



Reference No. : - 2411220318

Pt's Name : Mr. NAVPREET SINGH

Referred By : NA

Sample Collection Date/Time : 14-Nov-2024

Sample Receiving Date/Time : 14-Nov-2024 05:38AM

: KOS DIAG LAB Sample From

Age/Gender : 37 Yrs/Male

AMB-KOS

:15-Nov-2024 06:37PM

Date :14-Nov-2024

Approved Date

Report Print Time :17-Nov-2024 07:02PM

## **Molecular Biology**

Test Description	Observed Value	Biological Reference Interval
	MTHFR Gene PCR*	

## **MTHFR Mutation Detection**

C677T\* Homozygous mutant

A1298C\* Wild

Method: Real Time Polymerase chain Reaction (PCR)

MTHFR (Methylene Tetrahydrofolate Reductase) is an enzyme involved in the anabolism of methionine, where it converts 5,10 methylenetetrahydrofolate to 5methyltetrahydrofolate. This molecule, with homocysteine will form methionine in downstream processes. The gene for MTHFR expression is located on chromosome 1 of the human genome. Polymorphisms at 677 (C>T) and 1298 (A>C) reduce enzymatic activity. Reduction of MTHFR enzymatic activity results in increase of homocysteine levels and may produce hyperhomocysteinemia. Hyperhomocysteinemia is an independent risk factor for causing various blood vessel diseases including brain and/or heart blood vessel diseases and peripheral venous thrombosis. Because of the risks associated with reduction in MTHFR enzymatic activity, a method of determining the genetic staus of the MTHFR gene is necessary.

## **Interpretation:**

Limit of detection: 10 ng/ul

"Wild": No mutation in MTHFR gene

"Heterozygous mutant": Partial mutancy in MTHFR gene "Homozygous mutant": 100% mutancy in MTHFR gene

## Methodology details:

- \* DNA is extracted from samples by US FDA approved Automatic Extraction machine based on magnetic bead technology.
- \* Purified DNA is then Amplified and quantified using CE-IVD approved Real time PCR.
- \* Extraction and Amplification controls (IC) are incorporated in each run to ensure more accurate and precise detection of mutation.

\*\*\* End Of Report \*\*\*

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