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Supplementary Figure S5. Neighbor-Joining (NJ) phylogeny of *Metarhizium* fungi as reconstructed from MzIGS3 (DUF895) marker sequences. Terminal branches are labelled by genus, species and strain designations; "TYPE" denotes the nomenclatural type strain of a species. Accession numbers are indicated for reference sequences identified in similarity searches across the GenBank database. Numbers on branches indicate bootstrap support percentages. The size bar corresponds to 1 % sequence divergence. The orthologous sequences from the *M. majus* and *M. guizhouense* type strains have been used as outgroup.