

# Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 18, 2026

## MFEprimer

RRID:SCR\_003066

Type: Tool

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### Proper Citation

MFEprimer (RRID:SCR\_003066)

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### Resource Information

**URL:** <https://github.com/quwubin/MFEprimer/>

**Proper Citation:** MFEprimer (RRID:SCR\_003066)

**Description:** A fast thermodynamics-based software program for checking PCR primer specificity against genomic DNA and mRNA/cDNA sequence databases.

**Synonyms:** MFEprimer-2.0

**Resource Type:** software resource

**Defining Citation:** [PMID:22689644](#)

**Keywords:** standalone software, bio.tools

**Availability:** Free, Available for download, Freely available

**Resource Name:** MFEprimer

**Resource ID:** SCR\_003066

**Alternate IDs:** biotools:mfeprimer-2.0, OMICS\_02355

**Alternate URLs:** <https://www.mfeprimer.com/>

**Record Creation Time:** 20220129T080217+0000

**Record Last Update:** 20260912T195551+0000

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### Ratings and Alerts

No rating or validation information has been found for MFEprimer.

No alerts have been found for MFEprimer.

## Data and Source Information

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

## Usage and Citation Metrics

We found 21 mentions in open access literature.

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

Jelen*?*i? A, et al. (2026) Simple, fast, reliable: multiplex digital PCR quantification of 19 genetically modified soybean events. *GM crops & food*, 17(1), 2635816.

Simons RB, et al. (2025) Investigating Single-Molecule Molecular Inversion Probes for Medium-Scale Targeted DNA Methylation Analysis. *Epigenomes*, 9(1).

Sun M, et al. (2025) Development of a single-nucleotide polymorphism panel genotyping system for genetic analysis of Chinese hamsters. *Animal models and experimental medicine*, 8(5), 916.

Llanquinao-Sandoval J, et al. (2025) Identifying a novel Mecp2-mediated epigenetic mechanism controlling Lonp1 in the hippocampus and its disruption by aging. *Scientific reports*, 15(1), 40887.

Yu Y, et al. (2025) Engineered crRNA Drives RPA-T7-CRISPR/Cas14a Cascade for Ultrasensitive Detection of ctDNA PIK3CA H1047R. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 12(44), e07126.

Zhang Z, et al. (2024) MiPRIME: an integrated and intelligent platform for mining primer and probe sequences of microbial species. *Bioinformatics (Oxford, England)*, 40(7).

Tang C, et al. (2024) High-throughput nanopore targeted sequencing for efficient drug resistance assay of *Mycobacterium tuberculosis*. *Frontiers in microbiology*, 15, 1331656.

Caballero Méndez A, et al. (2024) Development and performance evaluation of a qPCR-based assay for the fully automated detection of group B *Streptococcus* (GBS) on the Panther Fusion Open Access system. *Microbiology spectrum*, 12(6), e0005724.

Kalinkin AI, et al. (2023) Epigenomic Profiling Advises Therapeutic Potential of Leukotriene Receptor Inhibitors for a Subset of Triple-Negative Breast Tumors. *International journal of molecular sciences*, 24(24).

Kandel S, et al. (2023) Genomic Surveillance of SARS-CoV-2 Using Long-Range PCR Primers. *bioRxiv : the preprint server for biology*.

O'Callaghan JL, et al. (2021) Limitations of 16S rRNA Gene Sequencing to Characterize *Lactobacillus* Species in the Upper Genital Tract. *Frontiers in cell and developmental biology*, 9, 641921.

Leal-Calvo T, et al. (2021) A new paradigm for leprosy diagnosis based on host gene expression. *PLoS pathogens*, 17(10), e1009972.

Dutra HLC, et al. (2021) The impact of artificial selection for Wolbachia-mediated dengue virus blocking on phage WO. PLoS neglected tropical diseases, 15(7), e0009637.

Carneiro Dutra HL, et al. (2020) Wolbachia and Sirtuin-4 interaction is associated with alterations in host glucose metabolism and bacterial titer. PLoS pathogens, 16(10), e1008996.

Kawahara Y, et al. (2020) Mikan Genome Database (MiGD): integrated database of genome annotation, genomic diversity, and CAPS marker information for mandarin molecular breeding. Breeding science, 70(2), 200.

Mahapatra S, et al. (2020) Character-based identification system of scombrids from Indian waters for authentication and conservation purposes. Mitochondrial DNA. Part B, Resources, 5(3), 3221.

Johnston AD, et al. (2019) PrimerROC: accurate condition-independent dimer prediction using ROC analysis. Scientific reports, 9(1), 209.

Descalsota GIL, et al. (2018) Genome-Wide Association Mapping in a Rice MAGIC Plus Population Detects QTLs and Genes Useful for Biofortification. Frontiers in plant science, 9, 1347.

Collins RA, et al. (2018) Persistence of environmental DNA in marine systems. Communications biology, 1, 185.

Youngblut ND, et al. (2018) SIPSIm: A Modeling Toolkit to Predict Accuracy and Aid Design of DNA-SIP Experiments. Frontiers in microbiology, 9, 570.