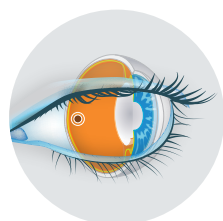


STARGARDT MASTR is a ready to use molecular research assay for early detection of *ABCA4*, *ELOVL4* and *CNGB3* variants associated with Stargardt macular dystrophy. This assay allows the analysis of SNVs, indels and CNVs in one single sequencing run.



Research application

- For the detection of variants (SNVs and CNVs), in 3 selected genes associated with Stargardt on blood-derived DNA using Illumina MiSeq®

Assay characteristics

| | |
|--------------------------------|---|
| Genes | <i>ABCA4</i> , <i>ELOVL4</i> and <i>CNGB3</i> |
| Genomic region analyzed | 21.7 kb |
| Number of amplicons | 91 |
| Amplicon length | 300 – 430 bp |
| Number of plexes | 4 |
| Verified with NGS System | Illumina MiSeq |
| Designed to be compatible with | Illumina NGS systems |

Performance Characteristics

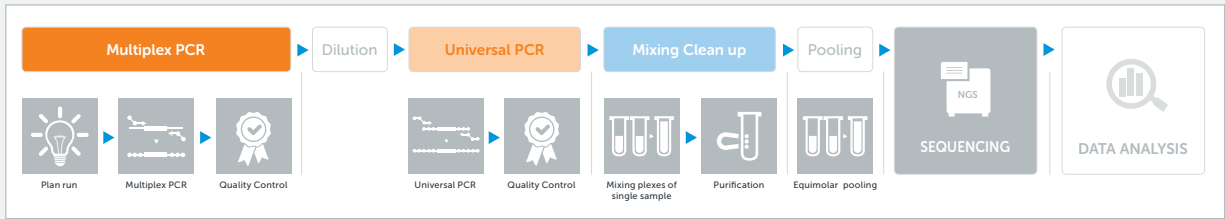
| | |
|--|--------------------------|
| Uniformity of amplification (0.2X mean coverage) | ≥ 98 % |
| On target read count | ≥ 96 % |
| DNA input | as low as 20 ng per plex |

Advised maximum number of samples per run:

| | | | |
|------------------------------------|-----------------------|------------------|------------------|
| Sequencing System | Illumina MiSeq® | | |
| | Reagent kit | | |
| Flow cell | Nano v2 2 x 250 bp | v2 2 x 250 bp | v3 2 x 300 bp |
| For SNV only | | | |
| Minimal coverage per allele: 20 | 46 | 690* | 1266* |
| For SNV and CNV | | | |
| Minimal coverage per amplicon: 200 | 9 | 139 | 254* |

*only 192 MID combinations available.

Workflow



Order info

| Cat. No. | Product Name | Product type | Reactions |
|-------------|-----------------|--------------|-----------|
| MR-0171.024 | STARGARDT MASTR | Research | 24 |

MID (Molecular Identifiers) kits are necessary to complete the workflow.

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

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