



Figure S10. Pangenome and core genome plot of eleven *Cylindrospermopsis raciborskii* strains from the Non Americas. (A) Shows the progression of the pangenome: the y-axis indicates the number of non-redundant genes and the x-axis indicates the number of genomes. (B) Shows the progression of the core genome: the y-axis indicates the number of genes present in all strains and the x-axis indicates the number of genomes. The estimated asymptote is 2.249 genes. The evident plateau from the eighth genome onward demonstrates statistical robustness in the estimation.