Staden Package
RRID:SCR_005629
Type: Tool

Proper Citation

Staden Package (RRID:SCR_005629)

Resource Information

**URL:** [http://staden.sourceforge.net/](http://staden.sourceforge.net/)

**Description:** A fully developed set of DNA sequence assembly (Gap4 and Gap5), editing and analysis tools (Spin) for Unix, Linux, MacOSX and MS Windows.

**Resource Name:** Staden Package

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**Resource Type:** Resource, software resource

**Keywords:** c, unix/linux, sequence assembly, dna/protein analysis, spin, sequence alignment, genome, genome viewer, c++, fortran, tcl

**Resource ID:** SCR_005629

**Parent Organization:** SourceForge

**References:** [PMID: 20513662](https://www.ncbi.nlm.nih.gov/pubmed/20513662)

**Availability:** BSD License

**Website Status:** Last checked up

**Alternate IDs:** OMICS_00894

**Abbreviations:** Staden Package

**Mentions Count:** 54
Ratings and Alerts

No rating or validation information has been found for Staden Package.

No alerts have been found for Staden Package.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 54 mentions in open access literature.

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


Valente U, et al. (2017) Is there any clinical relevant difference between non mosaic Klinefelter Syndrome patients with or without Androgen Receptor variations? Scientific reports, 7(1), 3358.