

Resource Summary Report

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Leibniz Institute of Plant Genetics and Crop Plant Research; Saxony-Anhalt; Germany

RRID:SCR_011341

Type: Tool

Proper Citation

Leibniz Institute of Plant Genetics and Crop Plant Research; Saxony-Anhalt; Germany
(RRID:SCR_011341)

Resource Information

URL: <http://www.ipk-gatersleben.de/en/>

Proper Citation: Leibniz Institute of Plant Genetics and Crop Plant Research; Saxony-Anhalt; Germany (RRID:SCR_011341)

Abbreviations: IPK

Synonyms: Leibniz-Institut für Pflanzengenetik und Kulturpflanzenforschung, Leibniz-Institut für Pflanzengenetik und Kulturpflanzenforschung (IPK), Leibniz-Institut für Pflanzengenetik und Kulturpflanzenforschung (IPK), Leibniz-Institut für Pflanzengenetik und Kulturpflanzenforschung, Leibniz Institute of Plant Genetics and Crop Plant Research, IPK Gatersleben

Resource Type: institution

Funding:

Resource Name: Leibniz Institute of Plant Genetics and Crop Plant Research; Saxony-Anhalt; Germany

Resource ID: SCR_011341

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Record Last Update: 20250420T014534+0000

Ratings and Alerts

No rating or validation information has been found for Leibniz Institute of Plant Genetics and Crop Plant Research; Saxony-Anhalt; Germany.

No alerts have been found for Leibniz Institute of Plant Genetics and Crop Plant Research; Saxony-Anhalt; Germany.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Junker A, et al. (2014) Optimizing experimental procedures for quantitative evaluation of crop plant performance in high throughput phenotyping systems. *Frontiers in plant science*, 5, 770.

Galili G, et al. (2014) The role of photosynthesis and amino acid metabolism in the energy status during seed development. *Frontiers in plant science*, 5, 447.

Kumar SP, et al. (2007) Comparative analysis of the 100 kb region containing the Pi-k(h) locus between indica and japonica rice lines. *Genomics, proteomics & bioinformatics*, 5(1), 35.