Non-Alcoholic Fatty Liver Disease Susceptibility (PNPLA3) Genotyping
ARUP test code 2014599

PNPLA3 Specimen: Whole Blood

PNPLA3 Variant: Homozygous *

PNPLA3 Interpretation:

See Note

Indication for testing: screening for genetic susceptibility to non-alcoholic fatty liver disease.

Positive: Two copies of the PNPLA3 c.444C>G; p.I148M variant were detected. This individual has a moderately increased genetic risk for the development and progression of non-alcoholic fatty liver disease. This genotype is also associated with a moderately increased risk for cirrhosis among individuals with alcoholic liver disease. Other genetic or environmental risk factors not detected by this assay may be present.

This result has been reviewed and approved by Pinar Bayrak-Toydemir, M.D., Ph.D.
BACKGROUND INFORMATION: Non-Alcoholic Fatty Liver Disease Susceptibility (PNPLA3) Genotyping

CHARACTERISTICS: Fatty liver disease is the accumulation of excessive triglycerides in the liver that may cause an inflammatory response, which can progress to fibrosis, cirrhosis, and liver cancer. The c.444C>G; p.I148M variant in the PNPLA3 gene confers an increased risk for the onset and progression of non-alcoholic fatty liver disease (NAFLD). This allele also confers an increased risk for the onset and progression of cirrhosis among individuals with alcoholic liver disease.

INCIDENCE: NAFLD occurs in approximately 20–30 percent of individuals in the US.

G ALLELE FREQUENCY: Varies by ethnicity; Latino 0.57, East Asian 0.38, European 0.23, South Asian 0.22, Africans 0.14.

CAUSE: Risk factors for non-alcoholic fatty liver disease include obesity, diabetes, insulin resistance and genetic risk factors including PNPLA3 c.444C>G; p.I148M.

INHERITANCE: Multifactorial.

CLINICAL SENSITIVITY: Unknown.

VARIANT TESTED: PNPLA3 c.444C>G; p.I148M (rs738409).

METHODOLOGY: Polymerase chain reaction followed by high-resolution melt analysis.

ANALYTICAL SENSITIVITY AND SPECIFICITY: Greater than 99 percent.

LIMITATIONS: Only the c.444C>G; p.I148M variant in the PNPLA3 gene will be targeted. Diagnostic errors can occur due to rare sequence variations.

Test developed and characteristics determined by ARUP Laboratories. See Compliance Statement C: aruplab.com/CS

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