

# Resource Summary Report

Generated by [kravitz2](#) on Aug 1, 2026

## CES Genome Quebec Core Facility

RRID:SCR\_017703

Type: Tool

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### Proper Citation

CES Genome Quebec Core Facility (RRID:SCR\_017703)

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### Resource Information

**URL:** <https://cesgq.com/>

**Proper Citation:** CES Genome Quebec Core Facility (RRID:SCR\_017703)

**Description:** Core for full DNA and RNA analysis services. Canada's largest integrated provider of sequencing, genotyping and biochips services. Offers scientific community around the world genomic services in support of research in human health, agrifood, forestry and sustainable development.

**Synonyms:** Genome Quebec Center of Expertise and Services

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

**Keywords:** DNA, RNA, analysis, service, core, Canada, sequencing, genotyping, biochip

**Availability:** Open

**Resource Name:** CES Genome Quebec Core Facility

**Resource ID:** SCR\_017703

**Alternate IDs:** ABRF\_115, SCR\_017718

**Record Creation Time:** 20220129T080336+0000

**Record Last Update:** 20260731T043018+0000

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### Ratings and Alerts

No rating or validation information has been found for CES Genome Quebec Core Facility.

No alerts have been found for CES Genome Quebec Core Facility.

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### Data and Source Information

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

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## Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Iqbal I, et al. (2023) Interaction and association analysis of malting related traits in barley. PloS one, 18(4), e0283763.

Clerissi C, et al. (2023) A core of functional complementary bacteria infects oysters in Pacific Oyster Mortality Syndrome. Animal microbiome, 5(1), 26.

Smith SR, et al. (2022) A chromosome-anchored genome assembly for Lake Trout (*Salvelinus namaycush*). Molecular ecology resources, 22(2), 679.

Ibeagha-Awemu EM, et al. (2021) Whole Genome Methylation Analysis Reveals Role of DNA Methylation in Cow's Ileal and Ileal Lymph Node Responses to *Mycobacterium avium* subsp. *paratuberculosis* Infection. Frontiers in genetics, 12, 797490.

Sauriol A, et al. (2020) Modeling the Diversity of Epithelial Ovarian Cancer through Ten Novel Well Characterized Cell Lines Covering Multiple Subtypes of the Disease. Cancers, 12(8).