

# BRAF V600 Mutation Detection Kit

Qualitative Detection of BRAF V600 mutations

50 Reactions

20°C

INTENDED USE

BRAF V600 Mutation Detection Kit is an *in-vitro* multiplex, Real-Time PCR assay for the qualitative detection of V600 Mutations and differentiation of V600E mutation in the BRAF gene in human genomic DNA extracted from fresh, frozen, or formalin-fixed paraffin-embedded (FFPE) tumor tissue. The Kit differentiates V600E mutation from the other V600 mutations like K/D/Ecomplex/M/R/G.

SAMPLE TYPE

Human genomic DNA extracted from fresh, frozen, or formalin-fixed paraffin-embedded (FFPE) tumor tissue

BACKGROUND

High-frequency BRAF mutations have been detected in specific cancers, with melanoma showing approximately 50 to 60% of cases. The presence of BRAF gene mutations is not limited to a particular body part or cell type; it is associated with various cancers, such as Melanoma, Hairy Cell Leukemia, Non-Hodgkin Lymphoma, Thyroid Cancer, Ovarian Cancer, Lung adenocarcinoma, Colorectal cancer, and certain brain cancers like glioblastoma. Around 90% of identified BRAF mutations in human cancer involve a T1799A transversion mutation in exon 15, leading to a V600E amino acid substitution. This V600E mutation accounts for 70 to 90% of BRAF mutant melanoma cases. Additionally, the T1799A alteration might be linked to a second nucleotide mutation (G1798A), resulting in a V600K mutation in an additional ~6% to 29% of patients with a BRAF mutation. The serine/threonine protein kinase BRAF plays a crucial role in the epidermal growth factor receptor (EGFR)-mediated mitogen-activated protein kinase (MAPK) pathway, where activation occurs through the RAS small GTPase.

PRODUCT DESCRIPTION

BRAF V600 Mutation Detection Kit is an *in-vitro* multiplex Real-Time PCR qualitative assay for the detection of V600 mutations and differentiation of V600E mutations in the BRAF gene in human genomic DNA extracted from fresh, frozen, or formalin-fixed paraffin-embedded (FFPE) tumor tissue.

KIT COMPONENTS		VOLUME
Master Mix		625 µL
Primer and Probe mix		65 µL
Nuclease Free Water		500 µL
Positive Test Control		50 µL
Instruction for use		1 No.

MATERIALS REQUIRED BUT NOT PROVIDED

Consumables

PCR Plates/tubes  
PCR plate covers/tube caps

COMPATIBLE INSTRUMENTS

Real-Time PCR, Instruments like, BIORAD-CFX96, THERMO-Q55, QIAGEN-ROTOR-GENE Q and other instruments which supports FAM (495 nm - 520 nm), Cy5 (649 nm - 670 nm) and HEX (535 nm - 556 nm).

TARGET	REPORTER	QUENCHER
V600 K/D/E complex/M/R/G	FAM	BHQ1
V600 E	Cy5	BHQ2
BRAF Wild	HEX	BHQ2

\*Select the quencher settings as BHQ/None

STORAGE AND HANDLING

- Store all BRAF Qualitative Kit reagents at -20°C.
- Do not repeatedly freeze-thaw reagents more than five times as it leads to reduced assay sensitivity. Thaw the reagents only on ice or at 4°C.
- Kit components are stable through the end of the expiration date indicated on the box when stored at -20°C. Shelf Life - 12 Months from date of manufacturing.

PRECAUTIONS

- It is recommended that this product is used by personnel specially instructed and trained in real-time PCR and *in-vitro* diagnostics procedures.
- Treat all the specimens as potentially infectious.
- Wear protective disposable powder-free gloves, a laboratory coat and eye protection when handling specimens.
- Store positive and/or potentially positive material separated from all other components of the kit.
- Keep separate areas for master mix and template preparation and work under biosafety cabinets.
- Use aerosol barrier pipette tips and frequently change the gloves.
- Do not open the reaction tubes/plates post-amplification, to avoid contamination with amplicons.
- Do not smoke, drink or eat in areas where kit reagents and/or human specimens are being used.
- Do not use kit components that have passed their expiration date.

REACTION MIXTURE - 25 µL

Reagents	1 Rxn	20 Rxn	50 Rxn
Master Mix	12.5 µL	250 µL	625 µL
Primer & Probe mix	1.25 µL	25 µL	62.5 µL
Nuclease Free Water	6.25 µL	125 µL	312.5 µL
Total	20 µL	400 µL	1000 µL

Add 5 µL of the test DNA per reaction

SAMPLE PREPARATION

DNA should be extracted from fresh, frozen, or formalin-fixed paraffin-embedded (FFPE) tumor tissue aspirates using any approved DNA extraction methods. Store the extracted DNA at -20 °C for further use.

TEST PREPARATION/REACTION SET-UP

- Thaw all components of the kit on ice, mix gently using vortex and spin down the contents for 5 secs and use it immediately.

- Prepare the reaction mix in a 1.5/2 mL tube for the calculated number of samples in Master Mix Preparation room.
- Spin down the tubes and dispense 20 µL reaction mix in each tube strips or 96 well plate. Before moving to template adding area, add 5 µL of nuclease free water in NTC wells.
- Carefully add 5 µL of DNA samples kept on ice in the designated wells in template addition room. Add 2 µL of PTC in a separate hood and make up the volume by adding 3 µL of nuclease free water. The assay should be run along with positive controls and negative controls.
- Seal the plate carefully, briefly spin down and use any qRT-PCR instrument which complies with the dyes specified in the kit insert.

THERMAL CYCLIC CONDITIONS				
STEP	TEMP °C	TIME	DETECTION	CYCLE
Hold	95	1 Min	Off	1
PCR and Detection	95	5 Sec	Off	40
	50	20 Sec	On	

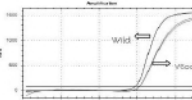
READING TEST RESULTS / DATA ANALYSIS

- NTCs should be negative and should not exhibit fluorescence amplified curves that cross the threshold line.
- If a false positive occurs with one or more of the primer and probe in NTC reactions, it indicates sample contamination.
- In that case, invalidate the run and repeat the assay with stricter adherence to the procedure guidelines.
- Positive control should produce a positive result with an expected Ct value for each target included in the test.
- If expected positive reactivity is not achieved, invalidate the run and repeat the assay with stricter adherence to procedure guidelines.
- After completion of the run, analyze the data as per the instrument manufacturer instructions.
- Analysis should be performed separately for each target using a manual threshold settings.
- In case internal control has not worked for a sample re-do the test with 2 or more dilutions.
- Negative results do not exclude possibility of infection and should not be used as the sole basis for the treatment.

RESULTS INTERPRETATION		
For V600 E & K/D/E complex/M/R/G	For BRAF Wild	Assay result
Ct < 40	Ct ≤ 40	Positive
Ct = Undetermined or Ct ≥ 40	Ct ≤ 40	Negative
Ct = Undetermined or Ct ≥ 40	Ct ≥ 40 Undetermined	Invalid. Re-purify the nucleic acid and repeat the test.

DATA INTERPRETATION			
V600E	V600 K/D/E complex/M/R/G	Wild	Results Interpretation
+	-	+	V600E detected
-	+	+	V600 K/D/E complex/M/R/G detected
-	-	+	No mutation detected
-	-	-	Invalid results repeat the nucleic acid extraction and re-run

PERFORMANCE CHARACTERISTICS



Performance characteristics has been established by testing the BRAF V600 Mutation Detection Kit with Horizon Discovery V600E standard .50 %

WASTE DISPOSAL

- Dispose all the waste/remains of the reagents used in reaction mixture preparation & expired kit components along with bio-waste as per the lab manual/general bio-waste management instruction.
- Dispose the PCR plates with patient samples "sealed" post run to avoid potential infection to the operators and contamination of the lab.

TROUBLESHOOTING

**Positive control showed no amplification**  
*Inappropriate storage of reagents*

- Store the reagents at recommended temperature for their optimal performance.
- Avoid repetitive freezing and thawing.
- Check the expiry of reagents.

**Negative controls are positive**  
*Causes - Cross-contamination*

- Follow good laboratory practices to avoid contamination issues.
- Use a new batch of reagents and repeat the experiment.

**Abnormal plot and/or low ΔRn values in amplification curve**  
*The baseline was set improperly (some samples have CT values lower than the baseline value)*

- Switch from manual to automatic baseline, or move the baseline stop value to a lower CT (2 cycles before the amplification curve for the sample crosses the threshold)

**An amplification signal is detected in the early cycles**  
*Dilute the sample to increase the CT value*