Resource Summary Report

Generated by Kravitz on Sep 10, 2026

University of Edinburgh Genomics Core Facility

RRID:SCR_025433
Type: Tool

Proper Citation

University of Edinburgh Genomics Core Facility (RRID:SCR_025433)

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: 20260905T133458+0000

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.

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 Kravitz .

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.