ALTER
RRID:SCR_015968
Type: Tool

Proper Citation

ALTER (RRID:SCR_015968)

Resource Information

URL: http://sing.ei.uvigo.es/ALTER/

Description: Web application to perform program-oriented conversion of DNA and protein alignments and transform between multiple sequence alignment formats. ALTER focuses on the specifications of mainstream alignment and analysis programs rather than on the conversion among more or less specific formats.

Resource Name: ALTER

Proper Citation: ALTER (RRID:SCR_015968)

Resource Type: Resource, image analysis software, data analysis software, data processing software, alignment software, software application, sequence analysis software, web application, software resource

Keywords: software, genome, sequence, DNA, alignments, protein, bioinformatics, analysis, format, phylogenetics

Resource ID: SCR_015968

Funding Agency: European Research Council, INBIOMED initiative, Spanish Ministry of Science and Education, University of Vigo, Xunta de Galicia

References: PMID:20439312

Availability: Freely available, Free, Available for download

Website Status: Last checked up
Alternate URLs: https://github.com/sing-group/ALTER

Mentions Count: 37

Ratings and Alerts

No rating or validation information has been found for ALTER.

No alerts have been found for ALTER.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 37 mentions in open access literature.

**Listed below are recent publications.** The full list is available at FDI Lab - SciCrunch Infrastructure.


