

# Resource Summary Report

Generated by SPARC SAWG on Sep 12, 2026

## methVisual

RRID:SCR\_006705

Type: Tool

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### Proper Citation

methVisual (RRID:SCR\_006705)

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### Resource Information

**URL:** <http://www.bioconductor.org/packages/2.12/bioc/html/methVisual.html>

**Proper Citation:** methVisual (RRID:SCR\_006705)

**Description:** Software package that allows the visualization of DNA methylation data after bisulfite sequencing.

**Abbreviations:** methVisual

**Synonyms:** methVisual - Methods for visualization and statistics on DNA methylation data

**Resource Type:** software resource

**Availability:** GNU General Public License, v2 or greater

**Resource Name:** methVisual

**Resource ID:** SCR\_006705

**Alternate IDs:** OMICS\_00604

**Record Creation Time:** 20220129T080237+0000

**Record Last Update:** 20260912T195644+0000

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### Ratings and Alerts

No rating or validation information has been found for methVisual.

No alerts have been found for methVisual.

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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 [SPARC SAWG](#) .

Tobias IC, et al. (2020) Targeted expression profiling reveals distinct stages of early canine fibroblast reprogramming are regulated by 2-oxoglutarate hydroxylases. Stem cell research & therapy, 11(1), 528.