

Resource Summary Report

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Kazusa Arabidopsis data opening site

RRID:SCR_013511

Type: Tool

Proper Citation

Kazusa Arabidopsis data opening site (RRID:SCR_013511)

Resource Information

URL: <http://www.kazusa.or.jp/kaos/>

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Description: This site has been developed by Kazusa DNA Research Institute for the purpose of offering the science community the analyzed sequence data produced by a multi-national Arabidopsis genome sequencing project coordinated by the Arabidopsis Genome Initiatives (AGI). The aim of this service is to enable users to browse the annotated sequence data produced by all the sequencing teams of AGI through an user-friendly graphic display system and search engines. Gene structures proposed on the annotated sequences as well as those predicted by computer programs are presented and each graphic item has a hyperlink to detailed information of the corresponding area. The nucleotide sequence data deposited in GenBank by AGI was downloaded, re-computer-analyzed at Kazusa and parsed results are displayed graphically.

Synonyms: KAOS

Resource Type: data or information resource, database

Keywords: annoated, arabidopsis, dna, genome, sequence

Resource Name: Kazusa Arabidopsis data opening site

Resource ID: SCR_013511

Alternate IDs: nif-0000-20958

Record Creation Time: 20220129T080316+0000

Record Last Update: 20260729T044317+0000

Ratings and Alerts

No rating or validation information has been found for Kazusa Arabidopsis data opening site.

No alerts have been found for Kazusa Arabidopsis data opening site.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Novinec L, et al. (2024) New achievements in orbital angular momentum beam characterization using a Hartmann wavefront sensor and the Kirkpatrick-Baez active optical system KAOS. Journal of synchrotron radiation, 31(Pt 5), 1058.

Attia S, et al. (2019) Evaluation of Implant Success in Patients with Dental Aplasia. BioMed research international, 2019, 1680158.

Hecker A, et al. (2015) The impact of early surgical intervention in free intestinal perforation: a time-to-intervention pilot study. World journal of emergency surgery : WJES, 10, 54.

Hayashi M, et al. (2000) AtPex14p maintains peroxisomal functions by determining protein targeting to three kinds of plant peroxisomes. The EMBO journal, 19(21), 5701.

Somers DE, et al. (2000) ZEITLUPE encodes a novel clock-associated PAS protein from Arabidopsis. Cell, 101(3), 319.

Fluet A, et al. (2000) All about Arabidopsis. Current biology : CB, 10(13), R468.