

Supplementary Figure 11. Principal components analysis of codon usage for the genes predicted from *Basidiobolus meristosporus* CBS 931.73 from the HGT assay. Rows represent secondary metabolite classes and columns putative origin. The two principal components with the highest variance contribution (PC1, 3% and PC2, 2%) are shown. Points represent the genes from the HGT analysis. Colors represent secondary metabolite classes.