

Génétique Quantitative et Évolution - Le Moulon

Director:

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Scientific goals

The main scientific focus of the UMR⁵, so-called « Le Moulon », is the genetics and evolution of complex traits (quantitative traits), with three complementary objectives:

- *Bringing knowledge in fundamental multifactorial genetics, within the following subjects: modeling of quantitative variation, evolution of quantitative traits, searches for genes underlying quantitative effects and the dynamics of genetic resources.*
- *Guiding selection methodologies by including genomic knowledge. Exploiting and enhancing genetic resources. Target traits are typically growth, development or adaptation traits.*
- *Decrypting the genetic, molecular and epigenetic mechanisms controlling both qualitative and quantitative traits, unveiling their importance in evolutionary history of species and associated adaptations.*

The main species studied are maize and wheat. Some projects involve other species, either as models (*Arabidopsis thaliana*, *Saccharomyces cerevisiae*, *Escherichia coli*), or to address specific questions (*Brassica*, basal Eudicots).

Context

The UMR is linked to INRA, to Université Paris-Sud, to CNRS and to AgroParisTech. It gathers about fifteen full time researchers (INRA and CNRS), a dozen faculty members (Université Paris-Sud and AgroParisTech) and a permanent staff of nearly thirty engineers and technicians; the lab hosts about thirty trainees or postdoctoral and doctoral scientists.

Le Moulon is one of the three constitutive laboratories of the institute « Diversité, Écologie et Évolution du Vivant » (IDEEV) associated with Université Paris-Sud and CNRS.

We participate in several organizations *d'excellence*⁶. The UMR is a member of two LabEx networks "Biodiversity, Agroecosystems Society, Climate" (BASC) and "Saclay Plant Sciences" (SPS). Le Moulon coordinates the project « AMAIZING » that rallies twenty four public and private partners: this project is funded by the ANR (National Funding Agency).

¹ DR : Directeur de Recherche = Research PI

² INRA: Institut National de Recherche Agronomique

³ AgroParisTech: Graduate Institute in Life and Environmental Science and Engineering

⁴ CNRS: Centre National de la Recherche Scientifique

⁵ UMR: Unité Mixte de Recherche = Research Unit

⁶ Laboratoire d'excellence (Labex) and Initiative d'excellence (Idex) = two national networks programs.

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Research Teams

- *Biologie et Adaptation des Systèmes en Évolution (BASE)* – J. LEGRAND, UNIVERSITÉ PARIS-SUD
Modeling of quantitative variation and its evolution by integrating genetic, molecular, genomic, metabolic and environmental levels.
- *Recombinaison des Allèles en Méiose : Déterminisme, Applications, Modélisation (RAMDAM)* – O. MARTIN, INRA
Study of the formation of meiotic crossovers by theoretical and experimental approaches. Integrative biology and modeling.
- *Structure et Évolution des Chromosomes Fongiques (SECF)* – C. FAIRHEAD, UNIVERSITÉ PARIS-SUD
Modes of reproduction of hemiascomycetes yeasts, role in the evolution history of *Candida glabrata*. Population genomics, comparative and functional genomics, and adaptation of yeast to their environment.
- *Dynamique du Génome et Adaptation des Plantes cultivées (DyGAP)* – M. TENAILLON, CNRS
Characterisation of nucleotidic, structural and epigenetic variations implicated in the evolution of genomes, the regulation of their expression and the adaptation of populations.
- *Génétique, Épigenétique et Évolution de la Morphogenèse Florale (GE2MorF)* – C. DAMERVAL, CNRS
Mechanisms driving morphological diversity of petals and perianth symmetry. Role of perianth diversity for adaptation and species diversification.
- *Diversité, Évolution et Adaptation des Populations (DEAP)* – J. ENJALBERT, INRA
Genetic bases of local adaptation; dynamic biodiversity management and uses for innovative and sustainable farming, participatory plant breeding.
- *Génétique Quantitative et Méthodologie de la Sélection (GQMS)* – A. CHARCOSSET, INRA
Study of the genetic bases of complex traits and mechanisms of response to selection, optimization of genetic resource management and of selection methodology.

Facilities

- *Atelier Cartographie, Expression et Polymorphisme* - M. FALQUE, INRA
Common projects in molecular biology, technological watch, methodological developments for specific projects.
- *Atelier de Bioinformatique et d'Informatique* – 3 groups
ABI-AnaSeq, J. JOETS, INRA (Analysis of genomes and sequences); *ABI-Soft*, D. STEINBACH, INRA (Development of programs and data-bases for genetics and genomics); *ABI-Sys*, O. LANGELLA, CNRS (Computing services).
- *Plate-forme de protéomique PAPPISO* - M. ZIVY, CNRS
Identification and quantification of proteins, analysis of post-translational modifications, methodological developments, bioinformatics for proteomics.
- *Installation Expérimentale* – P. BRABANT, AgroParisTech
Field experiments for the genetic research of the laboratory, agronomic experiments for the INRA networks working on maize and wheat, hosting of educational experiments for the University Paris-Sud and AgroParisTech.

