

THE MOST POWERFUL TOMATO GENOTYPING PLATFORM

Plant Biotech Division Beyond Seeds Biotech Group

GENOTYPING

CURRENT SOLUTIONS FOR PLANT BREEDING PROGRAMS

Services based on molecular low number of markers

Specific services on a case-by-case basis, not comprehensive

The panels of characteristics of agronomic interests are limited TRADITIONAL GENOTYPING PROCESS IN TOMATO

The information they generate is not enough to design crossbreeding panels based on genetic distance Impossible to obtain quality information on the genetic background of the varieties

Cannot be multiplexed

ÓRIEL

MASSIVE TOMATE GENOTYPING PLATFORM

It allows to find the combination of desired agronomic characters with high reliability in the seedling stage

Multiplexing is possible. Analysis of large volumes of plants, reducing the number of individuals to be evaluated in the field

Robustness in genetic fingerprint analysis thanks to the high number of sequenced markers



Combined evaluation of resistance, characters with agronomic interest, varietal purity and genetic distance in a single analysis The genetic background of a variety can be analyzed with high resolution

Crossbreeding planning based on their genetic distance

WHAT IS ÓRIEL

Molecular markers platform designed to **facilitate** the processes of **selection and plant breeding in tomato**

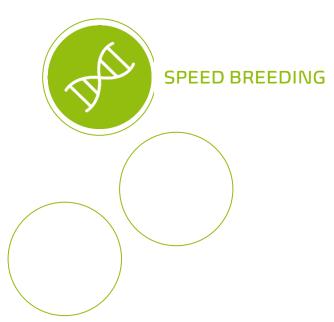
100 PHENOTYPE SPECIFIC

500 SNP

400 GENETIC BACKGROUND



WHAT SOLUTIONS ÓRIEL OFFERS



VARIETAL PURITY ANALYSIS

Using up to 500 SNP molecular markers, we obtain information on the level of purity

MOLECULAR PHENOTYPING

Prediction of the phenotype from the interpretation of the markers thanks to the development of 100 specific markers

PREDICTION OF BEST HYBRIDS

From the calculation of the genetic distance, a list of recommended crosses between plants is obtained, guaranteeing the maximum hybrid vigor in the offspring





WHAT SOLUTIONS ÓRIEL OFFERS

In addition to the standard report, our team of geneticists can issue a complete report and a detailed analysis of the results.

A population study is also included, calculating a matrix of genetic distances and a phylogenetic tree of the accessions o know the materials in a deep way.

FULL INTEGRATION WITH NOAH® PLANT BREEDING

ÓRIEL integrates with the leading plant germplasm management and plant breeding tool, NOAH.

Consult and visualize the data directly on your computer while you make decisions in your plant breeding programs.



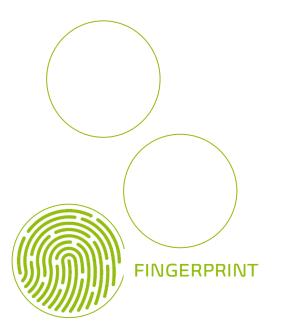
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ANALYTICS

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WHAT SOLUTIONS ÓRIEL OFFERS



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Obtaining a complete genetic fingerprint of each of the genotyped materials

Variety registration

Seed analysis





PHYSIOLOGICAL CHARACTERS

Chocolate color fruit	Accumulation of chlorophylls in the fruit skin
Purple color fruit	Accumulation of anthocyanins in the fruit skin
	Rounded
Fruit shape	Pear-shaped
	Determinate
Growth habit	Indeterminate
	Dwarf
Plant size	Normal

RESISTANCE CHARACTERS

TYLCV	Fusarium (Fol:1)
TSWV	Fusarium (Fol:2)
ТоМV	Verticillium (Vd:1; Va:1)
Pseudomonas syringae (Pst)	Stemphyllium spp.
Fusarium Crown Rot (FCR)	Nematodes (Ma, Mi, Mj)





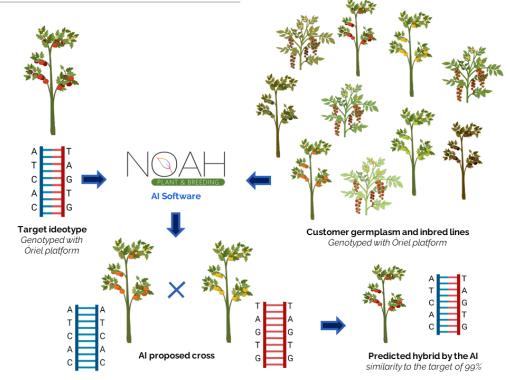
ARTIFICIAL INTELLIGENCE TO DESIGN CROSSBREEDING

REVERSE BREEDING 2.0

ÓRIEL

From a database of genotyped potential parents and knowing the genetic profile of the target ideotype, we can obtain a list of proposed crosses to reconstitute this ideotype from a genetic point of view.

Our AI platform uses the genotyping and phenotype data of certain characteristics of each tomato to be able to make elite tomato predictions, based on the comparison of these simulated genetic combinations.







"If you want to speed up the time and precision in the selection of your tomato breeding programs, ÓRIEL is the definitive tool".

Resistant to nematodes







For further information visit www.seeds4i.com res l

Determinate growth habit

Susceptible to Dseudomonas Agronomic Engineering · Software · Plant Biotech · Nanotechnology · Microbiology · Biopharma



Beyond Seeds

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