

Multimomics approaches for the control of plant-parasitic nematodes

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Cláudia Vicente



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25 years of Nematolab (Nematology Lab, MED-UE) – major research lines



Manuel Mota, PhD



Cláudia Vicente,
PhD – CEEC'18



Margarida Espada,
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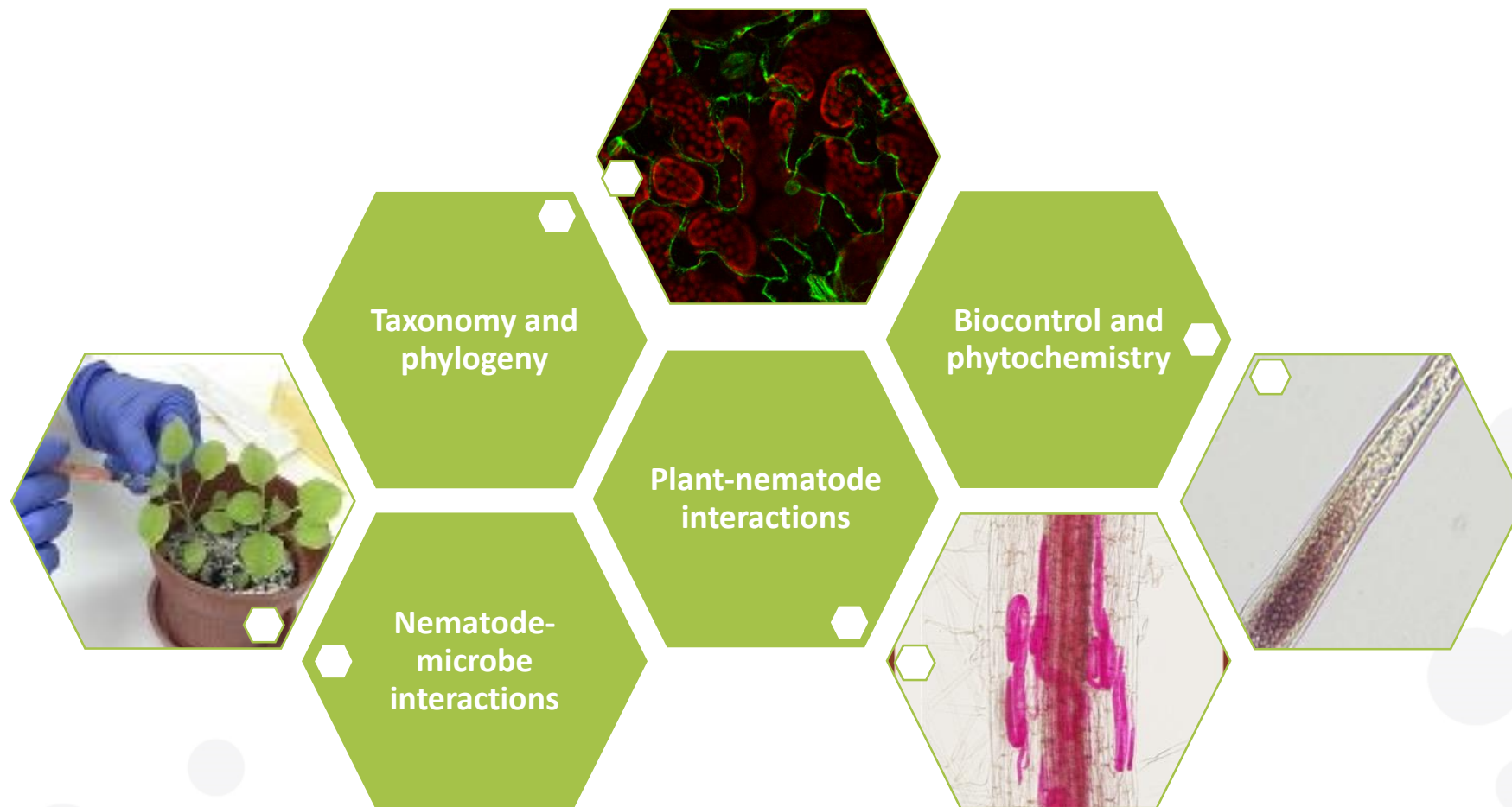
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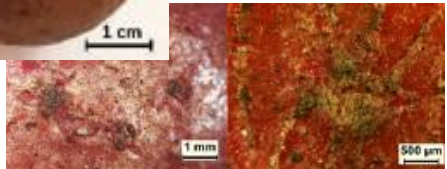
Plant-parasitic nematodes of interest

Root-lesion Nematodes



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Pratylenchus penetrans



Figureiredo et al., 2021, Plant Pathol. 1-9

Pinewood Nematode



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Bursaphelenchus xylophilus



Biotechnological approaches for the control of the root-lesion nematode *Pratylenchus penetrans*



Manuel Mota, PI



Paulo Vieira, co-PI

MAIN GOALS

- Study of the **molecular mechanisms** involved in the parasitism of *Pratylenchus penetrans* for the development new and species-specific strategies of control
- Generation of *Pratylenchus penetrans* **transcriptome infecting potato plants**
- Application of **silencing methodologies (RNAi soaking and *in planta* RNAi)** for the knock-down of target genes of *P. penetrans* involved in nematode-plant interaction

FCT funding: 199.555,20€



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 @PratyOmics



Cláudia Vicente, PI



Maria de Lurdes Inácio, co-PI
INIAV

Plant metabolomics for the control of the root-lesion nematode *Pratylenchus penetrans*

MAIN GOALS

- **Metabolome** comparison between **resistant** and **susceptible** potato cultivars in response to *P. penetrans* infection for the identification of resistant-related metabolites (e.g., plant secondary metabolites)
- **Bioactivity evaluation of secondary metabolites** in *P. penetrans*
- Study the **mode of action** of the most promising secondary metabolites in the **control** of *P. penetrans*



FCT funding: 249.936,90€



A *motif* to unveil mechanisms of parasitism gene regulation in the pinewood nematode as a target for disease control and plant resistance.

How is parasitism regulated?



Margarida Espada, PI



Manuel Mota, co-PI

MAIN GOALS

- Identify **transcription regulators** that are associated with secreted **parasitism-related genes**;
- Identify the **protein complexes that bind to non-coding DNA motif**
- **Knock-down** (RNAi) candidate regulators and evaluate the effect in nematode parasitism-related genes
- *In planta*: evaluate the effect of **knock-down or over-expression** of the regulators in nematode parasitism

FCT funding: 249.439,00€



Consultants



Plant metabolomics for the control of the root-lesion

nematode *Pratylenchus penetrans*

Claudia Vicente^{1,2*}, Jorge Faria¹, Dora Teixeira², Paulo Vieira³, Margarida Espada¹, Pedro Barbosa¹, Manuel Mota³, Joana Figueiredo⁴, Joana Sá Cardoso⁴, Ivánia Esteves⁴, Isabel Abrantes⁴, Rita Varela¹, Filomena Nóbrega², Maria de Lurdes Inácio¹

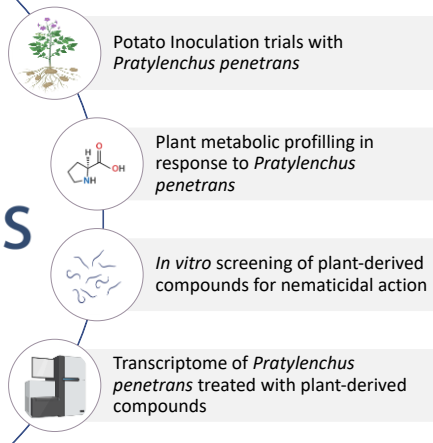
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Pratylenchus penetrans, one of the most detrimental root-lesion nematode species, greatly reduces the production in numerous important agronomic crops (e.g., corn, potato), ornamental plants (e.g., lily, roses) and fruit trees (e.g., almond, cherry orchards). In the EU, *P. penetrans* has been reported as the most damaging species associated with potato (*Solanum tuberosum* L.). In Portugal, this species was also detected in potato production fields across the country and often related with their low yield.

Plant metabolomics is an emerging approach to the study crop resistance against plant-parasitic nematodes, which can be applied to expedite traditional crop breeding programs and the development of novel pesticides.

PratyOmics



PratyOmics aims:

- (i) to compare global metabolomic profiling of resistant and susceptible potato cultivars in response to *P. penetrans* infection for the identification of host resistant-induced plant secondary metabolites (PSM) with potential anti-nematode activity;
- (ii) to evaluate the nematocidal bioactivity of the candidate PSM against *P. penetrans*; and
- (iii) to understand the mechanism of action of the most promising PSM by transcription profiling of nematode affected nematode molecular pathways, which can be translated into targets for the **development of new effective nematode control strategies**.

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NemaWAARS



Unveil mechanisms of parasitism gene regulation in the plant-parasitic nematode *Bursaphelenchus xylophilus*, as a target for disease control and plant resistance.



Margarida Espada¹, Cláudia Vicente¹, Jorge Faria², Pedro Barbosa¹, Isabel Velada¹, Margarida Santos², Filomena Nóbrega², Paulo Vieira³, Sebastian Eves-van den Akker⁴, Maria de Lurdes Inácio², Manuel Mota¹

¹NemaLab-MED – Mediterranean Institute for Agriculture, Environment and Development, Institute for Advanced Studies and Research, Universidade de Évora, Pólo da Mitra, Évora, Portugal | ²Nematology Lab-INIAV – Instituto Nacional de Investigação Agrária e Veterinária, Oeiras, Portugal | ³United States Department of Agriculture (USDA), Agricultural Research Services (ARS), Mycology & Nematology Genetic Diversity & Biology Lab, Beltsville, MD 20705, USA | ⁴Department of Plant Sciences, University of Cambridge, Cambridge, United Kingdom

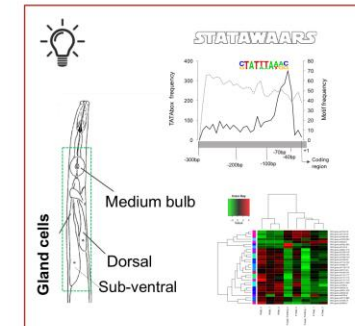


Figure 1 | Concept idea of NemaWAARS (Espada *et al.*, 2018; Eves van-den Akker *et al.*, 2016)

The interaction of plant-parasitic nematodes (PPN) with their plant hosts are mediated by parasitism proteins (effectors) that interact and/or modify host proteins to promote infection. NemaWAARS project focuses on mechanism(s) of regulation and gene control expression of parasitism genes of the migratory endoparasite *Bursaphelenchus xylophilus* (the pinewood nematode). Supported by genomic and transcriptomic data previously established for this pathogen we have identified a DNA motif - STATAWAARS - associated in the promoter region of secreted pharyngeal gland cells coding effector genes. Given that this non-coding genetic signature unifies many sequences of unrelated effectors, it implies the existence of a potential major regulator(s), that binds to this sequence to orchestrate the expression of downstream effector genes. A similar master regulator has been recently proposed for other PPNs (Tylenchida) non-related to migratory nematodes or to this clade of nematodes (Aphelenchida). We hypothesize that by disrupting this regulator(s), it would be possible to simultaneously disrupt the expression of a large number of associated effector genes. This could be a new attractive target for host induced gene silencing, as switching-off the regulator of numerous effectors at the same time, can reduce and inhibit the performance of this parasitic nematode. This project aims to identify transcription regulators associated with this motif and understand the impact of silencing enriched gland cells transcription factors on the PWN interaction with the host. The strategy in NemaWAARS will include innovative approaches to develop a better understanding of how the effectors are regulated, their function in the cell host and use their sequences as potential targets for genetic editing towards an effective nematode resistance in the plant. The ability to explore host-delivered mechanisms against the pathogens could have a tremendous impact as biotechnology standpoint in important forestry species.

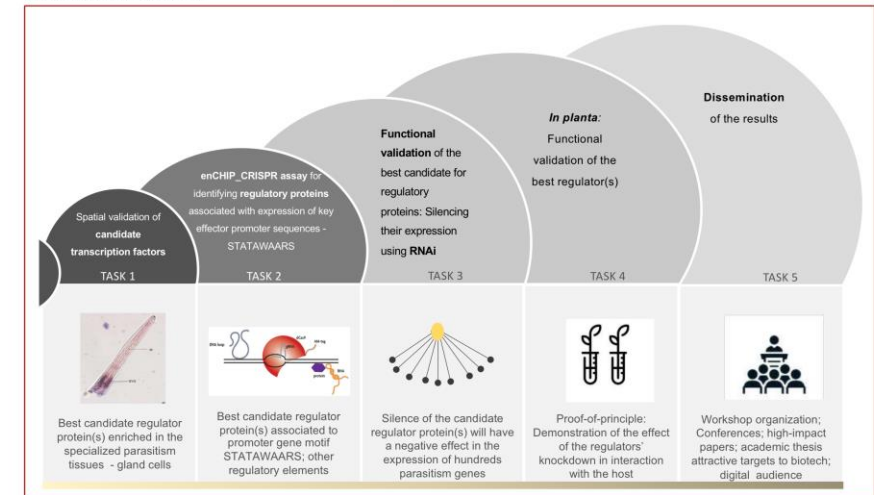
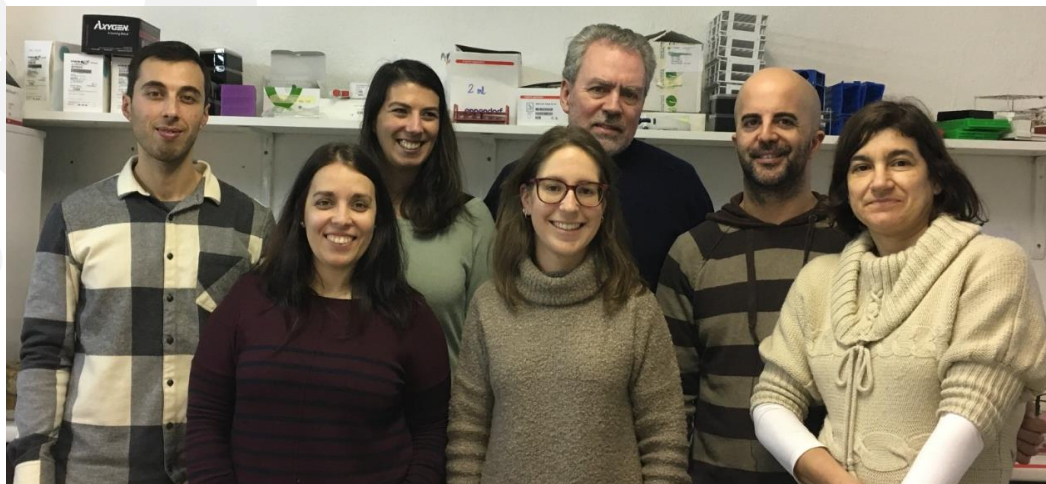


Figure 2 | Workflow of NemaWAARS project

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Our collaborations



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Ivânia Esteves
Joana Sá Cardoso
Joana Figueiredo



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Paulo Vieira
Lev G. Nemchinov



Ana C. Figueiredo



Fátima Duarte
Teresa Brás



Sebastian Eves-van den Akker



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Thank you!
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