

# HRD v2 Panel

The **oncoReveal® HRD v2 Panel** is designed for detection of SNVs, indels, and BRCA1 & BRCA2 exon level CNVs. 33 HRD-related genes can be assessed in either FFPE or blood samples. The panel uses proprietary Stem-Loop Inhibition-Mediated amplification (SLIMamp®) technology, a tiled amplicon-based library prep chemistry, designed by Pillar's AI-empowered VersaTile™ Primer Design tool, for efficient single-tube target enrichment.

## oncoReveal® HRD v2 Panel (33 genes)

|        |        |        |        |         |        |
|--------|--------|--------|--------|---------|--------|
| ARID1A | BRCA2  | ESR1   | FEN1   | PPP2R1A | RAD51D |
| ATM    | BRIP1  | FANCA  | KRAS   | PTEN    | RAD54L |
| ATR    | CDK12  | FANCC  | MRE11A | RAD50   | TP53   |
| BARD1  | CHEK2  | FANCD2 | NBN    | RAD51   |        |
| BRAF   | CTNNB1 | FANCE  | PALB2  | RAD51B  |        |
| BRCA1  | ERBB2  | FANCF  | PIK3CA | RAD51C  |        |

Genes marked in green indicate full CDS coverage

### Simple NGS library prep workflow

Maintain control of samples and results with single-tube, tiled amplification that can be performed in-house by any NGS lab

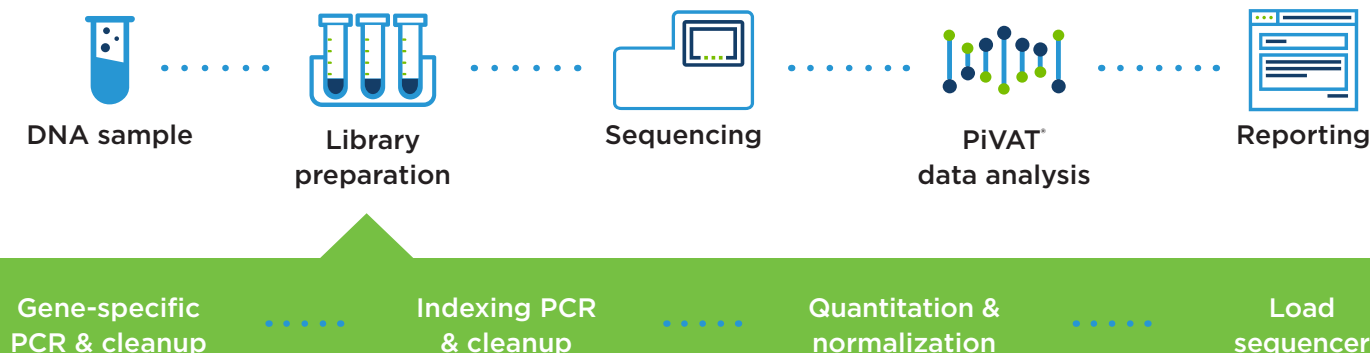
### Sensitive and robust chemistry

Achieve variant detection as low as 1% VAF<sup>†</sup> even with limited DNA input or poor sample quality

### Reduced fully-loaded lab costs

Improve lab efficiency and reduce “no calls”, repeat testing, and difficult interpretation decisions

### Simple, one-day workflow



<sup>†</sup> VAF, variant allele frequency

For Research Use Only. Not for use in diagnostic procedures.

## Panel specifications\*

|                                                             |                                                                                                                                |
|-------------------------------------------------------------|--------------------------------------------------------------------------------------------------------------------------------|
| Enrichment chemistry                                        | Multiplex PCR using tiled amplicons                                                                                            |
| Number of pools                                             | 1 pool                                                                                                                         |
| Number of genes/amplicons                                   | 33/1,314                                                                                                                       |
| Number of targets                                           | Full CDS coverage and exon-level CNVs for BRCA1 & BRCA2 genes, hotspots in 31 additional HRD-related genes; 114.5kb total size |
| Variant types                                               | SNVs, indels, exon level CNVs for BRCA1 & BRCA2                                                                                |
| Average amplicon size                                       | 157bp                                                                                                                          |
| Recommended DNA input range                                 | 20ng to 80ng                                                                                                                   |
| Sample types                                                | DNA from tissue, blood, or FFPE                                                                                                |
| Mapping rate                                                | 97.9% ± 1.7%                                                                                                                   |
| % on-target aligned reads                                   | 91.1% ± 0.8%                                                                                                                   |
| Coverage uniformity<br>(% targets with >0.2X mean coverage) | 92.5% ± 0.9%                                                                                                                   |
| Recommended Reads Per Sample                                | ~6.5 million to 7.5 million paired-end reads                                                                                   |
| Total assay time (from DNA to sequencer)                    | <8 hours                                                                                                                       |

\* Mapping rate, percentage of on-target aligned reads, and coverage uniformity metrics are based on internal testing performed using reference standard materials

## Ordering information

Select the panel AND one of the index kit options listed below.

| Panel                                   | Part number    |
|-----------------------------------------|----------------|
| oncoReveal® HRD v2 Panel (24 reactions) | HDA-HR-1002-24 |

| Pillar Index Kit options          | Reactions                      | Part number     |
|-----------------------------------|--------------------------------|-----------------|
| Pillar Custom Index Primers Kit A | 32 Combinations, 96 reactions  | IDX-PI-1001-96  |
| Pillar Custom Index Primers Kit D | 96 Combinations, 192 reactions | IDX-PI-1004-192 |

**TO ORDER OR LEARN MORE:**  
[pillarbiosci.com](https://pillarbiosci.com)

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