Melanin production in coelomycetous agents of black grain eumycetoma

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Background: Eumycetoma is a fungal infection characterised by the formation of black grains by causative agents. The melanin biosynthetic pathways used by the most common causative agents of black-grain mycetoma are unknown and unravelling them could identify potential new therapeutic targets.

Method: Melanin biosynthetic pathways in the causative fungi were identified by the use of specific melanin inhibitors.

Results: In Trematosphaeria grisea and Falciformispora tompkinsii, 1,8-dihydroxynaphthalene (DHN)-melanin synthesis was inhibited, while DHN-, 3,4-dihydroxyphenylalanine (DOPA)- and pyo-melanin were inhibited in Medicopsis romeroi and Falciformispora senegalensis.

Conclusion: Our data suggest that Me. romeroi and F. senegalensis synthesise DHN-, DOPA- and pyo-melanin, while T. grisea and F. tompkinsii only synthesise DHN-melanin.

Keywords: Falciformispora senegalensis, Falciformispora tompkinsii, Medicopsis romeroi, melanin, mycetoma, Trematosphaeria grisea

Introduction

Eumycetoma is a subcutaneous fungal disease characterised by the presence of black, white or occasionally yellow grains in infected tissue. It is recognised as a neglected tropical disease by the WHO and can be caused by more than 40 different fungal species. Four of the five most prevalent eumycetoma-causing agents form black grains, namely, Madurella mycetomatis, Falciformispora senegalensis, Trematosphaeria grisea and Medicopsis romeroi, while the fifth, Scedosporium boydii, forms white grains. Melanin was found to be responsible for the black colour in Ma. mycetomatis. Melanins are hydrophobic, negatively charged macromolecular pigments formed by oxidative polymerisation of phenolic or indolic compounds. They contribute to virulence and play a role in the protection against various environmental stresses, antifungal agents and host defences. It is known that melanised fungi are often more difficult to treat and have more relapses compared with non-melanised fungi. Three melanin biosynthesis pathways have been described in fungi: the 1,8-dihydroxynaphthalene (DHN)-, the 3,4-dihydroxyphenylalanine (DOPA)- and the pyo-melanin biosynthetic pathways. Interference of DHN-melanin synthesis can be achieved either by inhibiting tetrahydroxynaphthalene reductase with tricyclazole (TCZ) and pyraquilon (PYR), or by inhibiting scytalone dehydratase with carpropamid (CAR) and fenoxanil. Glyphosate (GLY) inhibits DOPA-melanin, while sulcotrione (SCT) inhibits pyo-melanin. We have shown that Ma. mycetomatis forms melanin via DHN- and pyo-melanin pathways; however, the pathways used in F. senegalensis, T. grisea and Me. romeroi are still unknown. To understand the types of melanin involved in grain synthesis, we used melanin inhibitors to determine the melanin biosynthetic pathways present in these prevalent agents of human eumycetoma, and in the related, less common agent of eumycetoma, F. tompkinsii.

Materials and methods

All the melanin inhibitors were obtained from Sigma-Aldrich (Sigma-Aldrich Chemie NV, Zwijndrecht, the Netherlands) and were dissolved in Dimethyl sulfoxide (DMSO). Three T. grisea, three Me. romeroi, two F. tompkinsii and one F. senegalensis were cultured on minimal media supplemented with 0.5, 5 or 50 mg/L melanin inhibitors as described...
Results and discussion

To determine the types of melanin involved in grain synthesis, melanin inhibition was performed on F. senegalensis, T. grisea, Me. romeroi and F. tompkinsii. Our data revealed that all four species were inhibited by DHN-melanin inhibitors, similar to Ma. mycetomatis in earlier studies (Figure 1).2 Trematosphaeria grisea and F. tompkinsii were inhibited by TCZ and PYR, while Me. romeroi and F. senegalensis were inhibited in two steps of the DHN-melanin pathway, as observed in TCZ, PYR, CAR and PYR inhibition. An orange hue was noted in isolates inhibited by TCZ and PYR due to the production of coloured by-products in the melanin biosynthesis pathway such as flaviloin and juglone.4 Inhibition in expansion growth was also noted. The highest reduction in growth was observed when DHN-melanin was inhibited by TCZ (median 4 mm) (Figure 1).

In addition to DHN melanin, Ma. mycetomatis also forms pyo-melanin. Using the pyo-melanin inhibitor SCT, a slight reduction of colony darkness and diameter was noted in Me. romeroi and F. senegalensis, while no reduction in melanisation or expansion growth was observed in T. grisea and F. tompkinsii, indicating that only Me. romeroi and F. senegalensis formed significant amounts of pyo-melanin. Medicopsis romeroi and F. senegalensis were also the only species inhibited by the DOPA-melanin inhibitor GLY. A diffusible pigment was also present in Me. romeroi inhibited with GLY, similar to what was observed in isolates inhibited by TCZ and PYR (Figure 1). Consequently, the pathways of melanin production were concluded to be species-dependent; in summary, DHN-melanin in T. grisea and F. tompkinsii and DHN-, DOPA- and pyo-melanin in Me. romeroi and F. senegalensis. Remarkably, sibling species F. tompkinsii and F. senegalensis differ in the presence of melanin pathways even although they are phylogenetically close, as shown by similarity in their rDNA Internal transcribed spacer (ITS) gene.3 The ITS sequences of F. senegalensis CBS198.79 are 86.03% (490 out of 580 bp) and 86.06% (500 out of 580 bp), identical to those of F. tompkinsii isolates CBS200.79 and CBS201.79, respectively. In F. senegalensis, pyo-melanin and DOPA-melanin were present, while in F. tompkinsii these were absent or present below the detection level of the applied method. Since the analysed fungi are able to synthesise melanin using multiple pathways, inhibition of a single pathway is not sufficient to completely stop melanin production. To determine if complete eradication of melanin pigmentation of the mycelium can be achieved by simultaneous inhibition of multiple biosynthetic pathways, the strains were subjected to four combined inhibition protocols, namely, inhibition of DHN and DOPA (PYR and GLY), pyo-melanin and DOPA (SCT and GLY), DHN and pyo-melanin (PYR and SCT), and DHN-, pyo- and DOPA-melanin (PYR, SCT and GLY). No enhanced melanin inhibition was noted compared with inhibition by single inhibitors. It may be hypothesised that these inhibitors alone were not enough to completely inhibit their respective melanin-biosynthesis pathways, leading to melanin production via a different route.

A clear limitation of this study is that our data were derived from only nine fungal isolates. While these nine fungal isolates do not reflect all eumycetoma-causing agents, there is unfortunately no larger collection of these fungi currently available in the world. With a larger collection also consisting of other eumycetoma-causing agents, we may be able to obtain a clearer understanding of the types of melanin involved in grain synthesis.

DHN-melanin is known to be present in Ma. mycetomatis and was found in all the mycetoma agents tested here. Since a reduction in expansion growth was noted with some of the DHN-melanin inhibitors and DHN-melanin is the prevalent type of melanin in all fungi investigated, it could be explored as a potential drug target for some of the causative agents. An interesting approach might be to enhance the efficacy of azole therapy for mycetoma since the presence of melanin is known to lead to decreased drug susceptibility. In Ma. mycetomatis, melanin was able to bind to itraconazole, the drug currently in use for mycetoma treatment and lower inhibitory concentrations.2 A similar phenomenon has also been observed with Aspergillus fumigatus DHN-melanin.2 Since DHN-melanin is present in all the causative agents tested here, it could be envisioned that similar protection would occur in these species as well. Therefore, to enhance the efficacy of the azoles currently used in mycetoma treatment, combination treatment with melanin inhibitors should be investigated.

Authors’ contributions: WWJS conceived the study; WWJS and FP designed the study protocol; FP and WL implemented the study; FP, WL and WWJS performed analysis and interpretation of the data. WL drafted the manuscript; FP, SH, AV and WWJS critically revised the manuscript for intellectual content. All the authors read and approved the final manuscript.
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