

Resource Summary Report

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University of Cape Town Centre for Proteomic and Genomic Research (CPGR) Core Facility

RRID:SCR_017158

Type: Tool

Proper Citation

University of Cape Town Centre for Proteomic and Genomic Research (CPGR) Core Facility (RRID:SCR_017158)

Resource Information

URL: <http://www.cpgr.org.za>

Proper Citation: University of Cape Town Centre for Proteomic and Genomic Research (CPGR) Core Facility (RRID:SCR_017158)

Description: Services to life science and biotech communities in South Africa. Based in Cape Town, combine information about genomic and proteomic technologies with bio computational pipelines to create fit for purpose offerings for customers in academia and industry.

Abbreviations: CPGR Core

Synonyms: Centre for Proteomic and Genomic Research, CPGR, core facility, University of Cape Town, Center for Proteomic and Genomic Research

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

Keywords: bioinformatics, omics, core, facility, genomic, proteomic

Availability: Restricted

Resource Name: University of Cape Town Centre for Proteomic and Genomic Research (CPGR) Core Facility

Resource ID: SCR_017158

Record Creation Time: 20220129T080333+0000

Record Last Update: 20260801T191222+0000

Ratings and Alerts

No rating or validation information has been found for University of Cape Town Centre for Proteomic and Genomic Research (CPGR) Core Facility.

No alerts have been found for University of Cape Town Centre for Proteomic and Genomic Research (CPGR) Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Taku O, et al. (2022) Molecular Identification of Cervical Microbes in HIV-Negative and HIV-Positive Women in an African Setting Using a Customized Bacterial Vaginosis Microbial DNA Quantitative PCR (qPCR) Array. Microbiology spectrum, 10(3), e0222921.

Baloyi NM, et al. (2017) Proteomic analysis of Arabidopsis plasma membranes reveals lipopolysaccharide-responsive changes. Biochemical and biophysical research communications, 486(4), 1137.