

Resource Summary Report

Generated by RRID on Aug 13, 2026

Illumina Experiment Manager

RRID:SCR_021202

Type: Tool

Proper Citation

Illumina Experiment Manager (RRID:SCR_021202)

Resource Information

URL:

https://emea.support.illumina.com/sequencing/sequencing_software/experiment_manager/download

Proper Citation: Illumina Experiment Manager (RRID:SCR_021202)

Description: Software to create and edit sample sheet in order to perform next generation sequencing on Illumina platforms. Used for MiniSeq, MiSeq, NextSeq 500/550, HiSeq, and NovaSeq 6000 Systems.

Abbreviations: IEM

Synonyms: Illumina Experiment Manager software, Illumina Experiment Manager v1.18.1, Illumina Experiment Manager v1.19.1

Resource Type: data management software, software resource, software application

Keywords: Create sample sheet, edit sample sheet, next generation sequencing, Illumina platforms

Availability: Free, Available for download, Freely available

Resource Name: Illumina Experiment Manager

Resource ID: SCR_021202

Record Creation Time: 20220129T080354+0000

Record Last Update: 20260812T044408+0000

Ratings and Alerts

No rating or validation information has been found for Illumina Experiment Manager.

No alerts have been found for Illumina Experiment Manager.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Brotsky AL, et al. (2026) Combination Therapy with Copanlisib and Niraparib in Patients with Recurrent Endometrial and Ovarian Cancer (COPANIRA): Efficacy, Toxicity, and Translational Insights. Clinical cancer research : an official journal of the American Association for Cancer Research, 32(7), 1234.

Sebastiani G, et al. (2024) A set of circulating microRNAs belonging to the 14q32 chromosome locus identifies two subgroups of individuals with recent-onset type 1 diabetes. Cell reports. Medicine, 5(6), 101591.

Grieco GE, et al. (2021) Protocol to analyze circulating small non-coding RNAs by high-throughput RNA sequencing from human plasma samples. STAR protocols, 2(3), 100606.