

# Resource Summary Report

Generated by [NIF](#) on Sep 3, 2026

## RPA

RRID:SCR\_012768

Type: Tool

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### Proper Citation

RPA (RRID:SCR\_012768)

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### Resource Information

**URL:** <http://bioconductor.org/packages/devel/bioc/html/RPA.html>

**Proper Citation:** RPA (RRID:SCR\_012768)

**Description:** A fully scalable online pre-processing algorithm for short oligonucleotide microarray atlases.

**Abbreviations:** RPA

**Synonyms:** RPA: Robust Probabilistic Averaging for probe-level analysis

**Resource Type:** software resource

**Resource Name:** RPA

**Resource ID:** SCR\_012768

**Alternate IDs:** OMICS\_00778

**Record Creation Time:** 20220129T080312+0000

**Record Last Update:** 20260829T182421+0000

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### Ratings and Alerts

No rating or validation information has been found for RPA.

No alerts have been found for RPA.

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### 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](#) .

Lahti L, et al. (2013) A fully scalable online pre-processing algorithm for short oligonucleotide microarray atlases. Nucleic acids research, 41(10), e110.