

Resource Summary Report

Generated by [ASWG](#) on Aug 3, 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: 20260801T190359+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.