

# Resource Summary Report

Generated by [kravitz2](#) on Sep 4, 2026

## University of Edinburgh Genomics Core Facility

RRID:SCR\_025433

Type: Tool

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### Proper Citation

University of Edinburgh Genomics Core Facility (RRID:SCR\_025433)

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### Resource Information

**URL:** <https://clinical-research-facility.ed.ac.uk/core-services/genetics>

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**Description:** Core provides secure receipt, processing, archiving and analysis of biological samples. Offers support for clinical research from sample collection to genetic analysis, working to the principles of Good Clinical Practice for Laboratories. Facility provides access to genomic technologies and services including next-generation sequencing (Illumina), third-generation sequencing (PacBio and Oxford Nanopore), bespoke bioinformatics analysis, and genomics and bioinformatics training.

**Synonyms:** Edinburgh Genomics Core Facility

**Resource Type:** access service resource, core facility, service resource

**Keywords:** Genomics, Sequencing, Microarrays, Digital PCR, DNA & RNA extractions; Illumina, Element, PacBio, ONT

**Availability:** Open

**Resource Name:** University of Edinburgh Genomics Core Facility

**Resource ID:** SCR\_025433

**Alternate IDs:** ABRF\_5723

**Alternate URLs:** <https://coremarketplace.org/?FacilityID=5723&citation=1>

**Record Creation Time:** 20240628T053249+0000

**Record Last Update:** 20260904T000942+0000

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### Ratings and Alerts

No rating or validation information has been found for University of Edinburgh Genomics Core Facility.

No alerts have been found for University of Edinburgh Genomics Core Facility.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Vancaester E, et al. (2025) Ghosts of symbionts past: the hidden history of the dynamic association between filarial nematodes and their Wolbachia endosymbionts. G3 (Bethesda, Md.), 15(12).

Durmaz Mitchell E, et al. (2025) A Balanced Inversion Polymorphism Exhibits a Dominance Reversal at the Gene Expression Level that Depends on Developmental Context. Molecular biology and evolution, 42(9).

Tijjani A, et al. (2024) Genomic Reference Resource for African Cattle: Genome Sequences and High-Density Array Variants. Scientific data, 11(1), 801.