

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 19, 2026

MED

RRID:SCR_013403

Type: Tool

Proper Citation

MED (RRID:SCR_013403)

Resource Information

URL: <http://ctb.pku.edu.cn/main/SheGroup/Software/MED2.htm>

Proper Citation: MED (RRID:SCR_013403)

Description: A non-supervised gene prediction algorithm for prokaryotic genomes with multivariate entropy distance method.

Abbreviations: MED

Resource Type: software resource

Defining Citation: [PMID:17367537](#)

Keywords: prokaryotic, genome, multivariate entropy distance

Resource Name: MED

Resource ID: SCR_013403

Alternate IDs: OMICS_01489

Record Creation Time: 20220129T080315+0000

Record Last Update: 20260919T195242+0000

Ratings and Alerts

No rating or validation information has been found for MED.

No alerts have been found for MED.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [PRECISE-TBI](#) .

Zhu H, et al. (2007) MED: a new non-supervised gene prediction algorithm for bacterial and archaeal genomes. BMC bioinformatics, 8, 97.