

Resource Summary Report

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INSDC

RRID:SCR_011967

Type: Tool

Proper Citation

INSDC (RRID:SCR_011967)

Resource Information

URL: <http://www.insdc.org/>

Proper Citation: INSDC (RRID:SCR_011967)

Description: International collaboration of the International Nucleotide Sequence Databases (INSD), DDBJ, ENA, and GenBank, maintained for over 18 years. Individuals submitting data to the international sequence databases should be aware of INSDC policy.

Abbreviations: INSDC

Synonyms: International Nucleotide Sequence Database Collaboration

Resource Type: community building portal, data or information resource, portal

Keywords: nucleotide sequence, FASEB list

Resource Name: INSDC

Resource ID: SCR_011967

Alternate IDs: OMICS_01653

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260821T193921+0000

Ratings and Alerts

No rating or validation information has been found for INSDC.

No alerts have been found for INSDC.

Data and Source Information

Usage and Citation Metrics

We found 240 mentions in open access literature.

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

Ahrens AK, et al. (2026) Germany as a key transit hub for the emergence and spread of high pathogenicity avian influenza H5 clade 2.3.4.4b reassortants in Europe. *Frontiers in microbiology*, 17, 1824729.

Tello-Ruiz MK, et al. (2026) Adoption of Standard Reference SNP Identifiers in Agricultural Genomics for Interoperability and Data Reuse. *Scientific data*, 13(1).

Suetsugu K, et al. (2026) Divergent mycorrhizal strategies and nutritional modes in the East Asian orchid genus *Ephippianthus*. *Mycorrhiza*, 36(4).

Yamato M, et al. (2026) Partial mycoheterotrophy in the arbuscular mycorrhizal *Gentiana squarrosa* (Gentianaceae) demonstrated by coculture assays using C3 and C4 plants. *Mycorrhiza*, 36(3).

Saghaei S, et al. (2026) VirJenDB: a FAIR (meta)data and bioinformatics platform for all viruses. *Nucleic acids research*, 54(D1), D912.

Beldade P, et al. (2026) LepEU: A Consortium to Study the Population Genomics of Butterflies and Moths Across Europe. *Evolutionary applications*, 19(5), e70246.

Sato R, et al. (2026) Dragonfly red opsins share a common tuning mechanism with mammalian red opsins and further enhancement of near-infrared sensitivity. *Cellular and molecular life sciences : CMLS*, 83(1), 66.

Wu S, et al. (2026) Environmental DNA (eDNA) Technology in Biodiversity and Ecosystem Health Research: Advances and Prospects. *Ecology and evolution*, 16(1), e72891.

Damgaard F, et al. (2026) Distinct prophage infections in colorectal cancer-associated *Bacteroides fragilis*. *Communications medicine*, 6(1).

Le Floch E, et al. (2026) Wheat historical phenotypic data from European genebanks as an important resource for research and breeding. *Scientific data*, 13(1).

Hanna SJ, et al. (2025) The Type 1 Diabetes T Cell Receptor and B Cell Receptor Repository in the AIRR Data Commons: a practical guide for access, use and contributions through the Type 1 Diabetes AIRR Consortium. *Diabetologia*, 68(1), 186.

Cannon EKS, et al. (2025) Guidelines for gene and genome assembly nomenclature. *Genetics*, 229(3).

Fukasawa Y, et al. (2025) Oak Wilt Disease May Reduce the Initial Decay Rate of Dead *Quercus serrata* Stems by Altering Fungal Communities in the Wood. *Environmental microbiology*, 27(1), e70026.

Fischer L, et al. (2025) Recombinant Myxoma Virus in European Brown Hares, 2023-2024. *Emerging infectious diseases*, 31(8), 1608.

Furuhashi R, et al. (2025) ?*Synodus autumnus*, a new species of lizardfish (Aulopiformes, Synodontidae) from the Indo-Pacific region, and a reassessment of distributional records of *Synodus rubromarmoratus*. *ZooKeys*, 1243, 191.

Gutiérrez-Hurtado IA, et al. (2025) Past, Present and Future Perspectives of Forensic Genetics. *Biomolecules*, 15(5).

Vancaester E, et al. (2025) Ghosts of symbionts past: the hidden history of the dynamic association between filarial nematodes and their *Wolbachia* endosymbionts. *G3* (Bethesda, Md.), 15(12).

Dyer SC, et al. (2025) Ensembl 2025. *Nucleic acids research*, 53(D1), D948.

Righi E, et al. (2025) The BioGenome Portal: a web-based platform for biodiversity genomics data management. *NAR genomics and bioinformatics*, 7(1), lqaf020.

Suetsugu K, et al. (2025) Mycoheterotrophic continuum in rhizoctonia associations: genetic divergence and carbon acquisition variation among *Odontochilus* orchids. *Annals of botany*, 136(2), 407.