

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## TMA-Combiner

RRID:SCR\_005600

Type: Tool

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### Proper Citation

TMA-Combiner (RRID:SCR\_005600)

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### 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:** 20260905T132538+0000

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### Ratings and Alerts

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

No alerts have been found for TMA-Combiner.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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.