

Resource Summary Report

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Arabidopsis Biological Resource Center

RRID:SCR_008136

Type: Tool

Proper Citation

Arabidopsis Biological Resource Center (RRID:SCR_008136)

Resource Information

URL: <http://www.biosci.ohio-state.edu/~plantbio/Facilities/abrc/abrchome.htm>

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Description: The mission of the Arabidopsis Biological Resource Center (ABRC) is to acquire, preserve and distribute seed and DNA resources that are useful to the Arabidopsis research community. More than 100,000 stocks are shipped annually to researchers in more than 60 countries, and modest fees for stocks are charged. The ABRC database functions and ordering system are incorporated into The Arabidopsis Information Resource (TAIR). Researchers can obtain information about Arabidopsis, perform stock searches, order stocks, and view current and past orders. Sponsors: ABRC is supported by the National Science Foundation under Grant No. 0542034.

Synonyms: ABRC

Resource Type: database, data or information resource

Keywords: arabidopsis, biological, dna, seed, FASEB list

Resource Name: Arabidopsis Biological Resource Center

Resource ID: SCR_008136

Alternate IDs: nif-0000-20954

Record Creation Time: 20220129T080245+0000

Record Last Update: 20260805T044206+0000

Ratings and Alerts

No rating or validation information has been found for Arabidopsis Biological Resource Center.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2583 mentions in open access literature.

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

Seluzicki A, et al. (2026) Molecular genetic characterization of CASEIN KINASE 1-LIKE 12 in Arabidopsis. Plant signaling & behavior, 21(1), 2610025.

Liu RX, et al. (2026) MdHMGB15-MdXERICO-MdNRP module mediates salt tolerance of apple by regulating the expression of salt stress-related genes. Journal of advanced research, 79, 61.

Kerstens M, et al. (2026) PLETHORAs shape Arabidopsis phyllotaxis through modulation of patterning robustness and accelerated inflorescence development. The New phytologist, 249(1), 495.

Nakamura M, et al. (2026) Chromatin-associated DEK proteins maintain H3K27me3 balance and coordinate developmental transitions in plants. The New phytologist, 249(2), 930.

Bai P, et al. (2025) Comparative Proteomic Profiling of Receptor Kinase Signaling Reveals Key Trafficking Components Enforcing Plant Stomatal Development. bioRxiv : the preprint server for biology.

Rusnak B, et al. (2025) Brassinosteroids mediate proper coordination of sepal elongation. bioRxiv : the preprint server for biology.

Cao Y, et al. (2025) Polyadenylation Complex CFII Recognizes Downstream Cis-element for Pre-mRNA Polyadenylation Through Interaction with an RNA-Binding Protein in Arabidopsis. Advanced science (Weinheim, Baden-Wurttemberg, Germany), 12(41), e04562.

He R, et al. (2025) Turnip mosaic virus selectively subverts a PR-5 thaumatin-like, plasmodesmal protein to promote viral infection. The New phytologist, 245(1), 299.

Grubb LE, et al. (2025) Quantitative Proteomic Analysis of Brassica Napus Reveals Intersections Between Nutrient Deficiency Responses. Plant, cell & environment, 48(2), 1409.

Tangpranomkorn S, et al. (2025) A land plant-specific VPS13 mediates polarized vesicle trafficking in germinating pollen. The New phytologist, 245(3), 1072.

Vo HM, et al. (2025) Dehydration rapidly induces expression of NCED genes from a single subclade in diverse eudicots. Planta, 261(2), 46.

Guo Y, et al. (2025) TaWI12 may be involved in pistillody and leaf cracking in wheat. *BMC plant biology*, 25(1), 123.

Park SH, et al. (2025) Trichostatin A promotes de novo shoot regeneration from *Arabidopsis* root explants via a cytokinin related pathway. *Scientific reports*, 15(1), 978.

Pang L, et al. (2025) The adaptor protein AP-3?? disassembles heat-induced stress granules via 19S regulatory particle in *Arabidopsis*. *Nature communications*, 16(1), 2039.

Yamaguchi M, et al. (2025) LIPID RICH 1 modulates allocation of carbon between starch and triacylglycerol in *Arabidopsis* leaves. *Journal of experimental botany*, 76(8), 2144.

Candela-Ferre J, et al. (2025) Plant BCL-DOMAIN HOMOLOG proteins play a conserved role in SWI/SNF complex stability. *Proceedings of the National Academy of Sciences of the United States of America*, 122(3), e2413346122.

Mou W, et al. (2025) Ethylene-independent modulation of root development by ACC via downregulation of WOX5 and group I CLE peptide expression. *Proceedings of the National Academy of Sciences of the United States of America*, 122(6), e2417735122.

Wang C, et al. (2025) Plant ubiquitin E2 enzymes UBC32, UBC33, and UBC34 are involved in ERAD and function in host stress tolerance. *BMC plant biology*, 25(1), 412.

Yang Q, et al. (2025) Comparative single-nucleus RNA-seq analysis revealed localized and cell type-specific pathways governing root-microbiome interactions. *Nature communications*, 16(1), 3169.

Liang X, et al. (2025) An intracellular CPK-ECA1 phosphoregulatory circuit couples calcium signatures to ABA homeostasis for plant osmosensitivity. *Science advances*, 11(40), eadz2428.