

Resource Summary Report

Generated by SPARC SAWG on Sep 16, 2026

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: 20260912T195748+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 [SPARC SAWG](#) .

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.