

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

Japan Bioinformatics

RRID:SCR_012250

Type: Tool

Proper Citation

Japan Bioinformatics (RRID:SCR_012250)

Resource Information

URL: <http://www.scienceexchange.com/facilities/japan-bioinformatics>

Proper Citation: Japan Bioinformatics (RRID:SCR_012250)

Description: Japan Bioinformatics KK is a Tokyo based biotechnology company specializing in bioinformatic software and data analysis. Products: Simbiot: Secure, easy-to-use, web and cloud-enabled software for management and analysis NGS data. SimGene: Free iOS and Android app for access to gene annotation information. SimAlign: Free iOS and Android app interfacing to NCBI BLAST+ service. Services include RNA-Seq, ChIP-Seq, miRNA-Seq, CAGE-Seq, mutation detection for amplicons, exomes and whole genomes as well as de-novo assembly and annotation of small genomes and transcriptomes. Project management services are also offered.

Abbreviations: Simbiot, SimAlign, SimGene

Synonyms: Simbiot, Japan Bioinformatics, SimAlign, SimGene

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

Keywords: Simbiot, SimAlign, Simgene

Availability: Available to external user

Resource Name: Japan Bioinformatics

Resource ID: SCR_012250

Alternate IDs: SciEx_11020

Record Creation Time: 20220129T080309+0000

Record Last Update: 20260805T044256+0000

Ratings and Alerts

No rating or validation information has been found for Japan Bioinformatics.

No alerts have been found for Japan Bioinformatics.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.