

# Resource Summary Report

Generated by [dkNET](#) on Aug 27, 2026

## Oryza Tag Line

RRID:SCR\_013254

Type: Tool

### Proper Citation

Oryza Tag Line (RRID:SCR\_013254)

### Resource Information

**URL:** <http://urgi.versailles.inra.fr/OryzaTagLine/>

**Proper Citation:** Oryza Tag Line (RRID:SCR\_013254)

**Description:** Oryza Tag Line consists in a searchable database developed under the Oracle management system integrating phenotypic data resulting from the evaluation of the G??noplante rice insertion line library. To display the sequence information (Flanking Sequence Tags) resulting from the molecular characterization of the mutagen insertion sites in the collection, Oryza Tag Line is linked to OryGenesDB a GGB-derived interface or FLAGDB++ a Java based interface. Aside from generic data ( production record and seed stock ) Oryza Tag Line includes textual and pictorial information resulting from the following observations: (i.) GUS/GFP expression assays conducted on the primary transformants (T0) and/or T1 progenies (ii.) morpho-physiological alterations detected during the growth of the T1 progenies either in phytotron -up to the 5-6 leaf stage- or under agronomical conditions in the field (collaboration with CIAT, Cali, Colombia).

**Synonyms:** Oryza Tag Line

**Resource Type:** data or information resource, database

**Resource Name:** Oryza Tag Line

**Resource ID:** SCR\_013254

**Alternate IDs:** nif-0000-03232

**Record Creation Time:** 20220129T080315+0000

**Record Last Update:** 20260821T194006+0000

### Ratings and Alerts

No rating or validation information has been found for Oryza Tag Line.

No alerts have been found for Oryza Tag Line.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Sozzani R, et al. (2011) High-throughput phenotyping of multicellular organisms: finding the link between genotype and phenotype. Genome biology, 12(3), 219.