Resource Summary Report

Generated by RRID on Aug 5, 2026

Agilent: Oligo Pro II System

RRID:SCR_019529
Type: Tool

Proper Citation

Agilent: Oligo Pro II System (RRID:SCR_019529)

Resource Information

URL: https://www.agilent.com/en/product/automated-electrophoresis/oligo-pro-ii-systems/oligo-pro-ii-system-365769

Proper Citation: Agilent: Oligo Pro II System (RRID:SCR_019529)

Description: Oligo Pro II System delivers accurate purity assessments of ssDNA and ssRNA oligonucleotides without use of dyes.

Resource Type: instrument resource

Keywords: Agilent, Oligo Pro II System, Instrument Equipment, USEDit,

Availability: Commercially available

Specification URL: https://raw.githubusercontent.com/SciCrunch/RRID-Instruments/main/PDF/SCR_019529.pdf

Resource Name: Agilent: Oligo Pro II System

Resource ID: SCR_019529

Alternate IDs: Model_Number_Oligo_Pro_II

Record Creation Time: 20220129T080345+0000

Record Last Update: 20260801T190626+0000

Ratings and Alerts

No rating or validation information has been found for Agilent: Oligo Pro II System.

No alerts have been found for Agilent: Oligo Pro II System.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 14 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

Ismail D, et al. (2024) Interstitial 11q deletion in a patient with Sprengel's deformity: a case report and review of the literature. Molecular cytogenetics, 17(1), 30.

Wijesiriwardhana P, et al. (2021) Copy Number Variants Captured by the Array Comparative Genomic Hybridization in a Cohort of Patients Affected with Hereditary Colorectal Cancer in Sri Lanka: The First CNV Analysis Study of the Hereditary Colorectal Cancer in the Sri Lankan Population. Asian Pacific journal of cancer prevention : APJCP, 22(6), 1957.

Raftopoulou C, et al. (2020) Karyotypic Flexibility of the Complex Cancer Genome and the Role of Polyploidization in Maintenance of Structural Integrity of Cancer Chromosomes. Cancers, 12(3).

Schleicher S, et al. (2020) Establishment and Characterization of a Sclerosing Spindle Cell Rhabdomyosarcoma Cell Line with a Complex Genomic Profile. Cells, 9(12).

Felicio PS, et al. (2018) Genetic alterations detected by comparative genomic hybridization in BRCAX breast and ovarian cancers of Brazilian population. Oncotarget, 9(44), 27525.

Liu D, et al. (2018) Integrated Genome-Wide Analysis of Gene Expression and DNA Copy Number Variations Highlights Stem Cell-Related Pathways in Small Cell Esophageal Carcinoma. Stem cells international, 2018, 3481783.

Ghosh S, et al. (2014) Copy number variation in the horse genome. PLoS genetics, 10(10), e1004712.

Chong WW, et al. (2014) Performance of chromosomal microarray for patients with intellectual disabilities/developmental delay, autism, and multiple congenital anomalies in a Chinese cohort. Molecular cytogenetics, 7, 34.

Ottolini B, et al. (2014) Evidence of convergent evolution in humans and macaques supports an adaptive role for copy number variation of the ??-defensin-2 gene. Genome biology and evolution, 6(11), 3025.

Qi LN, et al. (2013) Genome-wide and differential proteomic analysis of hepatitis B virus and aflatoxin B1 related hepatocellular carcinoma in Guangxi, China. PloS one, 8(12), e83465.

Tian M, et al. (2013) Copy number variants in locally raised Chinese chicken genomes determined using array comparative genomic hybridization. BMC genomics, 14, 262.

Dunn B, et al. (2012) Analysis of the Saccharomyces cerevisiae pan-genome reveals a pool of copy number variants distributed in diverse yeast strains from differing industrial environments. Genome research, 22(5), 908.

Lagerstedt KK, et al. (2010) Genes with relevance for early to late progression of colon carcinoma based on combined genomic and transcriptomic information from the same patients. Cancer informatics, 9, 79.

Junnila S, et al. (2010) Genome-wide gene copy number and expression analysis of primary gastric tumors and gastric cancer cell lines. BMC cancer, 10, 73.