

# Resource Summary Report

Generated by NIF on Jul 31, 2026

## ODB - Operon database

RRID:SCR\_007827

Type: Tool

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### Proper Citation

ODB - Operon database (RRID:SCR\_007827)

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### Resource Information

**URL:** <http://odb.kuicr.kyoto-u.ac.jp/>

**Proper Citation:** ODB - Operon database (RRID:SCR\_007827)

**Description:** ODB (Operon DataBase) aims to collect known operons in multiple species and to offer a system to predict operons by user definitions. All the known operons are derived from the literature and from publicly available database including operon information. This system provides candidates of operons based on the conditions that users choice and also provide its prediction accuracy. This database integrates both known literature-based operons and as well as operon prediction, to provide a useful system for bioinformatics researchers and experimental biologists.

**Synonyms:** ODB

**Resource Type:** data or information resource, database

**Resource Name:** ODB - Operon database

**Resource ID:** SCR\_007827

**Alternate IDs:** nif-0000-03210

**Record Creation Time:** 20220129T080244+0000

**Record Last Update:** 20260729T044154+0000

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### Ratings and Alerts

No rating or validation information has been found for ODB - Operon database.

No alerts have been found for ODB - Operon database.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Janga SC, et al. (2006) The distinctive signatures of promoter regions and operon junctions across prokaryotes. Nucleic acids research, 34(14), 3980.