

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Jul 31, 2026

## [IMA](#)

RRID:SCR\_010955

Type: Tool

### Proper Citation

IMA (RRID:SCR\_010955)

### Resource Information

**URL:** <http://www.rforge.net/IMA/>

**Proper Citation:** IMA (RRID:SCR\_010955)

**Description:** A software package designed to automate the pipeline for analyzing site-level and region-level methylation changes in epigenetic studies utilizing the 450K DNA methylation microarray.

**Abbreviations:** IMA

**Resource Type:** software resource

**Resource Name:** IMA

**Resource ID:** SCR\_010955

**Alternate IDs:** OMICS\_00795

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

**Record Last Update:** 20260725T190709+0000

### Ratings and Alerts

No rating or validation information has been found for IMA.

No alerts have been found for IMA.

### 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 [PRECISE-TBI](#) .

Wei S, et al. (2021) Ten Years of EWAS. Advanced science (Weinheim, Baden-Wurttemberg, Germany), 8(20), e2100727.

Li T, et al. (2020) Genome-wide analysis of DNA methylation identifies S100A13 as an epigenetic biomarker in individuals with chronic (? 30 years) type 2 diabetes without diabetic retinopathy. Clinical epigenetics, 12(1), 77.

Han Y, et al. (2016) Integrating Epigenomics into the Understanding of Biomedical Insight. Bioinformatics and biology insights, 10, 267.

Huang SK, et al. (2014) Lung fibroblasts from patients with idiopathic pulmonary fibrosis exhibit genome-wide differences in DNA methylation compared to fibroblasts from nonfibrotic lung. PloS one, 9(9), e107055.

Patil VK, et al. (2013) Interaction of prenatal maternal smoking, interleukin 13 genetic variants and DNA methylation influencing airflow and airway reactivity. Clinical epigenetics, 5(1), 22.

Wessely F, et al. (2012) Identification of DNA methylation biomarkers from Infinium arrays. Frontiers in genetics, 3, 161.