

### XXXV ECCO MEETING

# WHO/WHERE ARE WE IN 2016 AND BEYOND? JURY'S INN HOTEL ABERDEEN 2016

## Multilocus molecular characterization of a *Penicillium* strain isolated from maize in Nigeria

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#### Aims

Access the phylogeny of *Penicillium* strain (MUM 14.07) within the *P. sclerotiorum* complex and the relationship with *P. mallochii*.

#### Methods and results

The *Penicillium* strain (MUM 14.07) was subjected to multilocus molecular phylogenetic analysis using partial  $\beta$ -tubulin (*BenA*), calmodulin (*CaM*), cytochrome c oxidase subunit 1 (*cox1*), internal transcriber spacer (ITS) region and translation elongation factor 1- $\alpha$  (*tef1-\alpha*).

The multilocus phylogenetic analysis revealed that the *Penicillium* strain (MUM 14.07) clustered closely with *P. mallochii* (1), revealing an unconditional position within the section Sclerotia. The sole exception was for the *BenA* gene where the studied strain did not show good homology with any particular *Penicillium* species. The Basic Local Alignment Search Tool (BLAST) of the sequence obtained of *BenA* gave a higher similarity with species from the section Citrina.

#### Conclusions

With this result we cannot state that our strain is a *P. malochii* but we can consider that they are two related species that were separated geographically. It would be interesting to be able to isolate more strains.

#### Significance of study

This case study shows that many species have a common ancestor and that *Penicillium sp.* MUM 14.07 and *P. mallochii* might belong to the same monophyletic group.

#### References

(1) Rivera KG, Díaz J, Chavarría-Díaz F, Garcia M, Urb M, Thorn RG, Louis-Seize G, Janzen DH, Seifert KA (2011). *Penicillium mallochii* and *P. guanacastense*, two new species isolated from Costa Rican caterpillars. Mycotaxon **119**:315-328.