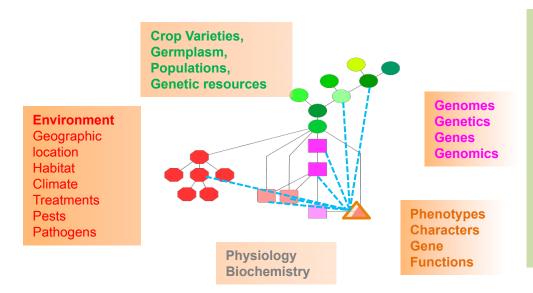
Planteome Overview

Common Reference Ontologies for Plants (cROP) and Planteome Tools for Integrative Plant Genomics

- Centralized platform where **reference ontologies for plants** will be used to access plant genomics data resources across a wide range of plant species
- Online informatics portal for **ontology-based**, **annotated data** for plant germplasm, gene expression, and non-model genomes
- Smart and semantic data query, analysis, visualization and community-based annotation and curation tools



- Plant Ontology (PO)
- Plant Trait Ontology (TO)
- Plant Stress Ontology (PSO)
- Plant Experimental Conditions
 Ontology (PECO/EO)
- Gene Ontology (plants)
- Phenotypic Qualities Ontology (PATO)
- Cell Type Ontology (CL)
- Chemicals (ChEBI)
- Protein Ontology (PRO)



Progress and Success Stories



Plant Ontology

www.plantontology.org

- A highly successful, collaborative, publicly-available resource with more than 50 collaborating partners
- Links plant genomics data sets to terms describing plant anatomy and developmental stages across all plant species for comparative studies and classroom teaching
- Currently over 1200 terms linked to more than 2.2 million data points

from 23 plant species Relations -is a [] - part_of ←has part develops_from ← L - located in ☐ 0 PO:0009011 : plant structure [110909] ■ PO:0025498 : cardinal part of multi-tissue plant structure [47530] O PO:0025497 : collective plant structure [103588] ■ PO:0025099 : embryo plant structure [41230] ⊕ PO:0000004 : in vitro plant structure [19145] n PO:0025496 : multi-tissue plant structure [103668 ⊕ PO:0009002 : plant cell [104207] D PO:0009072 : plant ovary [374] PO:0009007 : portion of plant tissue [98346] ■ PO:0030078 : rhizoid [0] ⊕ PO:0000282 : trichome [372] ■ PO:0000003: whole plant [104955]



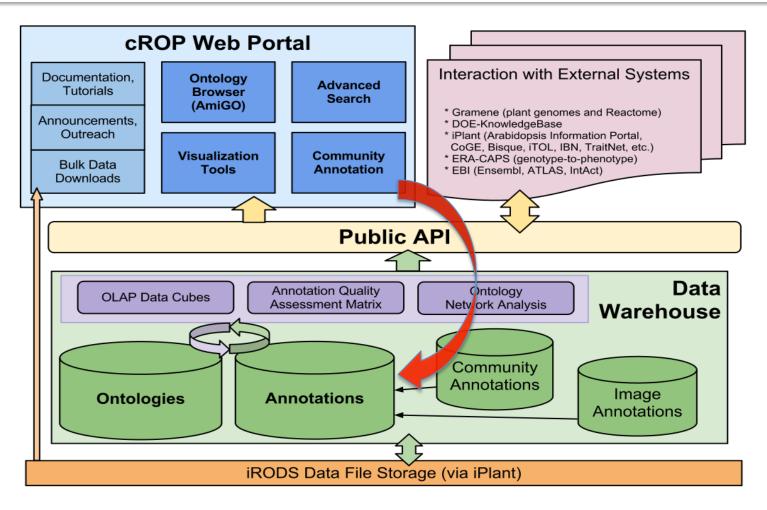
Cooper L et al. Plant Cell Physiol 2013;54:e1

Arnaud E, Cooper L, et al. (2012) Towards a Reference Plant Trait Ontology for Modeling Knowledge of Plant Traits and Phenotypes. PKEOD. Barcelona, Spain, pp 220–225.



Data Acquisition and Management





The Planteome informatics portal and the data store organization

Structure of the Planteome Project: Data Generation, Access and Sharing- Open and Collaborative

Planteome

iPlant Hosting platform

Expansion & Maintenance of the Reference Ontologies for Plants and US Outreach

- Coordinate the overall project and lead Ontology Development
- Use case for the semantic integration framework
- Develop data warehouse, visualization and analysis tools
- Manage online resource at iPlant

Dennis Stevenson, *Co-PI*New York Botanic

Garden.

Chris Mungall, Co-PI
Gene Ontology

Georgios Gkoutos, Co-PI PATO, Aberystwyth, UK

Barry SmithOBO-Foundry & NCBO

Pankaj Jaiswal, Pl **Laurel Cooper**,

Coordinator

Justin Elser, Sys. Admin Oregon State University

NSF

Gramene, KBase, SGN, MaizeGDB, AIP, UniProt, SoyBase, Oryzabase and many others...

EBI Paul Kersey

Ruth Bastow- GARnet Chris Rawlings-Rothamsted Research UK, BBSRC

Elizabeth Arnaud, Co-PI

Crop Ontology
CGIAR Consortium

Crop Ontology and the Integrated Breeding Platform

- Development of crop-specific ontologies to describe traits and annotations of
- wheat, rice, maize and other crops: cassava, sorghum, potato, yam and others

Divseek

- Contribution to the semantic framework
- Community Engagement, Breeding for Development

Phenotype RCN

John Doonan, National Plant Phenomics Center Aberystwyth Univ. Fabio Florani - Julich Bjorn Usadel – Aacehn Uli Schurr – Julich DROPS Germany

Cyril Pommier –BAP; Ephesis
Jacques Legouis – GEDEC,
Breedwheat

Francois Tardieu- Phenome, DROPS France, INRA Luke Ramsay
Dave Marshall
J. Hutton

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Semantic integration framework

- Develop Trait Ontology for wheat, barley & Brassicaceae
- Environment Ontology from Phenotyping platforms
- Warehouse, mirroring on server -EBI
- Community engagement

iDIV



Planteome and DivSeek



Many points of coincidence between Planteome and DivSeek:

- Developing common standards and descriptors that allow data to be structured, shared and reconciled with existing data management schemes and that facilitate use by diverse scientific, crop and breeding communities.- Phase 1 Case Studies and Pilot Projects
- 2. Establishing **new tools** and **approaches** that enable data exploration, mining and utilization
- 3. Characterize crop diversity using state-of-the-art genomic, phenomic and molecular technologies
- 4. Promote awareness of the value of crop diversity and mobilize the diverse resources required to accomplish these goals.