# Resource Summary Report

Generated by nidm-terms on Jul 29, 2026

## TMA-Combiner

RRID:SCR_005600
Type: Tool

### Proper Citation

TMA-Combiner (RRID:SCR_005600)

### Resource Information

**URL:** http://genome-www.stanford.edu/TMA/combiner/

**Proper Citation:** TMA-Combiner (RRID:SCR_005600)

**Description:** A Simple Software Tool to Permit Analysis of Replicate Cores on Tissue Microarrays., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

**Abbreviations:** TMA-Combiner

**Resource Type:** software resource

**Defining Citation:** PMID:16258508

**Keywords:** tissue microarray

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** TMA-Combiner

**Resource ID:** SCR_005600

**Alternate IDs:** OMICS_00822

**Record Creation Time:** 20220129T080231+0000

**Record Last Update:** 20260725T190607+0000

### Ratings and Alerts

No rating or validation information has been found for TMA-Combiner.

No alerts have been found for TMA-Combiner.

## 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 nidm-terms .

Espinosa I, et al. (2009) Coordinate expression of colony-stimulating factor-1 and colony-stimulating factor-1-related proteins is associated with poor prognosis in gynecological and nongynecological leiomyosarcoma. The American journal of pathology, 174(6), 2347.