

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 6, 2026

## Birdseed

RRID:SCR\_001122

Type: Tool

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### Proper Citation

Birdseed (RRID:SCR\_001122)

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### 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:** algorithm resource, software 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:** 20260806T054319+0000

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### 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 [PRECISE-TBI](#) .

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