Augustus
RRID:SCR_008417
Type: Tool

Proper Citation

Augustus (RRID:SCR_008417)

Resource Information

URL: http://bioinf.uni-greifswald.de/augustus/

Description: Software for gene prediction in eukaryotic genomic sequences. Serves as a basis for further steps in the analysis of sequenced and assembled eukaryotic genomes.

Resource Name: Augustus

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

Keywords: software, gene, prediction, eucaryotic, genomic, sequence

Resource ID: SCR_008417

Funding Agency: Deutsche Forschungsgemeinschaft (DFG), Ernst Moritz Arndt University of Greifswald, Institute for Mathematics and Computer Science, STA1009/6-1

References: PMID:23700307

Availability: Free, Available for download, Freely available

Website Status: Last checked up

Alternate IDs: nif-0000-30133, SCR_015981

Mentions Count: 684
No rating or validation information has been found for Augustus.

No alerts have been found for Augustus.

Data and Source Information
Source: SciCrunch Registry

Usage and Citation Metrics

We found 684 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch Infrastructure](#).


Gong N, et al. (2020) Divergent genes encoding the putative receptors for growth hormone and prolactin in sea lamprey display distinct patterns of expression. Scientific reports, 10(1), 1674.


Wu H, et al. (2020) Genome Assembly ofFurther Supports That and Belong to Different Sections. G3 (Bethesda, Md.), 10(2), 455-466.


