



Figure S2: Schematic overview illustrating expression distribution of genes within putative biosynthetic clusters of unknown metabolites (Sieber et al. 2014) above the $\log_2(\text{FPKM})$ thresholds demarcated by the 65 % of highest expressed genes within the respective growth condition, which are 1.99 in case of pathogenic (patho), 2.46 in case of saprophytic (sapro) and 1.92 in case of axenic (axo) growth. Three horizontal fields in one line represent one gene and grey indicates expression above the 65 % threshold.