

Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 2026

GABI-KAT

RRID:SCR_012751

Type: Tool

Proper Citation

GABI-KAT (RRID:SCR_012751)

Resource Information

URL: <http://www.GABI-Kat.de>

Proper Citation: GABI-KAT (RRID:SCR_012751)

Description: GABI-Kat is a database of flanking sequence tags (FSTs) from T-DNA mutagenised *A. thaliana* plants. Over time, an increasing number of lines will become available from NASC. The "show sequence" page of SimpleSearch will display if a GABI-Kat line for a given FST has already been donated to NASC. Lines that have so far not been regrown and confirmed are only available from GABI-Kat directly. We have used four vectors: pAC106 (GenBank:AJ537513), pAC161 (GenBank:AJ537514), pGABI1 (GenBank:AY529716) and pADIS1 (GenBank:AY529717). Sequence and overview map data of all vectors are available from the download page. Features of interest which are not included in the map should be deduced from the sequence. For a specified line, the vector is displayed in the "Show Sequence" page of SimpleSearch.

Synonyms: GABI-KAT

Resource Type: data or information resource, database

Keywords: *a. thaliana*, *a. thaliana* genome, mutagen, plant genome, FASEB list

Resource Name: GABI-KAT

Resource ID: SCR_012751

Alternate IDs: nif-0000-02865

Record Creation Time: 20220129T080312+0000

Record Last Update: 20260729T044310+0000

Ratings and Alerts

No rating or validation information has been found for GABI-KAT.

No alerts have been found for GABI-KAT.

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 [nidm-terms](#) .

Zhang Y, et al. (2025) ABA importers ABCG17 and ABCG18 redundantly regulate seed size in Arabidopsis. The Plant journal : for cell and molecular biology, 121(6), e70096.

Correa Córdoba S, et al. (2025) A data-integrative modeling approach accurately characterizes the effects of mutations on Arabidopsis lipid metabolism. Plant physiology, 197(2).

Lemke MD, et al. (2024) Investigating the mechanism of chloroplast singlet oxygen signaling in the Arabidopsis thaliana accelerated cell death 2 mutant. Plant signaling & behavior, 19(1), 2347783.

Delplace F, et al. (2024) The receptor MIK2 interacts with the kinase RKS1 to control quantitative disease resistance to Xanthomonas campestris. Plant physiology, 197(1).

Islam MM, et al. (2024) Involvement of Peptidoglycan Receptor Proteins in Mediating the Growth-Promoting Effects of Bacillus pumilus TUAT1 in Arabidopsis thaliana. Plant & cell physiology, 65(5), 748.

Li N, et al. (2024) Overexpression of vacuolar H⁺-pyrophosphatase from a recretohalophyte Reaumuria trigyna enhances vegetative growth and salt tolerance in transgenic Arabidopsis thaliana. Frontiers in plant science, 15, 1435799.

Cho Y, et al. (2024) Arabidopsis AGB1 participates in salinity response through bZIP17-mediated unfolded protein response. BMC plant biology, 24(1), 586.

Yu Y, et al. (2023) Arabidopsis thaliana: a powerful model organism to explore histone modifications and their upstream regulations. Epigenetics, 18(1), 2211362.

Traeger J, et al. (2023) Super-Resolution Imaging of Plant Receptor-Like Kinases Uncovers Their Colocalization and Coordination with Nanometer Resolution. Membranes, 13(2).

von Bismarck T, et al. (2023) Growth in fluctuating light buffers plants against photorespiratory perturbations. Nature communications, 14(1), 7052.

Luengwilai K, et al. (2022) Ectopic Expression of *Arabidopsis thaliana* zDof1.3 in Tomato (*Solanum lycopersicum* L.) Is Associated with Improved Greenhouse Productivity and Enhanced Carbon and Nitrogen Use. *International journal of molecular sciences*, 23(19).

Ito T, et al. (2022) Glutathione degradation activity of γ -glutamyl peptidase 1 manifests its dual roles in primary and secondary sulfur metabolism in *Arabidopsis*. *The Plant journal : for cell and molecular biology*, 111(6), 1626.

Lang L, et al. (2021) The DREAM complex represses growth in response to DNA damage in *Arabidopsis*. *Life science alliance*, 4(12).

Zhang Y, et al. (2021) ABA homeostasis and long-distance translocation are redundantly regulated by ABCG ABA importers. *Science advances*, 7(43), eabf6069.

Serrazina S, et al. (2021) Expression of *Castanea crenata* Allene Oxide Synthase in *Arabidopsis* Improves the Defense to *Phytophthora cinnamomi*. *Frontiers in plant science*, 12, 628697.

Mahesh S, et al. (2021) Functional analysis of a conserved domain in SWITCH1 reveals a role in commitment to female meiocyte differentiation in *Arabidopsis*. *Biochemical and biophysical research communications*, 551, 121.

Perry N, et al. (2021) RUS6, a DUF647-containing protein, is essential for early embryonic development in *Arabidopsis thaliana*. *BMC plant biology*, 21(1), 232.

Kamiyama Y, et al. (2021) *Arabidopsis* group C Raf-like protein kinases negatively regulate abscisic acid signaling and are direct substrates of SnRK2. *Proceedings of the National Academy of Sciences of the United States of America*, 118(30).

Ambastha V, et al. (2021) RabA2b Overexpression Alters the Plasma-Membrane Proteome and Improves Drought Tolerance in *Arabidopsis*. *Frontiers in plant science*, 12, 738694.

Liu S, et al. (2021) The ARRE RING-Type E3 Ubiquitin Ligase Negatively Regulates Cuticular Wax Biosynthesis in *Arabidopsis thaliana* by Controlling ECERIFERUM1 and ECERIFERUM3 Protein Levels. *Frontiers in plant science*, 12, 752309.