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Text S2: Dual mating in Botrytis cinerea (teleomorph Botryotinia fuckeliana).

B. cinerea is unusual among pezizomycete fungi in that some isolates are capable of 'dual mating'. Whereas most isolates act in a standard heterothallic fashion, being of MAT1-1 or MAT-1-2 identity, occasionally isolates are detected that can mate with both MAT1-1 and MAT1-2 isolates (Faretra et al., 1988; van der Vlugt-Bergmans et al., 1993). Some of these 'dual mater' isolates can self-fertilize and are truly homothallic. Thus, B. cinerea provides a rare example in which isolates exhibiting heterothallic, homothallic and dual mating ability are found within the same species. For example, genetic analysis of progeny from a cross between heterothallic B. cinerea isolates showed that the MAT1-1 and MAT1-2 alleles segregated largely in a 1:1 manner. However a small proportion (approximately 5%) of the progeny appeared to have been converted into dual maters (Faretra & Pollastro, 1996; van Kan, unpublished). Ordered tetrad analysis suggested that this was due to uni-directional change of MAT1-2 ascospores to the dual-mating form (Faretra & Pollastro, 1996). In general mating-type identity (and associated breeding system) is stable among filamentous ascomycete fungi, with switching of mating type only reported from very few species including Chromocrea spinulosa, Sclerotinia trifoliorum and certain species of Ceratocystis (Mathieson, 1952; Uhm & Fuji, 1983; Witthuhn et al., 2000; Debuchy et al 2010). The phenomenon of mating-type switching has long puzzled those working on mating and sexual development in ascomycete fungi, and has led to speculation concerning the genetic basis of this transition (Coppin et al., 1997). The only explanation to date has come from studies of Ceratocystis species in which a uni-directional change from a homothallic to a heterothallic state was correlated with loss of MAT1-2 gene sequence at the mating-type locus (Witthurn et al., 2000). However, changes leading to gain of dual-mating ability and homothallism, as observed in B. cinerea, remain unexplained.

The organisation of *MAT* loci in dual-mater strains of *B. cinerea* was investigated. One possible explanation was that dual mating and homothallism has arisen from a *MAT1-1-1* gene becoming incorporated into the genome of an individual already carrying a *MAT1-2-1* gene, with the subsequent presence of both alpha- and HMG-domain genes leading to ability to sexually cross to isolates of both mating type and ability to self-fertilitze (Coppin *et al.*, 1997; Paoletti *et al.*, 2007). Such gene incorporation might have occurred by aberrant cross over, translocation, gene conversion or swapping of a silent gene cassette (as in *S. cerevisiae*). The *MAT* locus and bordering region was therefore sequenced in five isolates that display dual-mater behaviour, comprised of two field isolates (Bc7 and Bc12; van der Vlugt-

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Bergmans et al. 1993), and two single ascospore progeny (RS11 and RS26) from a cross between heterothallic tester strains SAS56 and SAS405. As a control, the MAT loci were sequenced of two progeny from the same cross that displayed classical MAT1-2 behaviour. Surprisingly, no evidence was found for the joint presence of complete MAT1-1-1 and MAT1-2-1 genes in the genomes of the dual-mating strains. Instead, all four dual-mater isolates (Bc7, Bc12, RS11 and RS26) exhibited a MAT1-2 like locus. Polymorphisms were detected between the test and control isolates analysed, but none of the SNPs was consistently shared by all dual-mater isolates. Furthermore, neither PCR analysis nor Southern hybridization revealed any sequence homologous to the opposite MAT gene consensus domain. This excluded the possibility that the dual-mater strains had acquired a functional copy of the opposite mating-type gene elsewhere in their genome, as observed in some *Cochliobolus* and Aspergillus species (Yun et al., 1999; Paoletti et al., 2007; Rydholm et al., 2007). We conclude that dual mating in B. cinerea is not due to modifications at the MAT locus, but instead this phenomenon arose due to genetic changes elsewhere in the genome. It will be necessary to undertake an extended genetic crossing program to determine whether this form of mating-type switching has a monogenic or polygenic basis. It is perhaps significant that one of the few other documented examples of mating-type switching is in the related species S. trifoliorum. This suggests that heterothallic Sclerotiniaceae might be particularly prone to this phenomenon due to possible their origins from a homothallic ancestor, in which aspects of sexual development had become independent of MAT gene control as seen in A. nidulans (Paoletti et al, 2007). In parallel to our findings, there have been previous reports in which self fertility is possible in filamentous ascomyctee despite the presence of only one matingtype gene; namely a limited number of *Neurospora* species, very rare field isolates of Cryphonectria parasitica, a low percentage of Stemphylium species (Inderbitzin et al. 2005) and certain *Glomerella* species, (Coppin et al., 1997; McGuire et al., 2004; García-Serrano et al., 2008). Same-sex mating ability has also been reported as a rare occurrence in the yeast Candida albicans (Alby et al., 2009) and the basidiomycete Cryptococcus neoformans (Lin et al., 2005; Heitman, 2009).

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