



Illumina
ACCELERATOR

PLUGANDPLAY

TERRA
by Rabobank + RocketSpace

BIOME MAKERS



Biome Makers

We are geneticists, microbiologists, software engineers, bioinformaticians, communication and sales experts, and real winemakers.

we are...

20

team
members

30

years old
average

40%

PhD level



... in US and Europe

Biome Makers: Achievements

Illumina
ACCELERATOR

World Class
Genomic Accelerator

PLUGANDPLAY

World Class
Tech Accelerator

 **ERRA**
by Rabobank + RocketSpace

Agriculture
Accelerator



Winnovation Awards 2017
(San Francisco, CA)



AgFunder 2016
(San Francisco, CA)



Finalist AgTech Latam 2017
(Chile)



PYME INNOVADORA

Pyme Innovadora 2017
(Spain)

InnoSTARS

Semi-Finalist InnoStars
(China)

 **CB**INSIGHTS

nature
biotechnology

 **CLSA**
California Life Sciences Association

Our AgTech proposal

Microbial discovery

**Welcoming
future
agriculture**



Harness the Microbiome

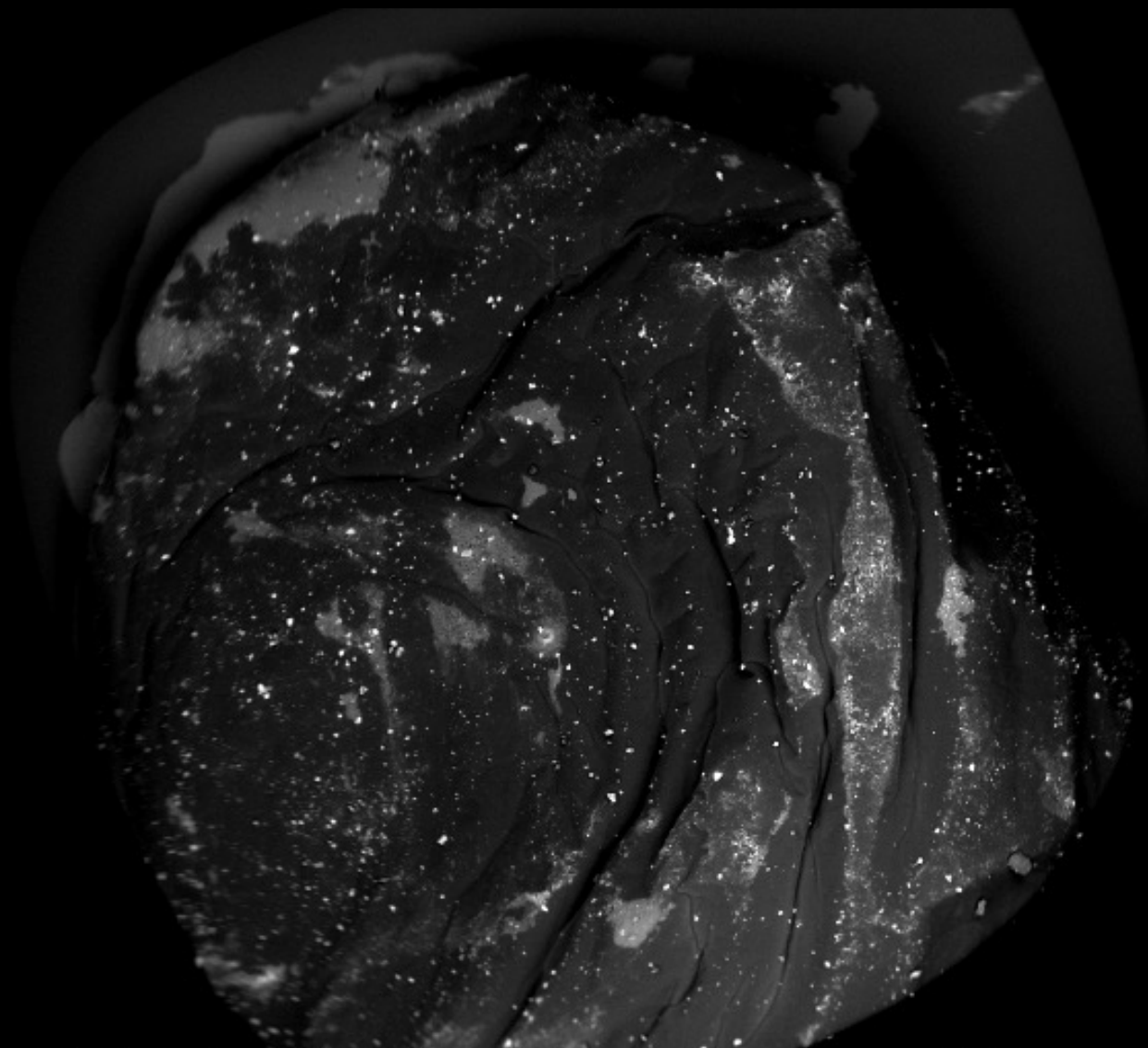
Healthy, productive, living soils improve and sustain productivity and quality



1,000,000,000 Microbes/oz

**FoodTech is already
in the wave!!**





10/10/2016 | det | spot | WD | HV | vac mode | 2 mm
12:13:23 PM | BSED | 6.0 | 13.1 mm | 19.00 kV | Low vacuum | UVA-1-12

Scanning Electron Microscopy – Grape Surface

The context


How much does it cost to solve AgProblems in the soil?

\$23.5B /year
Nutrition complements

\$15.2B /year
Chemical sprays



Even after the treatments
20-40% crop value is lost due to diseases



Technology requirements

- Accurate results**

Solution based on the microbiology of the soil-
- Customized suggestions**

Suggestions based on geo-location and metadata-
- Online portal**

Portal tools to easily explore all the information.
- Crop intelligence**

Custom algorithms for vineyard problems

accurate

Solution based on the microbiology of the soil-

Customized suggestions

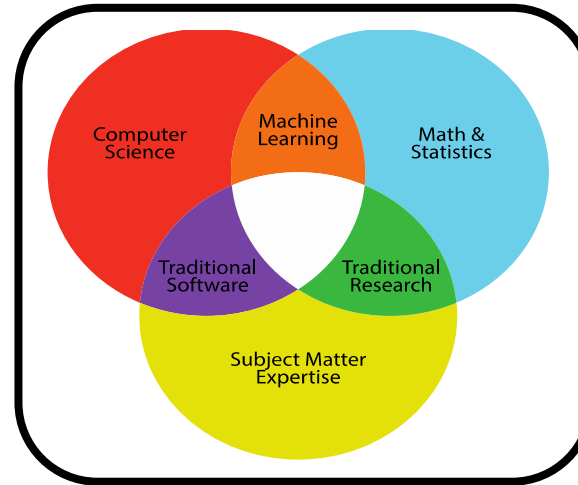
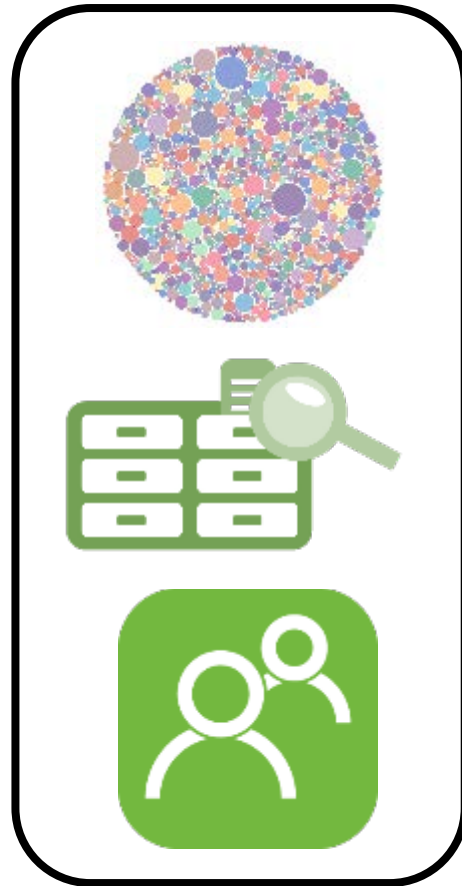
Suggestions based on geo-location and metadata-

online portal

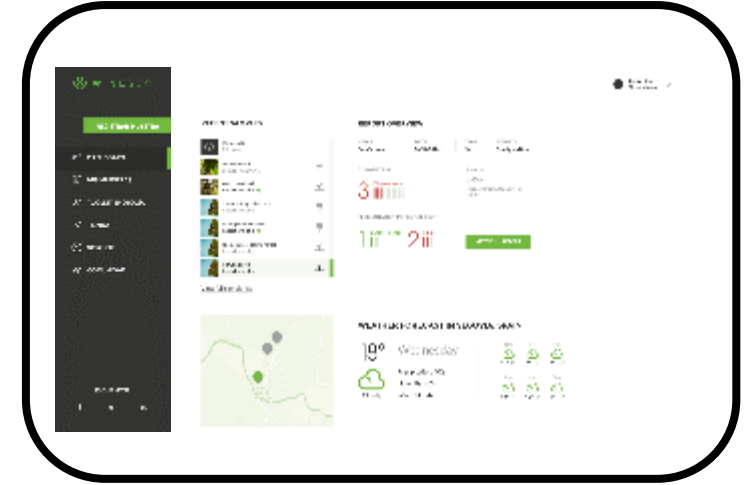
Portal tools to easily explore all the information.

op intelligence

Custom algorithms for vineyard problems



VERITAS

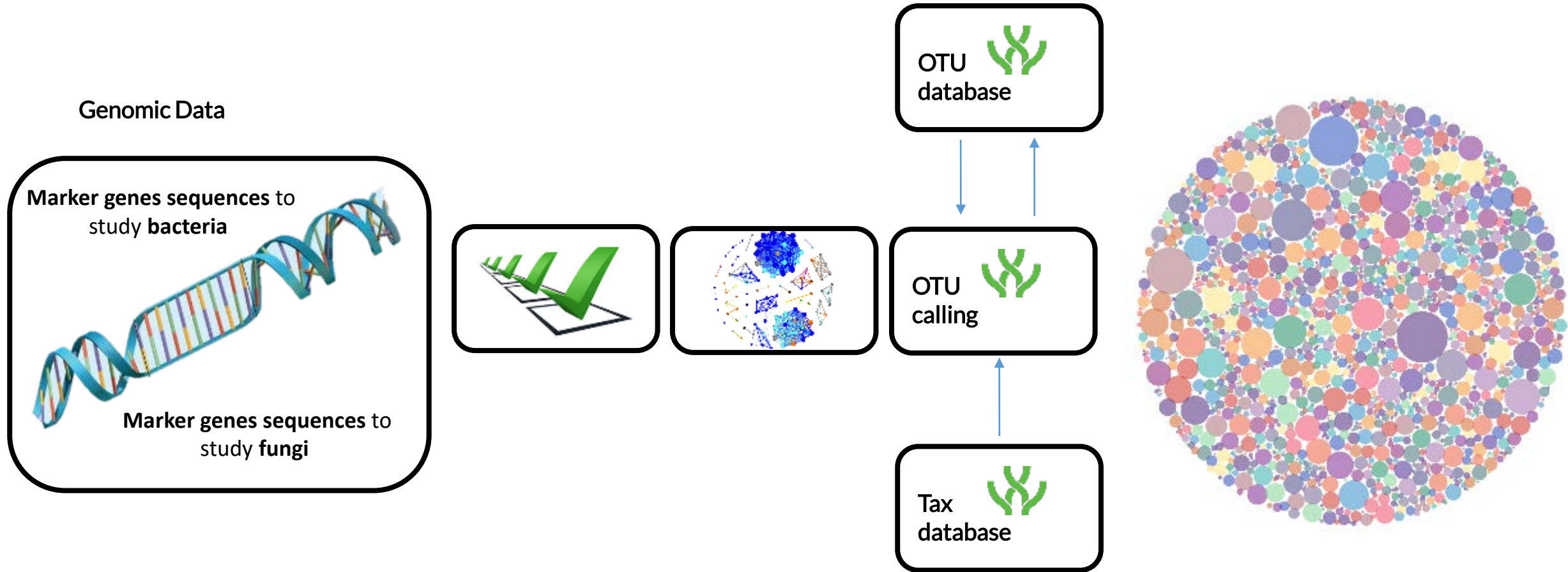


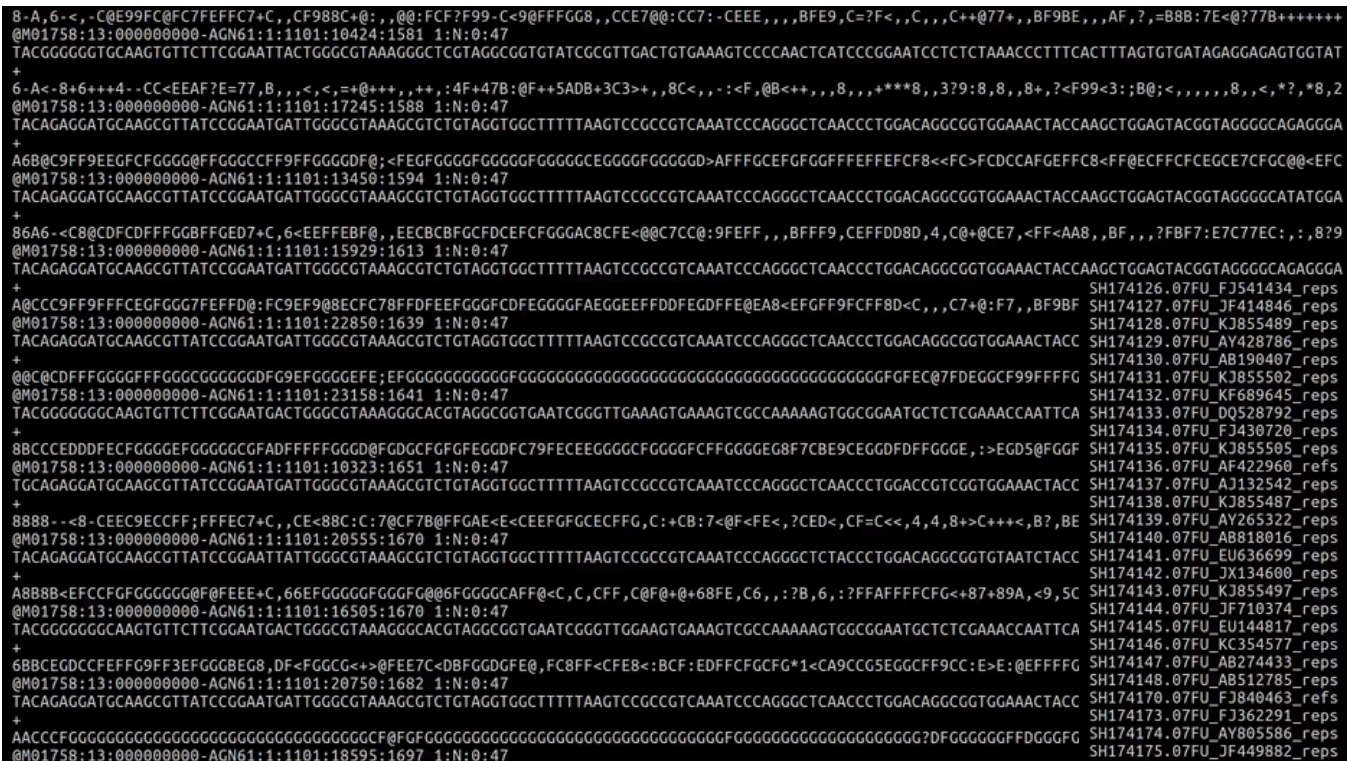
WineSeq Portal

WineSeq is a DNA based technology: microbiome discovery



E



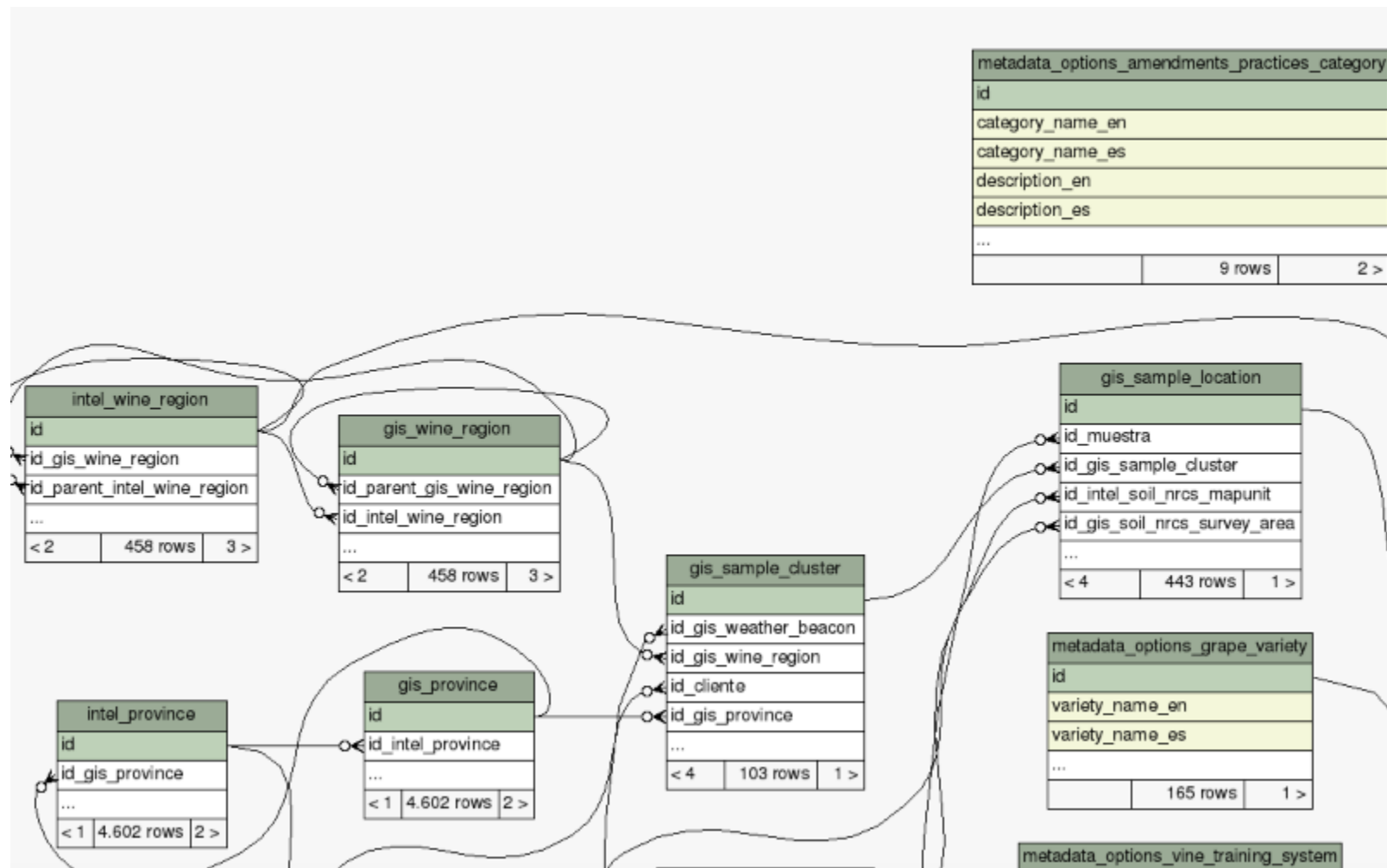


k_Fungi;p_Ascomycota;c_unidentified;o_unidentified;f_unidentified;g_unidentified;s_Ascomycota sp
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_Gaeumannomyces;s_Gaeumannomyces
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_Gaeumannomyces;s_Gaeumannomyces
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_Harpophora;s_Harpophora maydis
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_unidentified;s_Magnaporthaceae
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_unidentified;s_Magnaporthaceae
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_Magnaporthiopsis;s_Magnaporthin
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_Magnaporthes;s_Magnaporthes salu
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_unidentified;f_unidentified;g_unidentified;s_Sordariomycetes sp
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_unidentified;s_Magnaporthaceae
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_Buergenerula;s_Buergenerula spa
k_Fungi;p_Ascomycota;c_Eurotiomycetes;o_Chaetothyriales;f_Herpotrichiellaceae;g_Phalophora;s_Phalophora s
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_unidentified;s_Magnaporthaceae
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_Pyricularia;s_Pyricularia angul
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k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_Harpophora;s_Harpophora oryzae
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k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_Gaeumannomyces;s_Gaeumannomyces
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_unidentified;s_Magnaporthaceae
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_unidentified;f_unidentified;g_unidentified;s_Sordariomycetes sp
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_Pyricularia;s_Pyricularia zingi
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Magnaporthales;f_Magnaporthaceae;g_Pyricularia;s_Pyricularia sp MA
k_Fungi;p_Basidiomycota;c_Agaricomycetes;o_Boletales;f_Sclerodermataceae;g_Scleroderma;s_Scleroderma sp
k_Fungi;p_Basidiomycota;c_Agaricomycetes;o_Boletales;f_Boletaceae;g_Boletus;s_Boletus sp
k_Fungi;p_Ascomycota;c_unidentified;o_unidentified;f_unidentified;g_unidentified;s_Ascomycota sp
k_Fungi;p_Ascomycota;c_Leotiomycetes;o_Helotiales;f_unidentified;g_unidentified;s_Helotiales sp
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Hypocreales;f_Cordycipitaceae;g_unidentified;s_Cordycipitaceae sp
k_Fungi;p_Ascomycota;c_Sordariomycetes;o_Sordariales;f_unidentified;g_unidentified;s_Sordariales sp
k_Fungi;p_Ascomycota;c_Lecanoromycetes;o_Peltigerales;f_Pannariaceae;g_Psoroma;s_Psoroma fruticosum
k_Fungi;p_Ascomycota;c_Lecanoromycetes;o_Peltigerales;f_Pannariaceae;g_Psoroma;s_Psoroma buchananii
k_Fungi;p_Ascomycota;c_Lecanoromycetes;o_Peltigerales;f_Pannariaceae;g_Psoroma;s_Psoroma paleoderm
k_Fungi;p_Basidiomycota;c_Agaricomycetes;o_Agaricales;f_Inocybaceae;g_Inocybe;s_Inocybe chondroderma
k_Fungi;p_Basidiomycota;c_Agaricomycetes;o_Agaricales;f_Inocybaceae;g_Inocybe;s_Inocybe sororia
k_Fungi;p_Basidiomycota;c_Agaricomycetes;o_Agaricales;f_Inocybaceae;g_Inocybe;s_Inocybe glabripes
k_Fungi;p_Basidiomycota;c_Agaricomycetes;o_Agaricales;f_Inocybaceae;g_Inocybe;s_Inocybe sp
k_Fungi;p_Basidiomycota;c_Agaricomycetes;o_Agaricales;f_Inocybaceae;g_Inocybe;s_Inocybe cf microspora TAA18
k_Fungi;p_Basidiomycota;c_Agaricomycetes;o_Agaricales;f_unidentified;g_unidentified;s_Inocybaceae sp
k_Fungi;p_Basidiomycota;c_Agaricomycetes;o_Agaricales;f_unidentified;g_unidentified;s_Agaricales sp
k_Fungi;p_Basidiomycota;c_Agaricomycetes;o_Agaricales;f_Inocybaceae;g_Inocybe;s_Inocybe subnudipes
k_Fungi;p_Basidiomycota;c_Agaricomycetes;o_Agaricales;f_Inocybaceae;g_Inocybe;s_Inocybe posterula
k_Fungi;p_Basidiomycota;c_Agaricomycetes;o_Agaricales;f_unidentified;g_unidentified;s_Agaricales sp

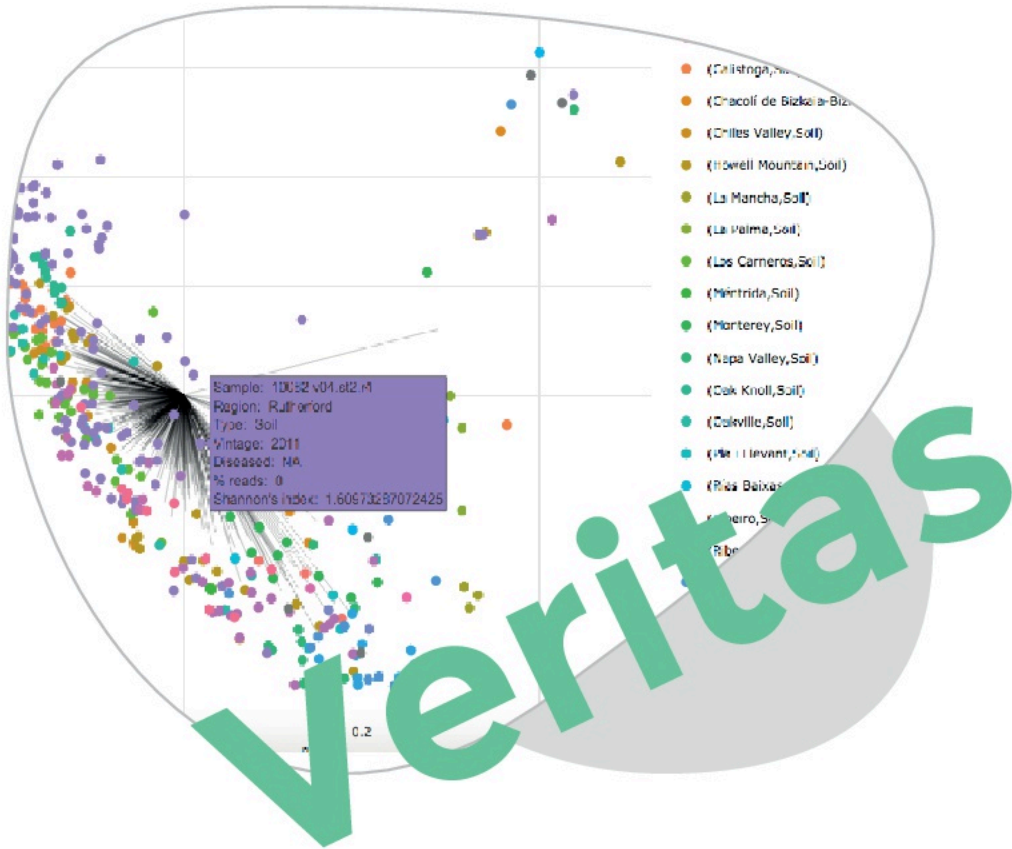
WineSeq Technology – metadata database



119 Boxes



Our internal intelligence platform



We take advantage of big data tools to:

- ✕ Understand global trends in plant diseases distribution and virulence
- ✕ Certificate sustainable farming practices
- ✕ Drive microbial discovery projects
- ✕ Create Biobanks of microbial resources

Understanding the Microbiome



E

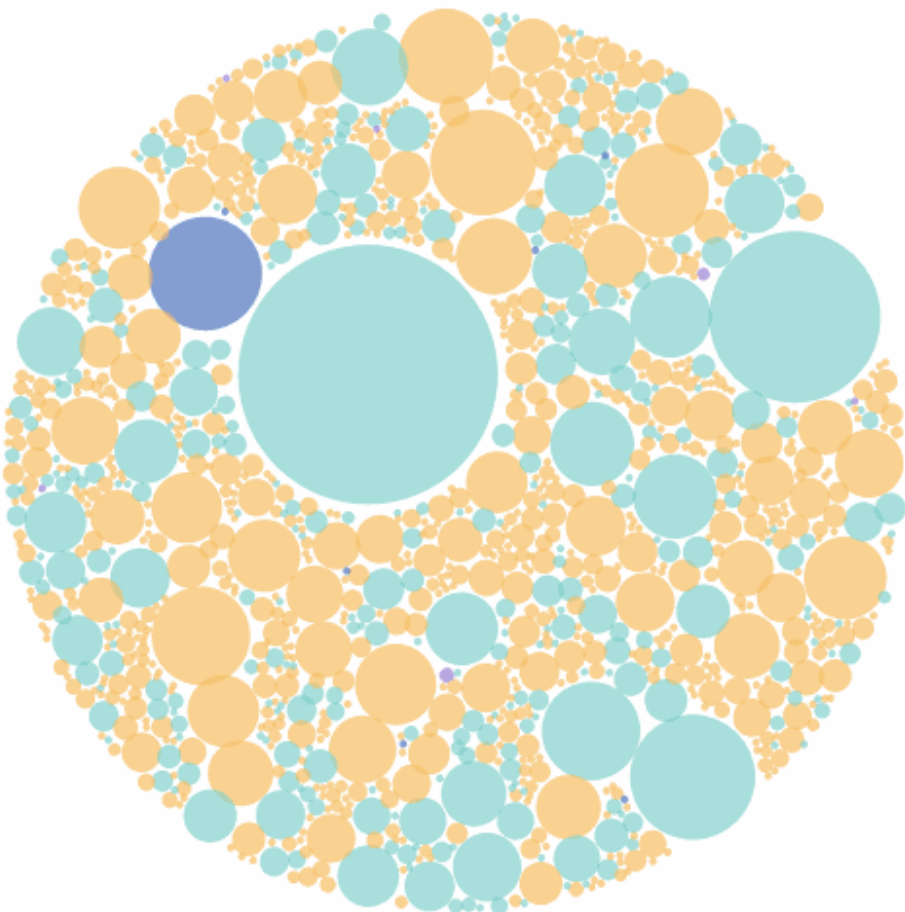
Fungus

Bacteria

Archaea

Yeast

Protist



Type here the key word...

624 ORGS

| | |
|--------------------------------|--------|
| <i>Cryptococcus podzolicus</i> | 26.03% |
| <i>Cryptococcus aerius</i> | 19.21% |
| <i>Fusarium oxysporum</i> | 10.17% |
| <i>Acinetobacter sp.</i> | 9.63% |
| <i>Arnim sp.</i> | 8% |
| <i>Pseudomonas putida</i> | 6.29% |
| <i>Cryptococcus terricola</i> | 5.27% |
| <i>Bacillus sp.</i> | 4.21% |
| <i>Ktedonobacter sp.</i> | 3.86% |
| <i>Nitrososphaera sp.</i> | 3.67% |
| <i>Penicillium restrictum</i> | 3.59% |
| <i>Singulisphaera sp.</i> | 3.45% |
| <i>Cometasp.</i> | 3.00% |

MWM INFO

X



Pyrenochaeta sp.

Fungus

ABUNDANCE

%

1.5081

CATEGORY

Potential plant pest

HIGHLIGHT

Leaf spot related

[Know more in Wikibiome](#)

INFLUENCE



Plant

4 INFLUENCES OF TERROIR

CLIMATE



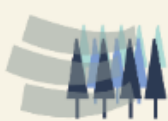
COLD



HOT



WINDY



Precipitaciónón

SOIL



ROCK + MINERAL DEPO



Estructura del suelo

TERRAIN



NORTH OR SOUTH
FACING

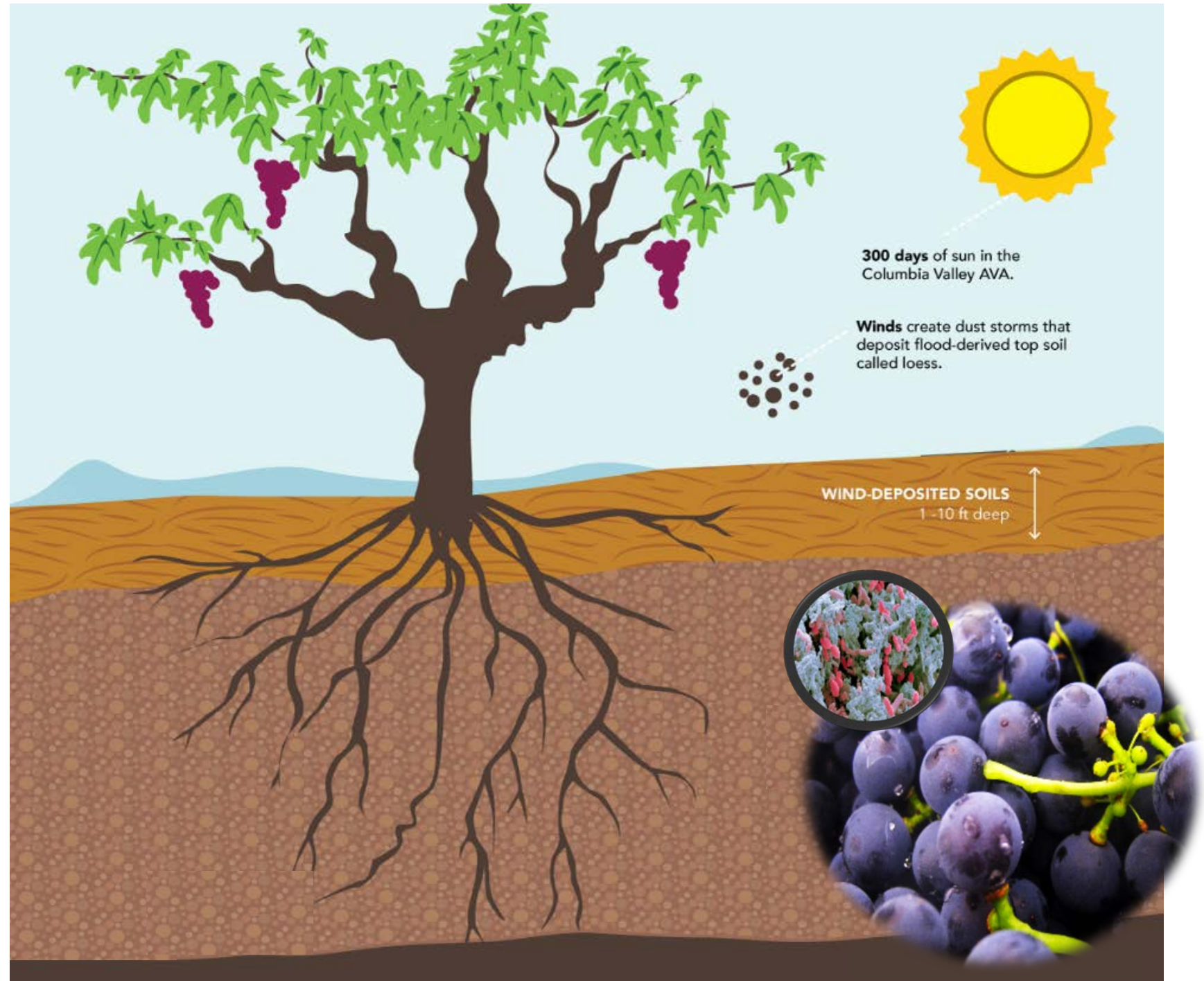


Altitud

TRADITION



TECHNIQUE + WINEMAKER



SUSTAINABLE MICROBIAL BIODIVERSITY CERTIFICATION

CONCLUSION



Biofertilizer MetaCrop Biome X respects the native microbial community, avoid most of crop pathogens and improve Biocontrol and Nitrogen Fixation Species.



PATHOGENS



NITROGEN FIXATORS



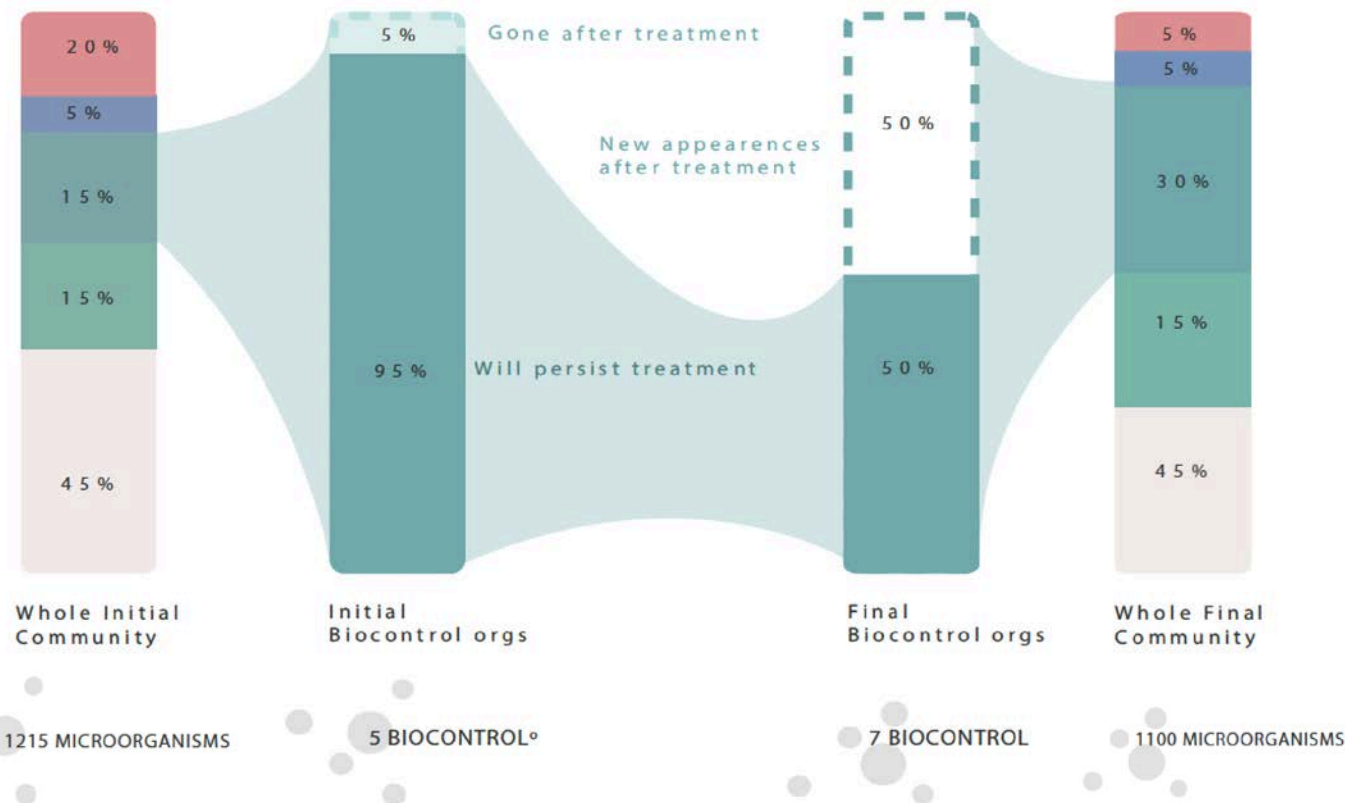
BIOCONTROL



PLANT GROWTH PROMOTERS

BEGIN
AAA001

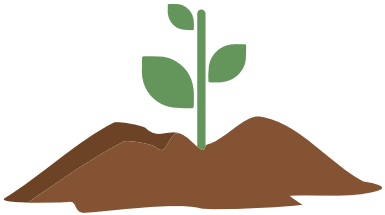
END
AAA002



Soil as BioMarker of Agriculture Practices



Soil

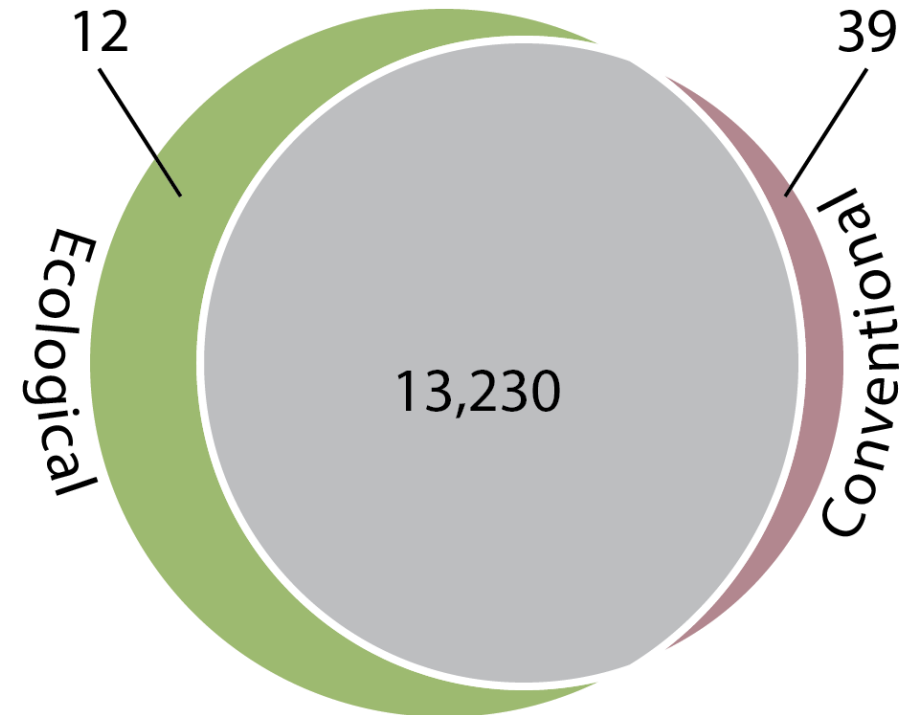


association with...

Farming Practices



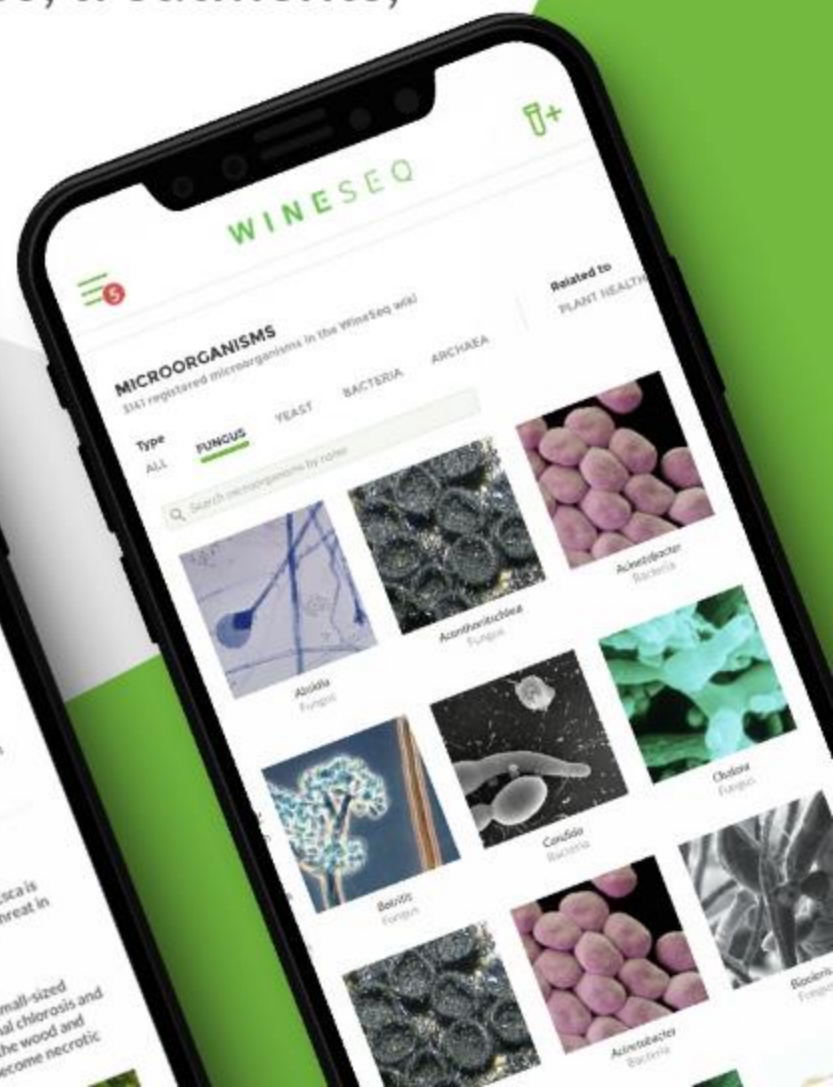
conventional
organic
biodynamic



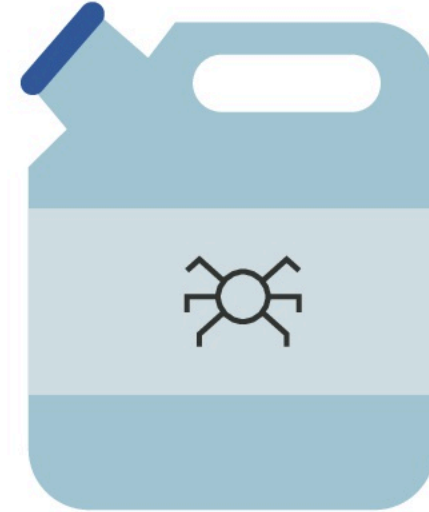
WEB PORTAL

Actionable information for users

WineSeq Portal provides information about sample results, diseases, treatments, and microorganisms



Applying smart discovery



**Bioactive product to
restore burn soil health**



WINESQ

PRECISION ENOLOGY



So far we have...

18

countries

600

users

30000

samples

Top wineries
working with us



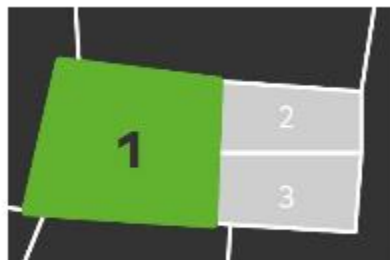
WineSeq is a Methodology: Sample collection

E



1

DOWNLOAD
THE APP
DESCARGA LA APP



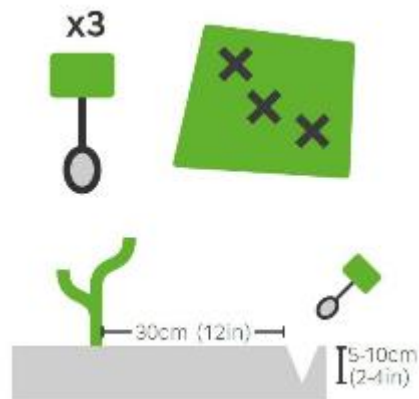
2

CHOOSE
A PLOT
*SELECCIONA UNA
PARCELA*



3

TAKE
A PICTURE
HAZ UNA FOTO



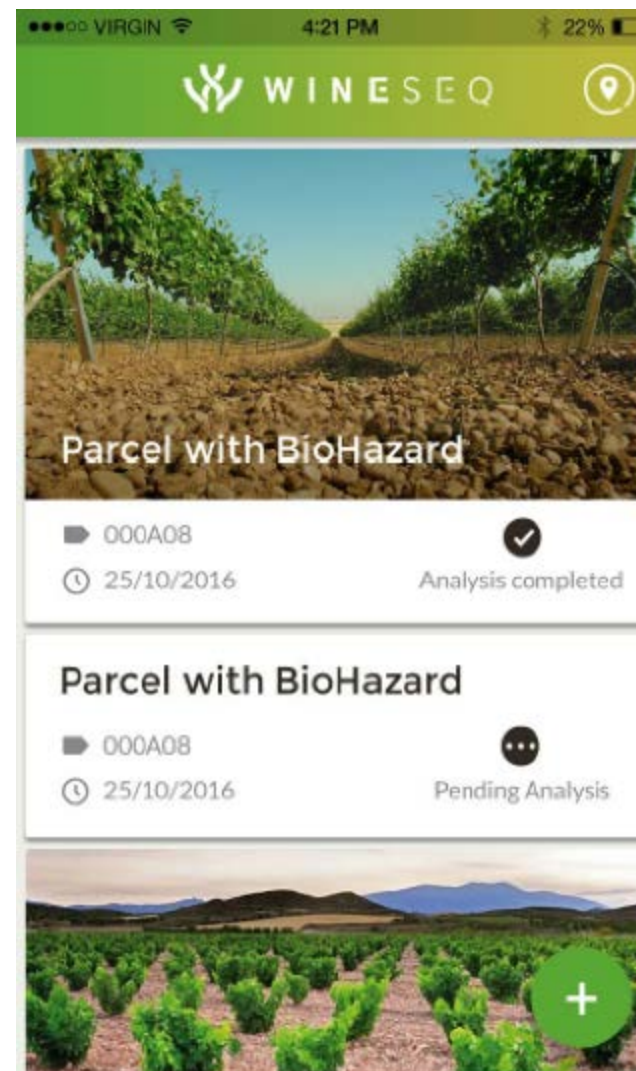
4

COLLECT
THE SAMPLE
RECOGE LA MUESTRA



5

SEND THE
TUBE BACK
ENVÍANOS EL TUBO



Crowding knowledge - Open Information



GRAPE VARIETIES



[298 grape varieties](#) registered in Wikibiome

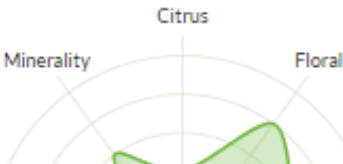
FEATURED GRAPE VARIETY

Verdejo
White

Very sensitive to powdery mildew. Moderately to botrytis.
Medium-alcohol content wines, greenish-yellow in hue, very aromatic, with aromas of bitter almonds. Medium to high acidity wines with body but with a



[Learn more](#)



VINE DISEASES



[61 vine diseases](#) registered in Wikibiome

FEATURED DISEASE

Pierce's disease
Treatable

Aggressive

Pierce's disease is known since 1892 after the description made by Newton Pierce. Pierce's disease is recognized as the most devastating grape disease caused by a bacterium. The disease is severe in countries like US, and is



5 TESTED TREATMENTS



PREVENTION INFO

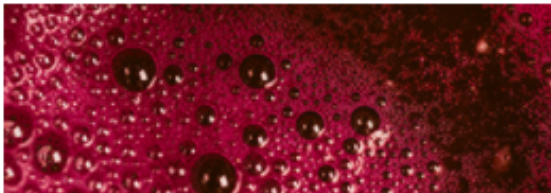
[Learn more](#)

EDITED RECENTLY



Angular leaf scorch / Rotbrenner

WINE MAKING MICROORGANISMS



[2672 microorganisms](#) registered in Wikibiome

FEATURED MICROORGANISM

Armillaria mellea
Fungus

Plant Health
Wine Making

Also known as "Honey fungus", *Armillaria* is a genus of parasitic fungi that lives on trees and woody shrubs. It includes about 10 species formerly lumped together as *A. mellea*. *Armillarias* are long lived and form some of the largest



NOT FOUND IN YOUR SAMPLES

[Learn more](#)

EDITED RECENTLY

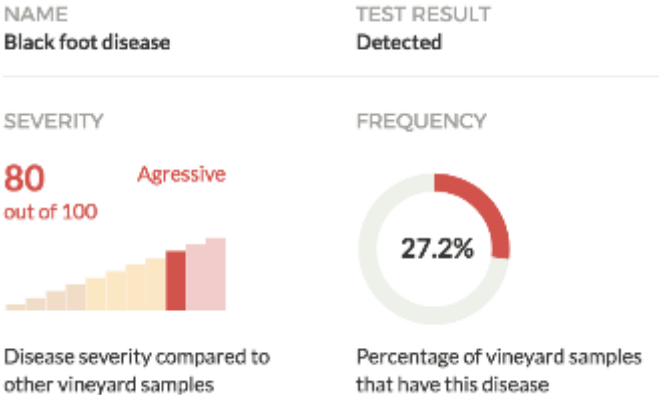
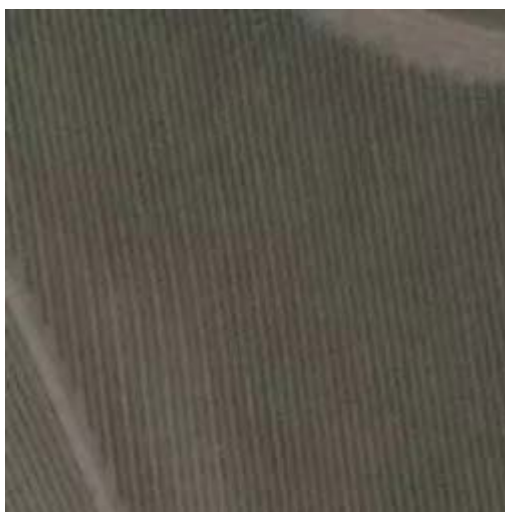


Plasmopara viticola
Protist

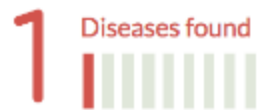
Success Cases & Curiosities

Heavy treated vineyard: Healthy soil or dead soil?

E



DISEASE RISK TEST



Based on

660
Microorganisms found in the sample

[Visualize these abundances in the toolkit](#)

Average in Organic soils: 1.067 sp

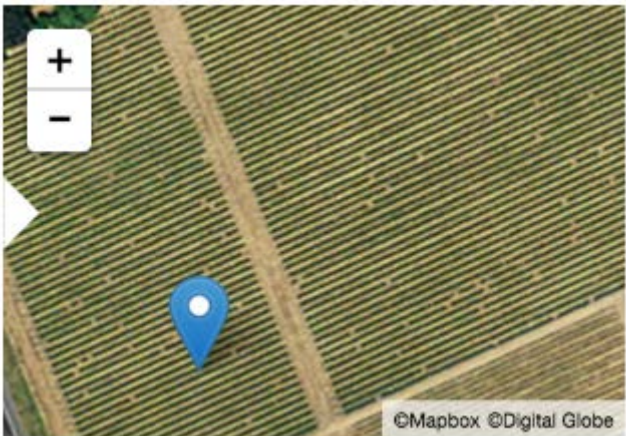
POTENTIAL FERMENTATION TEST



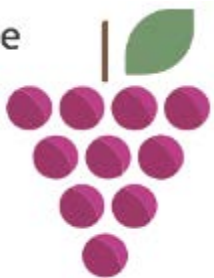
[VIEW FULL REPORT](#)

Brett detection in Soil, Grape and Wine

E



Grape



SEVERITY

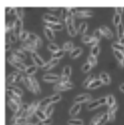
20
out of 100

Mild



Risk severity compared
to other vineyard
samples

Soil



Brettanomyces bruxellensis

Yeast

SEVERITY

70
out of 100

Agressive



Risk severity compared
to other vineyard
samples

FREQUENCY

1.1%



Percentage of vineyard
samples that have this
fermentation risk

BASED ON



1 Fungus

[View full
list](#)

Different
microorganisms found
in the sample



SEVERITY

<10
out of 100

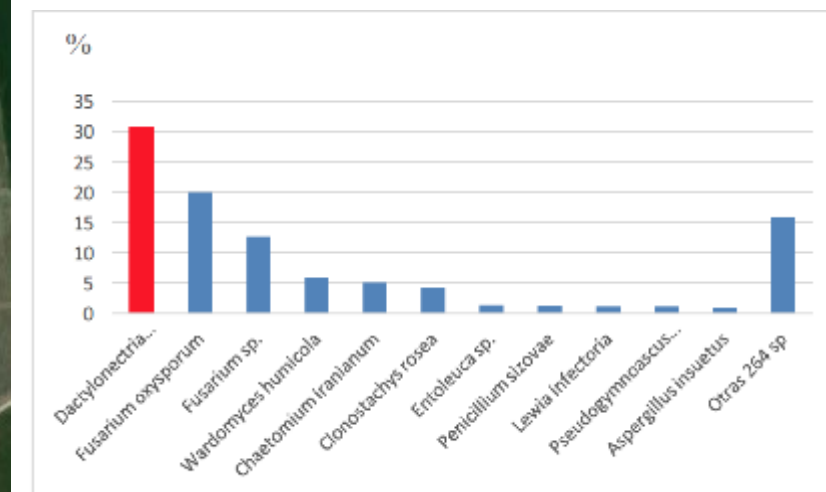
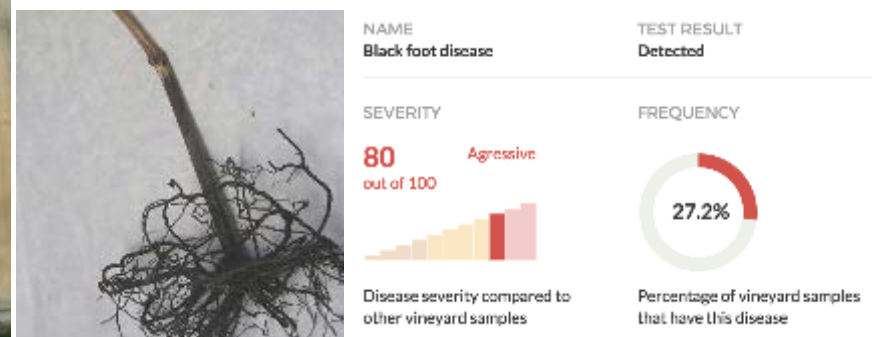
Mild



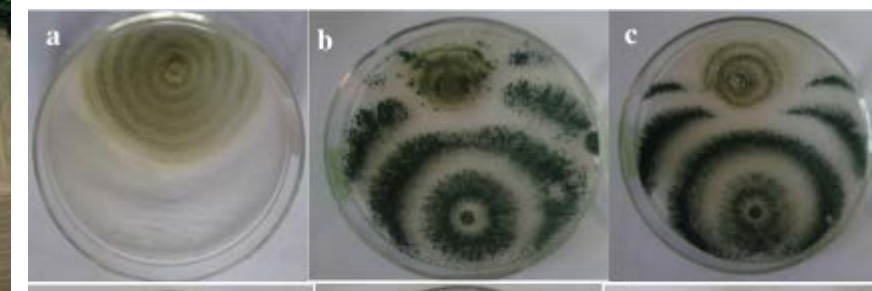
Risk severity compared
to other vineyard
samples

Grape pomace and Skin
compost





5 Biocontrol sp



Success cases

www.portal.wineseq.com/success_cases

WineSeq
works and the
proof is here



I  **MY**
MICROWINEMAKERS

 **WINESEQ**
www.wineseq.com

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