Dotter

RRID:SCR_016080
Type: Tool

Proper Citation

Dotter (RRID:SCR_016080)

Resource Information

URL: http://www.sanger.ac.uk/science/tools/seqtools

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Description: Software for sequence alignment that is a graphical dot-matrix program for detailed comparison of two sequences.

Synonyms: Seqtools Dotter

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

Defining Citation: PMID:26801397

Keywords: sequence, alignment, graphical, dot-matrix, program, comparison, two, detail

Funding Agency: Wellcome Trust Grant, NHGRI

Availability: Free, Available for download

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Ratings and Alerts

No rating or validation information has been found for Dotter.
No alerts have been found for Dotter.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 78 mentions in open access literature.

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


Gelali E, et al. (2019) iFISH is a publically available resource enabling versatile DNA FISH to
study genome architecture. Nature communications, 10(1), 1636.


