

## Cytochrome P450 1A2 Genotype, Varies

**Test ID:** 1A2Q

**Useful for:**

Identifying individuals who are poor, intermediate, normal (extensive) or rapid metabolizers of drugs metabolized by cytochrome P450 1A2 to assist drug therapy decision making

**Method:**

Real-Time Polymerase Chain Reaction (PCR) with Allelic Discrimination Analysis

**Advisory Information:**

Testing is available as the single gene assay (this test) or as a part of a focused pharmacogenomics panel, which includes testing for the following genes: *CYPs 1A2, 2C9, 2C19, 2D6, 3A4, 3A5, 4F2, SLCO1B1, and VKORC1*.

Order PGXQP / Focused Pharmacogenomics Panel, Varies if multiple pharmacogenomic genotype testing is desired.

**Reference Values:**

An interpretive report will be provided.

**Specimen Requirements:**

Multiple genotype tests can be performed on a single specimen after a single extraction.

**Submit only 1 of the following specimens:**

**Specimen Type:** Whole blood

**Container/Tube:** Lavender top (EDTA)

**Specimen Volume:** 3 mL

**Collection Instructions:**

1. Invert several times to mix blood.
2. Send specimen in original tube.

**Specimen Stability Information:** Ambient (preferred) 9 days/Refrigerated 30 days

**Specimen Type:** Saliva

**Patient Preparation:** Patient should not eat, drink smoke, or chew gum 30 minutes prior to collection.

**Supplies:** Saliva Swab Collection Kit (T786)

**Specimen Volume:** 1 Swab

**Collection Instructions:** Collect and send specimen per kit instructions.

**Specimen Stability Information:** Ambient 30 days

**Specimen Type:** Extracted DNA

**Container/Tube:** 2 mL screw top tube

**Specimen Volume:** 100 mcL (microliters)

**Collection Instructions:**

1. The preferred volume is 100 mcL at a concentration of 50 ng/mcL.
  2. Include concentration and volume on tube.
- Specimen Stability Information:** Frozen (preferred)/Ambient/Refrigerated

**Specimen Stability Information:**

Specimen Type	Temperature	Time
Varies	Varies	

**Cautions:**

Rare variants may be present that could lead to false-negative or false-positive results. If results obtained do not match the clinical findings (phenotype), additional testing should be considered.

Samples may contain donor DNA if obtained from patients who received non-leukoreduced blood transfusions or allogeneic hematopoietic stem cell transplantation. Results from samples obtained under these circumstances may not accurately reflect the recipient's genotype. For individuals who have received blood transfusions, the genotype usually reverts to that of the recipient within 6 weeks. For individuals who have received allogeneic hematopoietic stem cell transplantation, a pretransplant DNA specimen is recommended for testing.

*CYP1A2* genetic test results in patients who have undergone liver transplantation may not accurately reflect the patient's *CYP1A2* status.

This method may not detect all variants that result in altered *CYP1A2* activity. Therefore, absence of a detectable variant does not rule out the possibility that a patient has altered *CYP1A2* metabolism due to other *CYP1A2* variants that cannot be detected with this method. Furthermore, when 2 or more variants are identified, the cis-/trans- status (whether the variants are on the same or opposite chromosomes) is not always known. It should be noted that other laboratories may use different phenotype prediction methods as there is no consensus on this at this time. However, the method used here represents the findings of the majority of literature available at this time.

The frequency of variants which cause altered *CYP1A2* metabolism has not been fully characterized in all ethnic groups. *CYP1A2* enzyme activity may be inhibited or induced by a variety of substances, medications, or their metabolites.

**CPT Code:**

0031U

**Day(s) Setup:** Monday through Friday

**Analytic Time:** 3 days; not reported on Saturday or Sunday

**Questions**

Contact Heather Flynn Gilmer, Laboratory Technologist Resource Coordinator at 800-533-1710.