

Resource Summary Report

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

ApiDots

RRID:SCR_001778

Type: Tool

Proper Citation

ApiDots (RRID:SCR_001778)

Resource Information

URL: <http://www.cbil.upenn.edu/apidots/>

Proper Citation: ApiDots (RRID:SCR_001778)

Description: Note: ApiDots is currently unavailable. For data on apicomplexan EST assemblies, please see EuPathDB. ApiDots is a database integrating mRNA/EST sequences from numerous Apicomplexan parasites. ESTs and mRNAs were clustered and further assembled to generate consensus sequences. These consensus sequences were then subjected to database searches against protein sequences and protein domain sequences. The underlying relational structure of this database allows researchers to analyze these data and pose biologically interesting questions.

Resource Type: data or information resource, database

Defining Citation: [PMID:12618375](#)

Keywords: est, apicomplexan parasite, mrna, protein domain sequence, protein sequence

Funding: USDA ;
Amerman Family Foundation ;
Merck Research Laboratories ;
Burroughs Wellcome Fund ;
NRI 99-35204-8615;
NIAID AI45806;
NIAID AI45806A1;
NIAID AI48121

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: ApiDots

Resource ID: SCR_001778

Alternate IDs: nif-0000-02556

Record Creation Time: 20220129T080209+0000

Record Last Update: 20260801T190911+0000

Ratings and Alerts

No rating or validation information has been found for ApiDots.

No alerts have been found for ApiDots.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Khan A, et al. (2005) Composite genome map and recombination parameters derived from three archetypal lineages of *Toxoplasma gondii*. *Nucleic acids research*, 33(9), 2980.

Schwarz JA, et al. (2005) A novel rhoptry protein in *Toxoplasma gondii* bradyzoites and merozoites. *Molecular and biochemical parasitology*, 144(2), 159.