected, diabetes symptoms were observed in activity with *DUE(1)* and *DUE(1)*. patients with UNF14 and HNF4A variants. Treatment was changed in 37 (16%) of our patients after genetic diagnosis or during follow-up (5-9).

In the most largest series from Germany/Austria, MODY1 was found in 44, MODY2 in 609, MODY3 in 230, and MODY5 in 35 of 1047 MODY cases from 76,836 children under 20 years of age with diabetes (10). In the UK, MODY prevalence and subtype studies have been per ormed extensively and it has been reported that HNF1A-MODY is the most common type (52%) with GCK-MODY the second most common. While MODY due to *HNF1A* mutation is most common in European countries (14), GCK-MODY is the most common type in Japan, although there are few studies from Asian and Far Easterncountries (11).

Studies from Italy have shown that the selected patient population is important in determining the frequency of the MODY subtype. In the MODY subgroup in which patients withsymptomatic hyperglycemia were evaluated, HNF1A- and HNF1B-MODY were most prevalent, while in the group with incidental hyperglycemia, GCK-MODY was most common. This is also a factor in the difference in frequency of MODY subtypes in adult and pediatric studies. For an accurate ranking, both age groups should be evaluated together (12). Family history has been used as an essential criterion in MODY probability calculation models. This probability calculation reduces the probability of MODY in the absence of family history, but case reports and the present series show that there may be some exceptional cases. In a large series reported from Japan, 18% of the cases with genetically proven MODY did not have a family history (11). Furthermore, novel mutation was demonstrated in 7.3% of patients with the four most common MODY subtypes reported from Slovakia (13). In our series, the de novo mutation rate was 11%.

The use of anti-GAD positivity as a criterion for exclusion of MODY is a matter of debate. In a large series, anti-GAD was found to be positive in only 5 of 508 MODY cases, and this rate was similar to the normal population rate. However, in a Czech study, antiGAD positivity was found in 7 of 28 MODY cases, and IAC2 positivity was found in one of 28 MODY cases. In these patients, the course of diabetes was worse, and HbA1c values were higher. Autoantibody negativity has been reported with appropriate therapeutic control of diabetes. Schober et al. found a positive autoantibody rate of 17% in a German-Austrian series (14-16). In the present study, autoantibodies were positive in 25 patients. This may be due to a condition termed "double diabetes". Polygenic diabetes has been reported to affect 3-4% of MODY cases, and the coexistence of type 2 DM and MODY is more common than type 1 DM and MODY (17). In the present series, only three patients with GCK mutation had increased HbA1c values and insulin requirement in the follow-up. In

a Polish study, increased HbA1c values were also reported in patients with GCK mutation. This was explained by T1DM or T2DM accompanying monogenic diabetes. Monogenic DM does not preclude the existence of other types. In 8 of 285 cases with GCK mutation, HbA1c values above 7.5% were reported. Steel et al. also found HbA1c values between 8.2-9.5% in 6 of 235 cases with GCK mutation (18).

A CORPECTED PROC

Although obesity is used to differentiate cases with T2DM from MODY, obesity associated with rare types of MODY, such as those caused by variants in *IPF1/PDX1* (MODY4), *NEUROD1* (MODY6), *BLK* (MODY 11) and *ABCC8* (MODY12), and even common types (HNF1A- and HNF4A-MODY) have been reported in recent years. Therefore, obesity in MODY may be more common than previously thought, probably due to the general increase in global obesity prevalence (11). In the present series, obesity was found in 16 (%7) cases at diagnosis. In a Japanese study, the rate of overweight was 8.2%, and insulin resistance was 22% (18).

While MODY due to variants in *HNF4A*, *HNF1A* and *HNF1B* may present with DKA, *INS*, *NEUROD1* and *PDX*, which are rare types, may also present with DKA (19). Consistent with the literature, our patients also had these mutations and presented with acidosis due to insulin deficiency. It seems that using acidosis as anexclusion criterion for MODY may exclude some cases

The criteria of three generations affected with diabetes, autoantibodies being negative, and insulin reserve present do not always hold true. In the present series, autoantibodies were positive in 25 patients, seven patients presented with DKA, and no family history was found in 24 patients. Approximately 50% of positive MODY cases do not fulfill the classical MODY diagnostic criteria; some patients with genetically confirmed MODY may also have pancreatic antibodies, and some *de novo* mutations may have occurred (18).

Over time and with more extensive research, knowledge of the natural history of MODY has increased. The abilityto partially or completely change the treatment of people diagnosed with MODY from insulinuse to oral hypoglycaemic agents has saved patients from injections and unnecessary insulin therapy. However, while *GCK* mutations are followed up only with diet, *PDX*, *INS*, *BLK* and *APPL1* variant-associated MODY cases require earlyinsulin, and HNF4A, HNF1A, and PAX4, respectively, MODY due to *ABCCs* and *KCNJ11* mutations respond well to oral antidiabetics. In *HNF1B*, *NEUROD1*, *KLF11* and *CEL* mutations, the response to oral anticiabetics is variable and insulin requirement develops over time. In our series, treatment changes were made in 37 patients during follow-up. While 71% of the patients were followed up only by adjusting their diet, 10% were receiving intensive insulin trea ment, and 75 were receiving sulfonylurea.

Strengths of the present study include the high number of cases compared to early studies from Turkey, the detailed information about clinical characteristics of the identified cases and that this was a nationally supported study. However, the limitations include the fact that not all centers participated, the MODY panels used varied from center to center and the interpretation of the MODY reports were performed by several geneticists.

This large-series national study showed that the diagnostic criteria in the prediagnostic processof MODY should be reconsidered in the presence of DKA, the presence of antibody positivity and the absence of family history in patients with genetically diagnosed MODY. Therefore, it does not seems appropriate to use obesity and/or the absence of DKA as the absolute criterion for suggesting that a patient does not have MODY, and this will need to be considered in the development of unlated MODY screening guidelines. **AUTHOR CONTRIBUTIONS**

Elif Özsu and Zehra Aycan designed the study, Elif Özsu, Semih Bolu, Nihal Hatipoğlu, Şenay Savaş Erdeve, Olcay Evliyaoğlu, Firdevs Baş, Atilla Çayır, İsmail Dündar, Emine Demet Akbaş, Ahmet Uçaktürk, Şervan Özalkak, Nurse Muratoğlu Şahin, Melikşah Keskin, Ülkü Gül Şiraz, Hande Turan, Ayşe Pınar Öztürk, Eda Uçaktürk, Elif Sağsak, Fatma Dursur, Nesibe, kyürek, Sevinç Odabaşı Guneş, collected the data. Elif Özsu, Semra Çetinkaya and Zehra Aycan wrote the first draft of the paper, and Merih Berberoğlu, Zeynep şıklar and Zehra Aycan critically revised the paper. Elif Özsu is the guar- antor of the study and made the final decision to submit and publish the manuscript. ACKNOWLEDGEMENTS

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CONFLICT OF INTEREST

No conflict to interest

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Table 1 : Demographic characteristics of pediatric genetically confirmed MODY cases in a Turkish cohort (n=224)

Table I : Demo	HNF4A	GCK	HNF1A	PDX1	HNF1B	NEUROD1	KLF-	CELL	INS	ABCC8	KJNC11	Mean
	n=8	n=146	n=43	n=1	n=7	n=3	11 n=8	n=1	n=3	n=3	n=1	
Age at diagnosis (years)	10.3± 4.6	8.8 ±3.9	10.6±4.3	3.9	12.4± 3	9± 7.8	11.1 ±2	6	9.1 ±2.3	10.5 ±1.8	8.5	9.4± 4.1 (range 0.15-17)
Sex F/M	3/5	63/83	22/21	0/1	3/4	1/2	4/4	1/0	1/2	2/1	1	101/123(45%/55%)
Symptom (+) (-)	6 2	23 123	5 18	1 0	5 2	0 3	7 1	0 1	0 3	1 2	1 0	30% (49/224)
Presentation (n) IH DK DKA	5 1 2	145 0 0	31 9 2	0 0 1	5 1 2	2 0 1	3 3 0	1 0 0	3 1 0		0	197 (87%) 16 (7%) 8 (3.5%)
Negative Family History (n)	2	17	4	0	1	0	0	0	0	0	0	24/224(11%)

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IH: incidental hyperglycemia; DK,:diabetic ketosis; DKA: diabetic ketoacidosis

Table II a:Treatment PDX1 HNF1B KLF-11 CELL INS ABCC8 KJNC11 **JEU** n 1 n 1 n 1 n 7 3 n 8 n 3 n 3 Diet OAD İnsülin Literatüre D insulin D OAD OAD D OAD OAD OAD Insülin Insülin OAD (metformin, DPP4) İnsülin İnsülin Insulin Treatment D:1 D:1 D:2 D:1 D:1 OAD:1 OAD:2 OAD:1 intensive ins:4 intensive ins:1 M:1 İntensive Basal ins.+OAD:1 intensive ins:4 ins:1 Basal ins+OAD:2

OAD:Oral Anti-Diabetic M:metformin DPP4: dipeptidyl peptidase-4 inhibitors D:Diet SU: Sulphonylureas GLP1A: Glucagon-Like Peptide 1 Receptor Agonist SGLT-2: Sodium-Glucose Transport Protein 2 Inhibitors

Table II b: Treatment

Treatment	HNF4A (n=8)	GCK (n=146)	HNF1A (n=43)
Lecture	Diet SU GLP1-RA insulin	Diet	SU Meglitinide, GLP-1 RA, SGLT-2 insulin
In our cases	D:4 Su:2 intensive insulin:2	D:135 SU:3 OAD (M):3 insulin:2 intensive ins: 3	D:13 OAD (S):11 OAD (M):4 ins:7 Basal ms+OAD(2
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Table III: Characteristics of patients presenting with acidosis or developing acidosis during follow- up

Family HistoryNoNoYes grandfatherYes (brother, grandfather)Yes (father)Yes (father)Yes (father)Yes (father)Yes (father)Yes (father)Yes (father)Yes (father)Yes FatherGlucose at Dx (mg/dL)674548443210600281269C peptide (ng/mL)0,600,530,10,10,10,13,47HbA1c (%)11,7810,613,2987,511,2Treatment treatment continuedInt insulin the same treatment continuedInt insulin the same treatment continuedInt insulin the same treatment continuedInt insulin the same treatment continuedInt insulin the same treatment continuedInt insulin the same treatment continued
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Table IV : Characteristics of Patients on Insulin

Number of patients2Features in the applicationK: 1 DKA:1 DM sempt at dx: 2HbA1c %11,7 12,9	Diabet symptom:2 Diabet s 5.8, 6, 6, 7, 6.4, 7.4, 7.2, 8, 8.4,8	symptom: Diabet sympto 6, 7.5,	Diabet	K:3 DKA:0 Diabet symptom:6	1 1 K:0 K:0 DKA:0 DKA Diabet Didt symptom:0 symptom	
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HbA1c % 11,7 12,9	6, 7, 6.4, 7.4, 7.2, 8, 8.4,8	7.5,	7.4			
	7.4 9,11.30 12.5,13	7.7, 8, 9.5 .6,14.6,20 7.7, 8, 9.5 17.5		6.8, 9.8, 10, 11.9 14.7, 15.6	11.2 5.8	
				S S		
K:Ketosis DKA:Dia	is abetic Ketoasidosis		\sim			

	HNF4A n 8	GCK n 146	HNF1A n 43	PDX1 n 1	HNF1B n 7	NEUROD1 n 3	KLF-11 n 8	CELL n 1	INS n 3	ABCC8 n 3	KJNC11 n 1	Totall
Anti- GAD Anti- insulin Anti- islet	Anti- GAD:1 Anti- insulin:0 Anti-islet :0	Anti- GAD:0 Anti- insulin:1 Anti- islet:5	Anti- GAD:4 Anti- insulin:3 Anti-islet :3	Anti- GAD:0 Anti- insulin:0 Anti- islet:0	Anti- GAD:0 Anti- insulin:0 Anti- islet:0	Anti-GAD:0 Anti- insulin:0 Anti-islet:0	Anti- GAD:1 Anti- insulin:0 Anti- islet:0	Anti- GAD:0 Anti- insulin:0 Anti- islet:0	Anti- GAD:0 Anti- insulin:0 Anti- islet:0	Anti- GAD:1 Anti- insulin:0 Anti- islet:0	Anti- GAD:0 Anti- insulin:0 Anti- islet:0	13 4 8
	1	12	10	0	0	0	1	0	0	1	0	25

