

# Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

## Gentle

RRID:SCR\_016127

Type: Tool

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### Proper Citation

Gentle (RRID:SCR\_016127)

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### Resource Information

**URL:** <http://gentle.magnusmanske.de>

**Proper Citation:** Gentle (RRID:SCR\_016127)

**Description:** Software for DNA and amino acid editing, database management, plasmid maps, It can also be used for restriction and ligation, alignments, sequencer data import, calculators, gel image display, PCR, and more.

**Synonyms:** GENTle

**Resource Type:** data analysis software, data processing software, sequence analysis software, software application, software resource, software toolkit

**Keywords:** editing, database, management, plasmid maps, restriction, ligation, alignments, sequence, data, import, calculator, gel, image, display, PCR, cloning, genetic

**Funding:** Free Software Foundation

**Availability:** Free, Available for download

**Resource Name:** Gentle

**Resource ID:** SCR\_016127

**Alternate IDs:** OMICS\_18307

**Alternate URLs:** <https://sources.debian.org/src/gentle/>

**License:** GNU General Public License, version 2

**Record Creation Time:** 20220129T080329+0000

**Record Last Update:** 20260912T195835+0000

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### Ratings and Alerts

No rating or validation information has been found for Gentle.

No alerts have been found for Gentle.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Musah HS, et al. (2024) Antimicrobial resistance and molecular characteristics of *Neisseria gonorrhoea* isolates in Ghana. *Access microbiology*, 6(2).

de Carvalho JB, et al. (2019) miRNA Genetic Variants Alter Their Secondary Structure and Expression in Patients With RASopathies Syndromes. *Frontiers in genetics*, 10, 1144.

Havlicek J, et al. (2018) Rapid Microarray-Based Detection of Rifampin, Isoniazid, and Fluoroquinolone Resistance in *Mycobacterium tuberculosis* by Use of a Single Cartridge. *Journal of clinical microbiology*, 56(2).

Sao Emani C, et al. (2018) Generation and characterization of thiol-deficient *Mycobacterium tuberculosis* mutants. *Scientific data*, 5, 180184.

Havlicek J, et al. (2017) An application of competitive reporter monitored amplification (CMA) for rapid detection of single nucleotide polymorphisms (SNPs). *PloS one*, 12(8), e0183561.

Sirena D, et al. (2005) The nucleotide sequence and a first generation gene transfer vector of species B human adenovirus serotype 3. *Virology*, 343(2), 283.