

Resource Summary Report

Generated by SPARC SAWG on Aug 8, 2026

CoGenT++

RRID:SCR_000249

Type: Tool

Proper Citation

CoGenT++ (RRID:SCR_000249)

Resource Information

URL: <http://bioinformatics.oxfordjournals.org/content/21/19/3806.long>

Proper Citation: CoGenT++ (RRID:SCR_000249)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented August 26, 2016. A data environment for computational research in comparative and functional genomics, designed to address issues of consistency, reproducibility, scalability and accessibility., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Resource Type: database, data or information resource

Defining Citation: [PMID:16216832](#)

Keywords: computation, genomics, data environment, functional genomics

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: CoGenT++

Resource ID: SCR_000249

Alternate IDs: nif-0000-02674

Old URLs: http://cgg.ebi.ac.uk/cgg/cpp_sitemap.html

Record Creation Time: 20220129T080200+0000

Record Last Update: 20260808T190342+0000

Ratings and Alerts

No rating or validation information has been found for CoGenT++.

No alerts have been found for CoGenT++.

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](#) .

Goldovsky L, et al. (2005) CoGenT++: an extensive and extensible data environment for computational genomics. Bioinformatics (Oxford, England), 21(19), 3806.