Resource Summary Report

CUSHAW

RRID:SCR_005479
Type: Tool

Proper Citation

CUSHAW (RRID:SCR_005479)

Resource Information

URL: http://cushaw2.sourceforge.net/homepage.htm#latest

Proper Citation: CUSHAW (RRID:SCR_005479)

Description: Software package for next-generation sequencing read alignment that is fast and parallel gapped read alignment to large genomes, such as the human genome.

Abbreviations: CUSHAW

Synonyms: CUSHAW2, CUSHAW3

Resource Type: software resource

Defining Citation: PMID:22576173, PMID:24466273

Keywords: next-generation sequencing, read alignment, genome, alignment

Funding:

Resource Name: CUSHAW

Resource ID: SCR_005479

Alternate IDs: OMICS_00658

Record Creation Time: 20220129T080230+0000

Record Last Update: 20250420T014252+0000

Ratings and Alerts

No rating or validation information has been found for CUSHAW.

No alerts have been found for CUSHAW.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Schelkunov MI, et al. (2021) Genomic comparison of non-photosynthetic plants from the family Balanophoraceae with their photosynthetic relatives. PeerJ, 9, e12106.

Nath N, et al. (2020) Genome-Wide DNA Alterations in X-Irradiated Human Gingiva Fibroblasts. International journal of molecular sciences, 21(16).