

# Resource Summary Report

Generated by [kravitz2](#) on Sep 20, 2026

## FPSAC

RRID:SCR\_000555

Type: Tool

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### Proper Citation

FPSAC (RRID:SCR\_000555)

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### Resource Information

**URL:** <http://paleogenomics.irmacs.sfu.ca/FPSAC/>

**Proper Citation:** FPSAC (RRID:SCR\_000555)

**Description:** Sogftware for fast Phylogenetic Scaffolding of Ancient Contigs.

**Abbreviations:** FPSAC

**Synonyms:** Fast Phylogenetic Scaffolding of Ancient Contigs (FPSAC) and application to the medieval Black Death agent, Fast Phylogenetic Scaffolding of Ancient Contigs, FPSAC: fast phylogenetic scaffolding of ancient contigs

**Resource Type:** software resource

**Defining Citation:** [PMID:24068034](#)

**Keywords:** genome, scaffolding, bio.tools

**Resource Name:** FPSAC

**Resource ID:** SCR\_000555

**Alternate IDs:** biotools:fpsac, OMICS\_00041

**Alternate URLs:** <https://bio.tools/fpsac>

**Record Creation Time:** 20220129T080202+0000

**Record Last Update:** 20260919T194925+0000

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### Ratings and Alerts

No rating or validation information has been found for FPSAC.

No alerts have been found for FPSAC.

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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 [kravitz2](#) .

Duchemin W, et al. (2015) Reconstruction of an ancestral Yersinia pestis genome and comparison with an ancient sequence. BMC genomics, 16 Suppl 10(Suppl 10), S9.