

# Resource Summary Report

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## Core Gene Evaluation Script

RRID:SCR\_014604

Type: Tool

### Proper Citation

Core Gene Evaluation Script (RRID:SCR\_014604)

### Resource Information

**URL:** [http://hmpdacc.org/resources/tools\\_protocols.php](http://hmpdacc.org/resources/tools_protocols.php)

**Proper Citation:** Core Gene Evaluation Script (RRID:SCR\_014604)

**Description:** A tool used to screen for core gene sets as an indicator of completeness of draft genomes. The download includes a Perl script and required archaeal and bacterial core genes fasta and cluster files.

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

**Keywords:** microbiome, core gene, draft genome, indicator, microbe, microorganism, software

**Availability:** Available for download

**Resource Name:** Core Gene Evaluation Script

**Resource ID:** SCR\_014604

**Record Creation Time:** 20220129T080321+0000

**Record Last Update:** 20260805T044329+0000

### Ratings and Alerts

No rating or validation information has been found for Core Gene Evaluation Script.

No alerts have been found for Core Gene Evaluation Script.

### Data and Source Information

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

## Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Li J, et al. (2023) Arsenic-Containing Medicine Treatment Disturbed the Human Intestinal Microbial Flora. *Toxics*, 11(5).

Li S, et al. (2021) Helicobacter pylori infection is correlated with the incidence of erosive oral lichen planus and the alteration of the oral microbiome composition. *BMC microbiology*, 21(1), 122.

Patin NV, et al. (2020) The Role of the Gut Microbiome in Resisting Norovirus Infection as Revealed by a Human Challenge Study. *mBio*, 11(6).

Ji Y, et al. (2020) Analysis of by high-throughput sequencing: Helicobacter pylori infection and salivary microbiome. *BMC oral health*, 20(1), 84.

Richards PJ, et al. (2019) Phage Biocontrol of Campylobacter jejuni in Chickens Does Not Produce Collateral Effects on the Gut Microbiota. *Frontiers in microbiology*, 10, 476.

Ribeiro AA, et al. (2017) The oral bacterial microbiome of occlusal surfaces in children and its association with diet and caries. *PloS one*, 12(7), e0180621.