

# Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://www.fdi-lab.org/) on Apr 16, 2025

## Reptile

RRID:SCR\_013075

Type: Tool

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### Proper Citation

Reptile (RRID:SCR\_013075)

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### Resource Information

**URL:** <http://aluru-sun.ece.iastate.edu/doku.php?id=reptile>

**Proper Citation:** Reptile (RRID:SCR\_013075)

**Description:** A software developed in C++ for correcting sequencing errors in short reads from next-gen sequencing platforms.

**Abbreviations:** Reptile

**Resource Type:** software resource

**Defining Citation:** [PMID:20834037](https://pubmed.ncbi.nlm.nih.gov/20834037/)

**Keywords:** bio.tools

**Funding:**

**Resource Name:** Reptile

**Resource ID:** SCR\_013075

**Alternate IDs:** biotools:reptile, OMICS\_01109

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

**Record Creation Time:** 20220129T080314+0000

**Record Last Update:** 20250410T070330+0000

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### Ratings and Alerts

No rating or validation information has been found for Reptile.

No alerts have been found for Reptile.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 25 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch.org](#).

Bansal J, et al. (2025) EEG-Based ADHD Classification Using Autoencoder Feature Extraction and ResNet with Double Augmented Attention Mechanism. *Brain sciences*, 15(1).

Paphitis K, et al. (2024) Salmonella Vitkin Outbreak Associated with Bearded Dragons, Canada and United States, 2020-2022. *Emerging infectious diseases*, 30(2), 225.

Oskyrko O, et al. (2024) ReptTraits: a comprehensive dataset of ecological traits in reptiles. *Scientific data*, 11(1), 243.

Pan X, et al. (2024) The optimization path of agricultural industry structure and intelligent transformation by deep learning. *Scientific reports*, 14(1), 29548.

Wen G, et al. (2024) MMOSurv: meta-learning for few-shot survival analysis with multi-omics data. *Bioinformatics (Oxford, England)*, 41(1).

Deng T, et al. (2024) Data modeling analysis of GFRP tubular filled concrete column based on small sample deep meta learning method. *PloS one*, 19(7), e0305038.

Baum ZMC, et al. (2023) Meta-Learning Initializations for Interactive Medical Image Registration. *IEEE transactions on medical imaging*, 42(3), 823.

Yang J, et al. (2022) A survey of few-shot learning in smart agriculture: developments, applications, and challenges. *Plant methods*, 18(1), 28.

Liu H, et al. (2021) DNA methylation atlas of the mouse brain at single-cell resolution. *Nature*, 598(7879), 120.

Barrett R, et al. (2021) Investigating Active Learning and Meta-Learning for Iterative Peptide Design. *Journal of chemical information and modeling*, 61(1), 95.

Yao Z, et al. (2021) A transcriptomic and epigenomic cell atlas of the mouse primary motor cortex. *Nature*, 598(7879), 103.

Goldenberg J, et al. (2021) Substrate thermal properties influence ventral brightness evolution in ectotherms. *Communications biology*, 4(1), 26.

Williams RJ, et al. (2021) Climate and habitat configuration limit range expansion and patterns of dispersal in a non-native lizard. *Ecology and evolution*, 11(7), 3332.

Marshall BM, et al. (2020) Thousands of reptile species threatened by under-regulated global trade. *Nature communications*, 11(1), 4738.

Sethi A, et al. (2020) Supervised enhancer prediction with epigenetic pattern recognition and targeted validation. *Nature methods*, 17(8), 807.

He Y, et al. (2020) Spatiotemporal DNA methylome dynamics of the developing mouse fetus. *Nature*, 583(7818), 752.

Ramisch A, et al. (2019) CRUP: a comprehensive framework to predict condition-specific regulatory units. *Genome biology*, 20(1), 227.

Marshall BM, et al. (2019) Exploring snake occurrence records: Spatial biases and marginal gains from accessible social media. *PeerJ*, 7, e8059.

Tran BV, et al. (2019) Occurrence data of terrestrial vertebrates of Son Tra Peninsula, Da Nang City, Vietnam. *Biodiversity data journal*, 7, e39233.

Peña JF, et al. (2016) Conserved expression of vertebrate microvillar gene homologs in choanocytes of freshwater sponges. *EvoDevo*, 7, 13.