

Resource Summary Report

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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: 20260829T182219+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 [NIF](#) .

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.