

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

FPSAC

RRID:SCR_000555

Type: Tool

Proper Citation

FPSAC (RRID:SCR_000555)

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: 20260725T190447+0000

Ratings and Alerts

No rating or validation information has been found for FPSAC.

No alerts have been found for FPSAC.

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 [nidm-terms](#) .

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.