# **BRaCA** panel

Hereditary variants profiling in Breast and Ovary Cancer

# **4**bases

When Precision Matters

### **BRaCA PANEL**

4bases **BRaCA panel** is a **CE-IVD** kit for the identification of mutations in **BRCA1 and BRCA2 genes** (Table I) through a molecular protocol for Next Generation Sequencing (NGS) technologies on **Oxford Nanopore Technologies**' platforms. The kit is validated for germline and somatic analysis (SNVs, CNVs) of DNA extracted from blood samples and tissues (fresh, frozen or FFPE).

#### TECHNOLOGY

**BRaCA panel kit** contains all reagents needed for the amplification of BRCA1 and BRCA2 target genes and library multiplexing for NGS sequencing using **Oxford Nanopore Technologies**' platforms (Table 2).

#### WORKFLOW

The **BRaCA panel** kit is part of an amplicon-based family solution of 4bases molecular technologies. Its **accuracy and specificity** in detecting genetic variants (i.e. SNVs, CNVs), coupled with its **straightforward protocol** (<3 hours hands-on-time), are the key to its reliability. Thanks to **4bases proprietary analysis platform**, **4eVAR**, the clinical results are easily and efficiently accessible through a validated internal workflow.

#### SOFTWARE ANALYSIS

**4eVAR** is the analysis software developed by 4bases. The analysis is designed according to the characteristics and technologies of the kit used, to increase the **accuracy of the results** and to have **complete control over the entire process**. A **comprehensive report** is provided for each sample including SNV and CNV with ACMG and AMP classification for variant interpretation. Specific links to external databases are also provided for broader and more **complete classification** of **detected variants**, along with **quality metrics for each sample** in the run.

## VALIDATION

The **BRaCA** panel kit has been **validated** both internally and in clinical laboratories, and the whole protocol was tested on **DNA standards and on clinical samples.** 



Table 1: List of target genes in BRaCA panel.



Table 2: Samples per BRaCA panel sequencing run onOxford Nanopore Technologies' sequencers.

Flow Cell	Samples per Flow Cell (Coverage 100x)	Oxford Nanopore sequencer	Flow Cells per sequencer*
FLO-FLG114	8	Flongle	1
FLO-MIN114	96	MinION	1
		GridION	Up to 5
FLO- PRO114M	320	PromethION 2/P2 Solo	Up to 2
		PromethION 24	Up to 24
		PromethION 48	Up to 48

\*Multiple Flow Cells can be used during the same run session (GridION, PromethION), either with different libraries or with the same library to increase sequencing depth.

PLEASE NOTE: The volumes of the reagents in the kit are calculated to allow the subdivision into multiples of 8 samples per analysis session. Dividing the kit differently will decrease the total number of tests that can be performed. For different settings, please contact us at support@4bases.ch.

**PLEASE NOTE:** The number of multiplexable samples and coverage depend on starting gDNA degradation, presence of contaminants, external factors, library loading procedure.

#### **ORDERING INFORMATION \*\***

	Product	REF
LIBRARY AMPLIFICATION REAGENTS	BRaCA panel	H1070-16 (16 test)
	BRaCA panel	<b>H1070-96</b> (96 test)
MULTIPLEXING REAGENTS	BARCODE ONT Set 16	<b>R8001-16</b> (16 test)
	BARCODE ONT Set 96	<b>R8002-96</b> (96 test)
OXFORD NANOPORE ADAPTERS	NANO ADAPTER	<b>M1050-6</b> (6 sequencing runs)
	NANO ADAFTER	<b>M1050-24</b> (24 sequencing runs)

\*\*The reagents required for sequencing on Oxford Nanopore Technologies' platforms have to be purchased separately: contact 4bases for further information.

PLEASE NOTE: Should you need to run more than/multiple of 96 samples in the same library, please contact us.

For more detailed information regarding validated solutions for automatic library preparation, please contact us at info@4bases.ch

For the target regions of BRCA1 and BRCA2 genes (.bed file), please contact us at **support@4bases.ch** 

