

# Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

## GenomeRing

RRID:SCR\_011801

Type: Tool

### Proper Citation

GenomeRing (RRID:SCR\_011801)

### Resource Information

**URL:** [http://www-ps.informatik.uni-tuebingen.de/itNew/?page\\_id=1160](http://www-ps.informatik.uni-tuebingen.de/itNew/?page_id=1160)

**Proper Citation:** GenomeRing (RRID:SCR\_011801)

**Description:** Alignment visualization based on SuperGenome coordinates.

**Abbreviations:** GenomeRing

**Resource Type:** software resource

**Resource Name:** GenomeRing

**Resource ID:** SCR\_011801

**Alternate IDs:** OMICS\_00937

**Record Creation Time:** 20220129T080306+0000

**Record Last Update:** 20260725T190724+0000

### Ratings and Alerts

No rating or validation information has been found for GenomeRing.

No alerts have been found for GenomeRing.

### Data and Source Information

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

### Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Goormaghtigh F, et al. (2018) Reassessing the Role of Type II Toxin-Antitoxin Systems in Formation of Escherichia coli Type II Persister Cells. mBio, 9(3).

Tran TT, et al. (2018) Functional analysis of African Xanthomonas oryzae pv. oryzae TALomes reveals a new susceptibility gene in bacterial leaf blight of rice. PLoS pathogens, 14(6), e1007092.