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Characterization of *Emericella nidulans* RodA and DewA hydrophobin mutants

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Hydrophobins are small amphiphilic proteins containing an eight cysteine pattern only found in filamentous fungi. They are involved in the attachment of hyphae to hydrophobic structures and the formation of aerial structures. Five *Emericella nidulans* mutant strains were examined to study the two hydrophobins RodA and DewA. Individual knock-out mutants $rodA\Delta$, $dewA\Delta$ and the double deletion strain $rodA\Delta dewA\Delta$ were constructed. Furthermore, two strains containing a point mutation in the first of the cysteines of RodA (rodA-C57G), where one was coupled to the dewA deletion, were included. The reference strain (NID1) and $dewA\Delta$ displayed green conidia. However, $rodA\Delta$ and $rodA\Delta dewA\Delta$ showed a dark green/brown conidial pigmentation, while rodA-C57G and rodA-C57G dewA\Delta displayed lighter brown conidia. $rodA\Delta$ and $rodA\Delta dewA\Delta$ displayed a higher degree of hülle cells compared to the moderate amount observed for NID1 and $dewA\Delta$, while rodA-C57G and rodA-C57G dewA\Delta displayed a low number of hülle cells. NID1 and $dewA\Delta$ conidia were dispersed as spore chains. $rodA\Delta$, $rodA\Delta dewA\Delta$, rodA-C57G and rodA-C57G dewA\Delta spores were associated in large clumps, where the conidia seemed to adhere to one another. The largest degree of spore clustering was observed for $rodA\Delta$ and rodA-C57G dewA\Delta.