

Resource Summary Report

Generated by [ASWG](#) on Aug 23, 2026

AlignACE

RRID:SCR_010875

Type: Tool

Proper Citation

AlignACE (RRID:SCR_010875)

Resource Information

URL:

<http://www.webcitation.org/getfile?fileid=c6d148fcb4fde0ea6991ec319a7a3925d38f32bf>

Proper Citation: AlignACE (RRID:SCR_010875)

Description: A software program which finds sequence elements conserved in a set of DNA sequences.

Abbreviations: AlignACE

Resource Type: software resource

Defining Citation: [PMID:10698627](#)

Resource Name: AlignACE

Resource ID: SCR_010875

Alternate IDs: OMICS_00475

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260815T182402+0000

Ratings and Alerts

No rating or validation information has been found for AlignACE.

No alerts have been found for AlignACE.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Koramutla MK, et al. (2016) Strength, Stability, and cis-Motifs of In silico Identified Phloem-Specific Promoters in Brassica juncea (L.). Frontiers in plant science, 7, 457.

Sato MP, et al. (2016) Natural selection in a population of Drosophila melanogaster explained by changes in gene expression caused by sequence variation in core promoter regions. BMC evolutionary biology, 16, 35.

Gao D, et al. (2016) The role of regulator Eha in Edwardsiella tarda pathogenesis and virulence gene transcription. Microbial pathogenesis, 95, 216.

Karnik R, et al. (2015) Identification of Predictive Cis-Regulatory Elements Using a Discriminative Objective Function and a Dynamic Search Space. PloS one, 10(10), e0140557.

Benz C, et al. (2005) Messenger RNA processing sites in Trypanosoma brucei. Molecular and biochemical parasitology, 143(2), 125.