Electronic Supplementary Materials and Methods

Figure Supplementary 1
Figure S1. Alignment of the predicted catalytic domain of DmtA homologues.

Conserved DMT motifs are indicated by roman numerals. Clustal W with default settings [1] was used to align the amino acid sequences of the catalytic domains. Identical and highly conserved (i.e., present in eight or more species) are boxed together. Species abbreviations are: An, A. nidulans; Ao, A. oryzae; Af, A. fumigatus; At, A. terreus; Ci, C. immitis; Ur, Unicocarpus reesii; Nc, N. crassa; Nt, N. tetrasperma; Ni, N. intermedia; Cg, C. globosum; Gz, G. zeae; Nh, N. haematococca; Mg, M. grisea; Bf, B. fuckeliana; Ss, S. sclerotiorum; Ai, A. immersus.

References.