

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 2, 2026

## PyroBayes

RRID:SCR\_003757

Type: Tool

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### Proper Citation

PyroBayes (RRID:SCR\_003757)

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### Resource Information

**URL:** <http://bioinformatics.bc.edu/marthlab/wiki/index.php/PyroBayes>

**Proper Citation:** PyroBayes (RRID:SCR\_003757)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE. Documented on January 6, 2023. A base caller for pyrosequences from the 454 Life Sciences sequencing machines.

**Abbreviations:** PyroBayes

**Resource Type:** software resource

**Defining Citation:** [PMID:18193056](#)

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** PyroBayes

**Resource ID:** SCR\_003757

**Alternate IDs:** OMICS\_01155

**Record Creation Time:** 20220129T080220+0000

**Record Last Update:** 20260801T190233+0000

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### Ratings and Alerts

No rating or validation information has been found for PyroBayes.

No alerts have been found for PyroBayes.

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### Data and Source Information

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

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## Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Crisol-Martínez E, et al. (2017) Sorghum and wheat differentially affect caecal microbiota and associated performance characteristics of meat chickens. PeerJ, 5, e3071.

Stanley D, et al. (2016) Bacteria within the Gastrointestinal Tract Microbiota Correlated with Improved Growth and Feed Conversion: Challenges Presented for the Identification of Performance Enhancing Probiotic Bacteria. Frontiers in microbiology, 7, 187.

Erzurumluoglu AM, et al. (2015) Identifying Highly Penetrant Disease Causal Mutations Using Next Generation Sequencing: Guide to Whole Process. BioMed research international, 2015, 923491.