

# Resource Summary Report

Generated by [RRID](#) on Aug 2, 2026

## [digitagCT](#)

RRID:SCR\_012027

Type: Tool

### Proper Citation

digitagCT (RRID:SCR\_012027)

### Resource Information

**URL:** <http://cractools.gforge.inria.fr/software/digitagct>

**Proper Citation:** digitagCT (RRID:SCR\_012027)

**Description:** Software that combines digital gene expression (DGE) tags, RNA-Seq, tiling array expression data and species-comparison to explore new transcriptional regions and their specific biological features, particularly tissue expression or conservation.

**Abbreviations:** digitagCT

**Resource Type:** software resource

**Defining Citation:** [PMID:24357408](#)

**Keywords:** digital gene expression, rna-seq

**Resource Name:** digitagCT

**Resource ID:** SCR\_012027

**Alternate IDs:** OMICS\_02178

**Record Creation Time:** 20220129T080308+0000

**Record Last Update:** 20260801T190434+0000

### Ratings and Alerts

No rating or validation information has been found for digitagCT.

No alerts have been found for digitagCT.

### Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Philippe N, et al. (2014) Combining DGE and RNA-sequencing data to identify new polyA+ non-coding transcripts in the human genome. Nucleic acids research, 42(5), 2820.