

Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 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: 20260729T044314+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.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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