

Supplementary Figure 10. Principal components analysis of 5-meres 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, 14% and PC2, 8%) are shown. Points represent the genes from the HGT analysis. Colors represent secondary metabolite classes.