Correction to: Linking Genes to Traits in Fungi

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This addendum is to clarify that we used the Joint Genome Institute MycoCosm Fungal Genomics Resource (<https://mycocosm.jgi.doe.gov>) to query publicly available genomes for lists of genes associated with seven gene families. The numbers of genes returned by the search engine were manually counted from the lists and then recorded in an Excel spreadsheet for each queried species. In our paper, we call this spreadsheet “a database”; however, no genomic sequence data were downloaded from MycoCosm. Supplementary Table 1 includes publications referencing queried

genomes, when provided on the MycoCosm portal. The genomes queried through the MycoCosm portal were produced by the Joint Genome Institute in collaboration with their user community. The work conducted by the Joint Genome Institute, a DOE Office of Science User Facility, is supported by the Office of Science of the U.S. Department of Energy under Contract No. DE-AC02-05CH11231.

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