

# Inherited Insulin Resistance Syndromes (*INSR*) Genetic Testing

## Indications for Ordering

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- Confirm diagnosis of inherited insulin resistance syndromes in individuals with clinical and/or biochemical evidence
- Not intended for evaluation of individuals with nonsyndromic forms of insulin resistance, such as isolated diabetes mellitus with no other physical features

## Test Description

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Polymerase chain reaction, followed by bidirectional sequencing of the entire coding region and intron/exon boundaries of the *INSR* gene

## Tests to Consider

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### Primary test

[Inherited Insulin Resistance Syndromes \(\*INSR\*\) Sequencing 2006274](#)

### Related tests

Initial biochemical tests for suspicion of inherited insulin resistance syndrome

- [Insulin, Fasting 0070063](#)
- [Fasting Glucose, Plasma or Serum 0020024](#)

## Disease Overview

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**Incidence** – unknown, but estimated to be rare

## Clinical presentation

Variants in the *INSR* gene cause three main syndromes

- Donohue syndrome (most severe)
  - Intrauterine growth restriction
  - Failure to thrive
  - Loss of glucose homeostasis
  - Hyperinsulinemia
  - Enlarged heart and kidneys
  - Dysmorphic features
    - Prominent eyes
    - Thick lips
    - Upturned nostrils
    - Low-set, posteriorly rotated ears
  - Thick skin with lack of subcutaneous fat
  - Distended abdomen
  - Enlarged genitalia
  - Polycystic ovaries in females
  - Death often occurs prior to age 1
- Rabson-Mendenhall syndrome (intermediate phenotype)
  - Growth retardation
  - Hyperinsulinemia
  - Acanthosis nigricans
  - Diabetes mellitus
  - Dysmorphic features
    - Premature or dysplastic teeth
    - Gingival hyperplasia
    - Pineal hyperplasia
  - Survival ranges from early childhood to adolescence
- Type A insulin resistance syndrome (least severe)
  - Hirsutism
  - Reduced subcutaneous fat
  - Diabetes mellitus
  - Acanthosis nigricans
  - Hyperinsulinemia
  - Amenorrhea and polycystic ovaries in females
  - Survival often beyond middle age

## Pathophysiology

Insulin resistance occurs when insulin receptors are unable to bind insulin

- Results in decreased insulin action on target organs
  - Decreased insulin is compensated for by the pancreas with increased insulin release
    - Pancreatic beta cells become unable to compensate
      - Leads to increased glucose production by the liver and lipolysis of adipose tissue, resulting in ketoacidosis

## Diagnosis

Inherited insulin resistance syndrome is often diagnosed based on

- Clinical features
- Glucose and insulin levels
- Fibroblast studies for insulin binding
- Genetic testing

## Genetics

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**Gene** – *INSR*

### Inheritance

- Donohue syndrome – autosomal recessive
- Rabson-Mendenhall syndrome – autosomal recessive
- Type A insulin resistance syndrome – autosomal recessive or autosomal dominant
  - Recessive forms are more severe
  - Dominant forms may require contribution of other genetic or environmental factors to produce phenotype

**Penetrance** – unknown, but expected to be reduced for individuals with a dominant variant

## Test Interpretation

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### Sensitivity/specificity

- Clinical sensitivity – predicted to be >90% in individuals with a clinical diagnosis
- Analytical sensitivity/specificity – 99%

### Results

- Positive
  - One copy of pathogenic *INSR* gene variant detected
    - Predicts carrier status for an inherited insulin resistance syndrome
    - In some cases, one copy of a variant may indicate an increased likelihood for developing type A insulin resistance
      - Depends on other genetic and environmental factors
  - Two pathogenic *INSR* gene variants detected
    - Predicts a diagnosis of an inherited insulin resistance syndrome
- Negative
  - No pathogenic variants detected
    - Likelihood is reduced that the individual is a carrier of or is affected with an inherited insulin resistance syndrome
- Inconclusive
  - *INSR* gene variants of unknown clinical significance may be detected by this test

### Limitations

- Diagnostic errors can occur due to rare sequence variations
- Not detected
  - Regulatory region and deep intronic mutations
  - Large deletions and duplications
- Genes other than *INSR* will not be evaluated
- Medical management of patient should rely on clinical and/or biochemical findings