

# Resource Summary Report

Generated by [dkNET](#) on Sep 12, 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](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:** 20260905T132839+0000

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## 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.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

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.