

Resource Summary Report

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Sequentia Biotech SL

RRID:SCR_012435

Type: Tool

Proper Citation

Sequentia Biotech SL (RRID:SCR_012435)

Resource Information

URL: <https://www.sequentiabiotech.com/>

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Description: Core aims to empower research institutes and private companies in the fields of bioinformatics, biostatistics, biotechnologies, data management and R&D. Core is organized into four specialized divisions: Sequentia Green, Sequentia Health, Sequentia Research, and Sequentia Training.

Abbreviations: Sequentia Biotech, Sequentia

Synonyms: , Sequentia Biotechnologies, Sequentia

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

Keywords: transcriptomics, Metagenomics, Epigenomics, Genomics, biostatistics, data management

Availability: Available to external user

Resource Name: Sequentia Biotech SL

Resource ID: SCR_012435

Alternate IDs: SciEx_13316

Alternate URLs: <http://www.scienceexchange.com/facilities/sequentia-biotech-sl>

Record Creation Time: 20220129T080310+0000

Record Last Update: 20260903T235146+0000

Ratings and Alerts

No rating or validation information has been found for Sequentia Biotech SL.

No alerts have been found for Sequentia Biotech SL.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 11 mentions in open access literature.

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

Tamburino R, et al. (2025) Genotype-specific transcriptomic response to drought stress in potato cultivars modulated by microbial biostimulants. *BMC plant biology*, 25(1), 1777.

Puente-Sánchez F, et al. (2024) Bacterioplankton taxa compete for iron along the early spring-summer transition in the Arctic Ocean. *Ecology and evolution*, 14(6), e11546.

Ruggiero A, et al. (2022) Transcriptomic and splicing changes underlying tomato responses to combined water and nutrient stress. *Frontiers in plant science*, 13, 974048.

Maradonna F, et al. (2022) A zebrafish HCT116 xenograft model to predict anandamide outcomes on colorectal cancer. *Cell death & disease*, 13(12), 1069.

Velasco-Aviles S, et al. (2022) A genetic compensatory mechanism regulated by Jun and Mef2d modulates the expression of distinct class IIa Hdacs to ensure peripheral nerve myelination and repair. *eLife*, 11.

Sánchez-Botet A, et al. (2021) Atypical cyclin P regulates cancer cell stemness through activation of the WNT pathway. *Cellular oncology (Dordrecht)*, 44(6), 1273.

Pinoli P, et al. (2021) VirusLab: A Tool for Customized SARS-CoV-2 Data Analysis. *Biotech (Basel (Switzerland))*, 10(4).

Warnecke A, et al. (2021) Successful Treatment of Noise-Induced Hearing Loss by Mesenchymal Stromal Cells: An RNAseq Analysis of Protective/Repair Pathways. *Frontiers in cellular neuroscience*, 15, 656930.

Thodberg S, et al. (2020) A flavin-dependent monooxygenase catalyzes the initial step in cyanogenic glycoside synthesis in ferns. *Communications biology*, 3(1), 507.

Russo MT, et al. (2018) MRP3 is a sex determining gene in the diatom *Pseudo-nitzschia multistriata*. *Nature communications*, 9(1), 5050.

Basu S, et al. (2017) Finding a partner in the ocean: molecular and evolutionary bases of the response to sexual cues in a planktonic diatom. *The New phytologist*, 215(1), 140.