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Agilent Technologies

Human Genome CGH Microarray 105A

The Agilent Human Genome CGH Microarray 105A is a high-resolution tool for genome-wide DNA variation profiling without amplification or complexity reduction. Comprehensive probe coverage spans both coding and noncoding regions, with emphasis on well-known genes, promoters, miRNAs, and telomeric regions. Probe design and selection have been carefully optimized and validated for maximal sensitivity and specificity. Because the 105A format has the flexibility to run two samples on a single chip, it can significantly reduce costs

Features

- 99,000+ coding and noncoding human sequences represented
- Content sourced from UCSC hg17 human genome (NCBI build 35, May 2004)
- **15.0 KB** average probe spatial resolution
- **0.5 µg** total genomic DNA input requirement
- 2 x 105K slide format printed.

For further information or a quote, please contact m.herman@dnavigation.be