

TOWARD A SUSTAINABLE VITICULTURE IMPROVED GRAPEVINE PRODUCTIVITY AND TOLERANCE TO ABIOTIC AND BIOTIC STRESSES BY COMBINING RESISTANT CULTIVARS AND BENEFICIAL MICROORGANISMS

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sustainable viticulture: **Toward Improved** grapevine productivity and tolerance to abiotic and biotic stresses by combining resistant cultivars and beneficial microorganisms

Aït Barka E., <u>Aziz A.</u>, Sanchez L., Trotel-Aziz P., Jacquard C., Clément C., Gaveau-Vaillant N., Tzortzakis N., Topfer R., Kicherer A., Escalona J-M., HÖFTE M. REY P., Gardiman M., De Nardi B., Grando M.S., Fusco L., Maciejczak M., Vermunt A., Falcão Salles J., Agstner B., Taglietti F., and other members of Vitismart consortium

GOALS: to improve resistance/tolerance strategies of grapevine cultivars to pathogens and to mitigate the undesirable effect of climatic change

Outcomes:

Identification of susceptible and resistant genotypes for their tolerance/resistance to pathogens



 Identification of efficient Biocontrol agent (BCA) against mean grapevine diseases





Induced

Tolerance

Physiology, primary

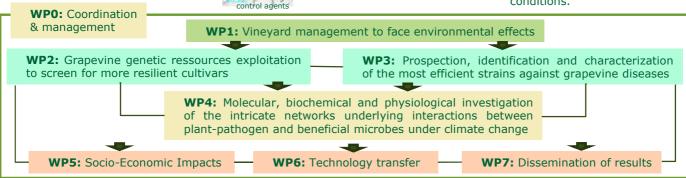
metabolism, mineral status....



Enhanced Resilience toward abiotic and bioticustouss

Establishment of the best association between bacteria and resilient cultivars in order to deliver a new generation of grapevines adapted to climate change

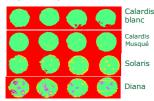
 Investigate the physiological and molecular responses of grapevine inoculated with selected beneficial bacteria, under heat and water deficit conditions.



WP 2: Resilient cultivars

Hyperspectral imaging to early detect DM

Downy mildew: Leaf disc assays with 27 different genotypes (susceptible; resistant and breeding lines). The hyperspectral tool consists of five different models to achieve best prediction performance.

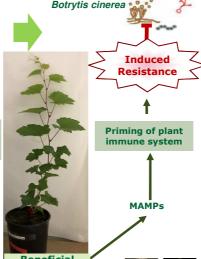


Solaris



Water shortage High temperatures

Identification of genotypes more adapted to abiotic stress



Plasmopara viticola

Beneficial microbes

WP 4: Mechanisms underlying the tripartite

interaction under climate change

Characterization of microbiome from resilient cultivars by molecular tools

WP 3: Efficient microbes



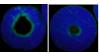
Microbial co-occurrence in the rhizosphere and endosphere using 16S rRNA based metagenomics metatranscriptomics

Identification of efficient strains

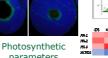
- Pseudomonas fluorescens sp. Paraburkholderia phytofirmans
- Pythium oligandrun

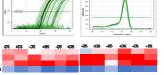




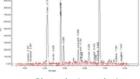


conditions





Gene expression

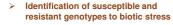






Induced systemic resistance

Establishment of the best association between bacteria and





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LDA using VNIR spectra for cultivar classification

Grey mold: Mapping of physical

barrieres against Botrytis/Drought



















