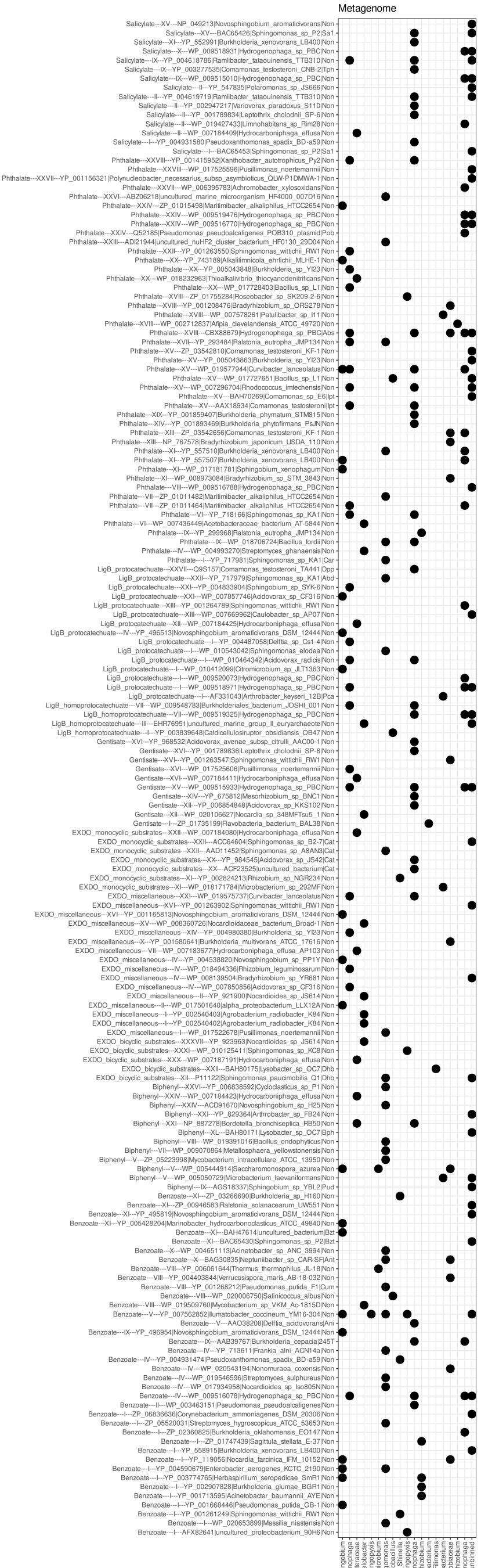
∄



gene counts

groopm_bin_34_Hydrogenophaga-groopm_bin_36_Sinobacteraceae-groopm_bin_43_Sphingopyxis-groopm_bin_48_Hyphomicrobium-groopm_bin_48_Hyphomicrobium-groopm_bin_50_Sphingomonas-maxbinout.010_sub_Shingopyxis-maxbinout.016_sub_Hydrogenophaga-maxbinout.015_sub_Sphingopyxis-maxbinout.016_sub_Hydrogenophaga-maxbinout.017_Meschizobium-metabat_bin21_sub_Filimonas-metabat_bin25_Microbacterium-metabat_bin5_Bradyrhizobiaceae-solidbin_4270_sub_Bradyrhizobium-unbinned-unbinga-groophaga-gro