

# The University of Chicago Genetic Services Laboratories



5841 S. Maryland Ave., Rm. G701, MC 0077, Chicago, Illinois 60637  
Toll Free: (888) UC GENES ☐ (888) 824 3637  
Local: (773) 834 0555 ☐ FAX: (773) 702 9130  
ucgslabs@genetics.uchicago.edu ☐ dnatesting.uchicago.edu  
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## GCK Analysis for Maturity Onset Diabetes of the Young

### Clinical Features

Heterozygous inactivating mutations in *GCK* [OMIM#138079] have been described in patients with maturity onset diabetes of the young type 2 (GCK-MODY) [OMIM#125851], which is characterized by mild fasting hyperglycemia (1). Hyperglycemia is present at birth but often only detected later in life, when individuals undergo routine screening tests (1). Affected individuals rarely, if ever, show progression of disease, or develop the microvascular or macrovascular complications typically associated with diabetes (1). These patients typically therefore can be managed by diet alone, and treatment with oral medications or insulin can actually cause poorer outcomes as patients have an altered counter-regulatory response to hypoglycemia (2). Homozygous inactivating *GCK* mutations are associated with permanent neonatal diabetes mellitus (PNDM) (1). In addition, heterozygous activating mutations in *GCK* have also been observed, which lead to hypoglycemia (1).

### Molecular Genetics

*GCK* encodes for the enzyme glucokinase, which has a central role in the regulation of blood glucose and acts as a "glucose sensor" in pancreatic  $\beta$ -cells (3). Mutations in *GCK* associated with GCK-MODY typically result in a modest decrease in glucokinase activity, which in turn leads to mild fasting hyperglycemia (4).

### Inheritance

GCK-MODY is inherited in an autosomal dominant manner. The majority of mutations are inherited, although *de novo* mutations have also been described. Recurrence risk for children of an affected individual is 50%.

### Test methods

Comprehensive sequence coverage of the coding regions and splice junctions of the *GCK* gene is performed. Targets of interests are captured and amplified using Agilent HaloPlex target enrichment system. The constructed genomic DNA library is sequenced using Illumina technology and reads are aligned to the reference sequence. Variants are identified and evaluated using a custom collection of bioinformatic tools and comprehensively interpreted by our team of directors and genetic counselors. All novel and/or potentially pathogenic variants are confirmed by Sanger sequencing. The technical sensitivity of this test is estimated to be >99% for single nucleotide changes and insertions and deletions of less than 20bp. Deletion/duplication analysis of the *GCK* gene by oligonucleotide array-CGH identifies copy number changes involving one or more exons. Partial exonic copy number changes and rearrangements of less than 400 bp may not be detected by this methodology. Array-CGH will not detect low level mosaicism, balanced translocations, inversions, or point mutations that may be responsible for the clinical phenotype. The sensitivity of this assay may be reduced when DNA is extracted by an outside laboratory.

#### GCK sequencing

Sample specifications: 3 to 10 cc of blood in a purple top (EDTA) tube  
Cost: \$2,200  
CPT codes: 81405  
Turn-around time: 4 - 6 weeks

#### GCK deletion/duplication analysis

Sample specifications: 3 to 10 cc of blood in a purple top (EDTA) tube  
Cost: \$1000  
CPT codes: 81404  
Turn-around time: 4 weeks

**Note: The sensitivity of our assay may be reduced when DNA is extracted by an outside laboratory.**

#### Testing for a known mutation in additional family members by sequence analysis

Sample specifications: 3 to 10 cc of blood in a purple top (EDTA) tube  
Cost: \$390  
CPT codes: 81403  
Turn-around time: 3-4 weeks

#### Prenatal testing for a known mutation by sequence analysis

Sample specifications: 2 T25 flasks of cultured cells from amniocentesis or CVS  
or 10 mL of amniotic fluid  
Cost: \$540  
CPT codes: 81403  
Turn-around time: 1-2 weeks

### Results

Results, along with an interpretive report, will be faxed to the referring physician. Additional reports will be provided as requested. All abnormal results will be reported by telephone.

## References

1. Osbak KK, Colclough K, Saint-Martin C et al. Update on mutations in glucokinase (GCK), which cause maturity-onset diabetes of the young, permanent neonatal diabetes, and hyperinsulinemic hypoglycemia. *Hum Mutat* 2009; 30: 1512-1526.
2. Guenat E, Seematter G, Philippe J et al. Counterregulatory responses to hypoglycemia in patients with glucokinase gene mutations. *Diabetes Metab* 2000; 26: 377-384.
3. Negahdar M, Aukrust I, Johansson BB et al. GCK-MODY diabetes associated with protein misfolding, cellular self-association and degradation. *Biochim Biophys Acta* 2012; 1822: 1705-1715.
4. Froguel P, Zouali H, Vionnet N et al. Familial hyperglycemia due to mutations in glucokinase. Definition of a subtype of diabetes mellitus. *N Engl J Med* 1993; 328: 697-702.

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