Resource Summary Report

Generated by Kravitz on Sep 10, 2026

Acellera

RRID:SCR_012200
Type: Tool

Proper Citation

Acellera (RRID:SCR_012200)

Resource Information

URL: https://www.acellera.com/

Proper Citation: Acellera (RRID:SCR_012200)

Description: Acellera is company focused on developing high-throughput molecular dynamics techniques that deliver solutions for estimating common physico-chemical properties such as binding affinities, kinetics, poses and pathways with validated accuracy.Company mission is to accelerate transition to rational, computerized drug discovery via simulations and machine learning.

Abbreviations: Acellera

Synonyms: Acellera Ltd

Resource Type: commercial organization, service resource

Resource Name: Acellera

Resource ID: SCR_012200

Alternate IDs: SciEx_10567

Old URLs: http://www.scienceexchange.com/facilities/acellera-ltd

Record Creation Time: 20220129T080309+0000

Record Last Update: 20260905T133351+0000

Ratings and Alerts

No rating or validation information has been found for Acellera.

No alerts have been found for Acellera.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 6 mentions in open access literature.

Listed below are recent publications. The full list is available at Kravitz .

Sabanés Zariquiey F, et al. (2025) QuantumBind-RBFE: Accurate Relative Binding Free Energy Calculations Using Neural Network Potentials. Journal of chemical information and modeling, 65(8), 4081.

Nicoli A, et al. (2023) Modeling the Orthosteric Binding Site of the G Protein-Coupled Odorant Receptor OR5K1. Journal of chemical information and modeling, 63(7), 2014.

Nicoli A, et al. (2023) Structure-Based Discovery of Mouse Trace Amine-Associated Receptor 5 Antagonists. Journal of chemical information and modeling, 63(21), 6667.

Wall MJ, et al. (2022) Selective activation of G?ob by an adenosine A1 receptor agonist elicits analgesia without cardiorespiratory depression. Nature communications, 13(1), 4150.

Standley M, et al. (2022) Experimental and In Silico Analysis of TEM ?-Lactamase Adaptive Evolution. ACS infectious diseases, 8(12), 2451.

Di Pizio A, et al. (2020) In Silico Molecular Study of Tryptophan Bitterness. Molecules (Basel, Switzerland), 25(20).