

# Resource Summary Report

Generated by [RRID](#) on Aug 2, 2026

## PAFScaff

RRID:SCR\_017976

Type: Tool

### Proper Citation

PAFScaff (RRID:SCR\_017976)

### Resource Information

**URL:** <https://github.com/slimsuite/pafscaff>

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**Description:** Software as Pairwise mApping Format reference based Scaffold anchoring and super scaffolding tool. Dsigned for mapping genome assembly scaffolds to closely related chromosome level reference genome assembly.

**Synonyms:** Pairwise mApping Format reference-based Scaffold

**Resource Type:** data analysis software, software application, software resource, sequence analysis software, data processing software

**Keywords:** Pairwise, mapping, reference, scaffold, genomics, scaffolding, assembly, genome, chromosome, bio.tools

**Availability:** Free, Freely available

**Resource Name:** PAFScaff

**Resource ID:** SCR\_017976

**Alternate IDs:** biotools:PAFScaff

**Alternate URLs:** <https://github.com/slimsuite/pafscaff/blob/master/PAFScaff.md>, <https://slimsuite.github.io/pafscaff/>, <https://bio.tools/PAFScaff>

**License:** GNU General Public License v3.0

**Record Creation Time:** 20220129T080338+0000

**Record Last Update:** 20260801T190609+0000

### Ratings and Alerts

No rating or validation information has been found for PAFScaff.

No alerts have been found for PAFScaff.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Field MA, et al. (2022) The Australian dingo is an early offshoot of modern breed dogs. Science advances, 8(16), eabm5944.

Edwards RJ, et al. (2021) Chromosome-length genome assembly and structural variations of the primal Basenji dog (Canis lupus familiaris) genome. BMC genomics, 22(1), 188.

Martin BT, et al. (2021) ClineHelpR: an R package for genomic cline outlier detection and visualization. BMC bioinformatics, 22(1), 501.

Field MA, et al. (2020) Canfam\_GSD: De novo chromosome-length genome assembly of the German Shepherd Dog (Canis lupus familiaris) using a combination of long reads, optical mapping, and Hi-C. GigaScience, 9(4).