



Extended Data Fig. 12 | Heat map of the top-40 genes enriched in the palaeofaeces, the industrial and the non-industrial samples. Related to Fig. 4, complete results are provided in Supplementary Table 8. Functions were

annotated using PROKKA³⁸ with the UniProtKB database¹⁰⁸. Enriched genes were identified using one-tailed Wilcoxon rank-sum tests with Bonferroni correction. RPKM values are shown on a log scale without scaling.