

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 9, 2026

Birdseed

RRID:SCR_001122

Type: Tool

Proper Citation

Birdseed (RRID:SCR_001122)

Resource Information

URL: <http://www.broadinstitute.org/mpg/birdsuite/birdseed.html>

Proper Citation: Birdseed (RRID:SCR_001122)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 23, 2022. SNP genotyping algorithm that runs on the Affymetrix 500K, SNP5.0, and SNP6.0 platforms. This software runs a clustering algorithm and generally should be run on a sample at least 50 times. It utilizes an Expectation-Maximization (EM) algorithm to produce genotypes and confidence scores for every individual at every SNP.

Resource Type: software resource, algorithm resource

Defining Citation: [PMID:21734406](#), [PMID:20865519](#)

Keywords: snp, genotype, clustering, expectation maximization, algorithm, confidence score, single nucleotide polymorphism

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Birdseed

Resource ID: SCR_001122

Alternate IDs: OMICS_00704

Record Creation Time: 20220129T080205+0000

Record Last Update: 20260808T190449+0000

Ratings and Alerts

No rating or validation information has been found for Birdseed.

No alerts have been found for Birdseed.

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 [SPARC SAWG](#) .

, et al. (2017) Comprehensive and Integrated Genomic Characterization of Adult Soft Tissue Sarcomas. Cell, 171(4), 950.