Resource Summary Report

Generated by nidm-terms on Jul 28, 2026

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 fur Pflanzengenetik und Kulturpflanzenforschung, Leibniz-Institut fur 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

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

Resource ID: SCR_011341

Record Creation Time: 20220129T080304+0000

Record Last Update: 20260725T190714+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 nidm-terms .

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.