

Resource Summary Report

Generated by [kravitz2](#) on Aug 23, 2026

Revvity: LabChip GX Touch Nucleic Acid Analyzer

RRID:SCR_026830

Type: Tool

Proper Citation

Revvity: LabChip GX Touch Nucleic Acid Analyzer (RRID:SCR_026830)

Resource Information

URL: <https://www.revvity.com/gb-en/product/labchip-gx-touch-24-cls138162>

Proper Citation: Revvity: LabChip GX Touch Nucleic Acid Analyzer (RRID:SCR_026830)

Description: Automated microfluidic electrophoresis system that rapidly quantifies and sizes DNA and RNA with high sensitivity using minimal sample volume.

Synonyms: LabChip GX Touch Nucleic Acid Analyzer

Resource Type: instrument resource

Keywords: microfluidic electrophoresis system, DNA quantification, RNA quantification, nucleic acid analyzer,

Specification URL: https://raw.githubusercontent.com/SciCrunch/RRID-Instruments/refs/heads/main/PDF/SCR_026830.pdf

Resource Name: Revvity: LabChip GX Touch Nucleic Acid Analyzer

Resource ID: SCR_026830

Alternate IDs: Model_Number_LabChip_GX_Touch

Record Creation Time: 20250423T060241+0000

Record Last Update: 20260815T183251+0000

Ratings and Alerts

No rating or validation information has been found for Revvity: LabChip GX Touch Nucleic Acid Analyzer.

No alerts have been found for Revvity: LabChip GX Touch Nucleic Acid Analyzer.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Yoshimoto D, et al. (2026) Downregulation of HLA Class I Expression through HLA-A DNA Methylation Is Associated with Reduced CD8+ T-cell Infiltration in Cervical Cancer. Cancer immunology research, 14(1), 77.

Manso L, et al. (2025) A Phase IB Study of Binimetinib and Palbociclib in Molecularly Selected Advanced Triple-Negative Breast Cancer. Cancer research communications, 5(9), 1728.

Teng K, et al. (2021) Modeling High-Grade Serous Ovarian Carcinoma Using a Combination of In Vivo Fallopian Tube Electroporation and CRISPR-Cas9-Mediated Genome Editing. Cancer research, 81(20), 5147.

Zhang J, et al. (2020) Combining mechanistic and machine learning models for predictive engineering and optimization of tryptophan metabolism. Nature communications, 11(1), 4880.

Jan?? P, et al. (2020) Interactomic affinity profiling by holdup assay: Acetylation and distal residues impact the PDZome-binding specificity of PTEN phosphatase. PloS one, 15(12), e0244613.

Morin A, et al. (2016) Immunoseq: the identification of functionally relevant variants through targeted capture and sequencing of active regulatory regions in human immune cells. BMC medical genomics, 9(1), 59.