

Resource Summary Report

Generated by [NIF](#) on Sep 11, 2026

Pipeline Pilot

RRID:SCR_014917

Type: Tool

Proper Citation

Pipeline Pilot (RRID:SCR_014917)

Resource Information

URL: <http://accelrys.com/products/collaborative-science/biovia-pipeline-pilot/>

Proper Citation: Pipeline Pilot (RRID:SCR_014917)

Description: Software used to automate the process of accessing, analyzing and reporting scientific data. This software can be used by a person with little or no software development experience can create scientific protocols that can be executed through a variety of interfaces including: BIOVIA Web Port, other BIOVIA solutions such as BIOVIA Electronic Lab Notebook, Isentris, Chemical Registration and third-party applications such as Microsoft SharePoint. The protocols aggregate and provide immediate access to volumes of research data, they automate the scientific analysis of data and allow researchers to explore, visualize and report results.

Resource Type: data analysis software, data processing software, software application, software resource

Keywords: automation, accessing, analysis, analyzing, scientific data, aggregation. aggregate, research, scientific

Availability: Commercial

Resource Name: Pipeline Pilot

Resource ID: SCR_014917

Record Creation Time: 20220129T080323+0000

Record Last Update: 20260905T133006+0000

Ratings and Alerts

No rating or validation information has been found for Pipeline Pilot.

No alerts have been found for Pipeline Pilot.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 358 mentions in open access literature.

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

Akabane T, et al. (2026) Chemical inhibition of INO1 reduces phytic acid in rice and wheat grains for enhanced micronutrient bioavailability. *Nature food*, 7(2), 163.

Marreddy RKR, et al. (2026) Development of a high-throughput screening platform for C. difficile toxin synthesis inhibitors unveils meclizine as an antivirulence agent. *Antimicrobial agents and chemotherapy*, 70(2), e0096025.

Calvo Fernández E, et al. (2026) Systematic design of combination therapy by targeting master regulators of coexisting diffuse midline glioma cell states. *Nature genetics*, 58(5), 1112.

Wu YW, et al. (2026) Consensus Pharmacological Interactions for PLK2 Inhibitor Identification in Colorectal Cancer Treatment. *Journal of chemical information and modeling*, 66(1), 577.

Bahia IAF, et al. (2026) Comparative pharmacoinformatic and quantum descriptor insights from BFM/GBTLI guidelines to phase I/II compounds for acute lymphoblastic leukemia (ALL). *Scientific reports*, 16(1).

Svensson E, et al. (2025) Network Analysis of the Organic Chemistry in Patents, Literature, and Pharmaceutical Industry. *Molecular informatics*, 44(7), e202500011.

Yang Z, et al. (2025) A force-sensitive adhesion GPCR is required for equilibrioception. *Cell research*, 35(4), 243.

Okabe A, et al. (2025) Discovery of Highly Potent Noncovalent Inhibitors of SARS-CoV-2 Main Protease through Computer-Aided Drug Design. *Journal of medicinal chemistry*, 68(20), 21330.

Cho NC, et al. (2025) The first South Korean data challenge for drug discovery using human and mouse liver microsomal stability data. *Journal of cheminformatics*, 17(1), 139.

Wang J, et al. (2025) Decoding the selective chemical modulation of CYP3A4. *Nature communications*, 16(1), 3423.

Yang X, et al. (2025) Predicted molecules followed by experimental validation for protecting human neurons from oxidative stress-induced cytotoxicity. *Proceedings of the National Academy of Sciences of the United States of America*, 122(45), e2505359122.

Esparza M, et al. (2025) Contrasting interferon-mediated antiviral responses in human lung adenocarcinoma cells. *Journal of virology*, 99(6), e0046925.

Dvorak NM, et al. (2025) Enhanced motivated behavior mediated by pharmacological targeting of the FGF14/Nav1.6 complex in nucleus accumbens neurons. *Nature communications*, 16(1), 110.

Yang CL, et al. (2025) Identification and Biological Evaluation of a Novel CLK4 Inhibitor Targeting Alternative Splicing in Pancreatic Cancer Using Structure-Based Virtual Screening. *Advanced science* (Weinheim, Baden-Wurttemberg, Germany), 12(19), e2416323.

Scott T, et al. (2025) Drugit: crowd-sourcing molecular design of non-peptidic VHL binders. *Nature communications*, 16(1), 3548.

Mettu A, et al. (2025) Functionally active modulators targeting the LRRK2 WD40 repeat domain identified by FRASE-bot in CACHE Challenge #1. *Chemical science*, 16(8), 3430.

McSwiggen DT, et al. (2025) A high-throughput platform for single-molecule tracking identifies drug interaction and cellular mechanisms. *eLife*, 12.

Khalid AQ, et al. (2025) Protocol for predicting γ -tocotrienol and δ -tocotrienol binding to colorectal cancer-related proteins using Schrödinger's Maestro and Glide. *STAR protocols*, 6(4), 104144.

Shan P, et al. (2025) Discovery of a novel potent tubulin inhibitor through virtual screening and target validation for cancer chemotherapy. *Cell death discovery*, 11(1), 392.

Hsieh TH, et al. (2025) Hinokiflavone is a novel CK2 inhibitor promoting apoptosis and synergizing with chemotherapeutic agents in cisplatin resistant bladder cancer cells. *Scientific reports*, 15(1), 20922.