

# Resource Summary Report

Generated by [ASWG](#) on Aug 23, 2026

## FastDMA

RRID:SCR\_010954

Type: Tool

### Proper Citation

FastDMA (RRID:SCR\_010954)

### Resource Information

**URL:** <http://bioinfo.au.tsinghua.edu.cn/software/fastdma/>

**Proper Citation:** FastDMA (RRID:SCR\_010954)

**Description:** A software analyzing Illumina Infinium HumanMethylation450 BeadChip data, which is featured as multiple core parallel computing.

**Abbreviations:** FastDMA

**Resource Type:** software resource

**Resource Name:** FastDMA

**Resource ID:** SCR\_010954

**Alternate IDs:** OMICS\_00794

**Record Creation Time:** 20220129T080301+0000

**Record Last Update:** 20260815T182403+0000

### Ratings and Alerts

No rating or validation information has been found for FastDMA.

No alerts have been found for FastDMA.

### 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 [ASWG](#) .

Wang D, et al. (2019) Integrative molecular analysis of metastatic hepatocellular carcinoma. BMC medical genomics, 12(1), 164.

Wu D, et al. (2013) FastDMA: an infinium humanmethylation450 beadchip analyzer. PloS one, 8(9), e74275.