

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

Illumina: LabChip GXII Touch protein characterization system

RRID:SCR_018593

Type: Tool

Proper Citation

Illumina: LabChip GXII Touch protein characterization system (RRID:SCR_018593)

Resource Information

URL: <https://www.perkinelmer.com/Product/ht-labchip-gx-ii-touch-clis138160>

Proper Citation: Illumina: LabChip GXII Touch protein characterization system (RRID:SCR_018593)

Description: LabChip GXII Touch is a protein characterization system that provides reproducible quantitation, molecular weight sizing and percent purity analysis of protein samples. Leveraging microfluidic electrophoretic separation technology, the LabChip GXII Touch system enables rapid characterization with minimal sample preparation setup and sample input volume.

Synonyms: LabChip GXII Touch

Resource Type: instrument resource

Keywords: Protein Characterization System, Instrument, Equipment, Illumina, USEDit, ABRF

Availability: Restricted

Resource Name: Illumina: LabChip GXII Touch protein characterization system

Resource ID: SCR_018593

Alternate IDs: Model_Number_GX_II, SCR_020135

Alternate URLs: https://resources.perkinelmer.com/lab-solutions/resources/docs/prd_labchip-gxii-touch-protein-portfolio_ag011808_27_pn.pdf

Record Creation Time: 20220129T080341+0000

Record Last Update: 20260801T190622+0000

Ratings and Alerts

No rating or validation information has been found for Illumina: LabChip GXII Touch protein characterization system.

No alerts have been found for Illumina: LabChip GXII Touch protein characterization system.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Jeremian R, et al. (2025) Investigating the cell of origin and novel molecular targets in Merkel cell carcinoma: a historic misnomer. Molecular oncology.

Conn VM, et al. (2024) Use of synthetic circular RNA spike-ins (SynCRS) for normalization of circular RNA sequencing data. Nature protocols.