

# Inherited Insulin Resistance Syndromes (INSR) Genetic Testing

# **Indications for Ordering**

- 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**

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

# **Tests to Consider**

#### **Primary test**

Inherited Insulin Resistance Syndromes (INSR) Sequencing 2006274

#### **Related tests**

Initial biochemical tests for suspicion of inherited insulin resistance syndrome

- Insulin, Fasting 0070063
- Glucose, Plasma or Serum 0020024

## **Disease Overview**

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
  - o Hyperinsulinemia
- Enlarged heart and kidneys
- Dysmorphic features
  - Prominent eyes
  - Thick lips
  - Upturned nostrils
  - Low-set, posteriorly rotated ears
- $\circ$  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
- o Hyperinsulinemia
- Acanthosis nigricans
- o Diabetes mellitus
- o Dysmorphic features
  - Premature or dysplastic teeth
  - Gingival hyperplasia
  - Pineal hyperplasia
- o Survival ranges from early childhood to adolescence
- Type A insulin resistance syndrome (least severe)
  - o Hirsutism
  - $\circ$  Reduced subcutaneous fat
  - o Diabetes mellitus
  - $\circ \text{Acanthosis nigricans}$
  - $\circ$  Hyperinsulinemia
  - o 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

## Gene – INSR

## Inheritance

- Donohue syndrome autosomal recessive
- Rabson-Mendenhall syndrome autosomal recessive
- Type A insulin resistance syndrome autosomal recessive or autosomal dominant
  - $\circ\, \text{Recessive}$  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**

## 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
  - $\circ$  Two pathogenic *INSR* gene variants detected
    - Predicts a diagnosis of an inherited insulin resistance syndrome
- Negative
  - $\circ\, \text{No}$  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 variants
    Large deletions and duplications
- Genes other than INSR will not be evaluated
- Medical management of patient should rely on clinical and/or biochemical findings