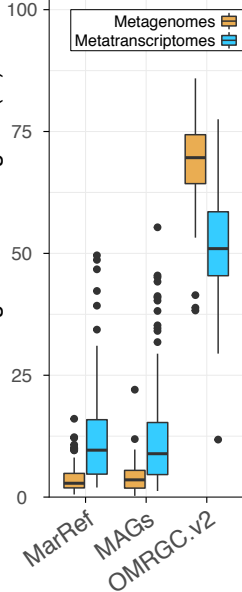
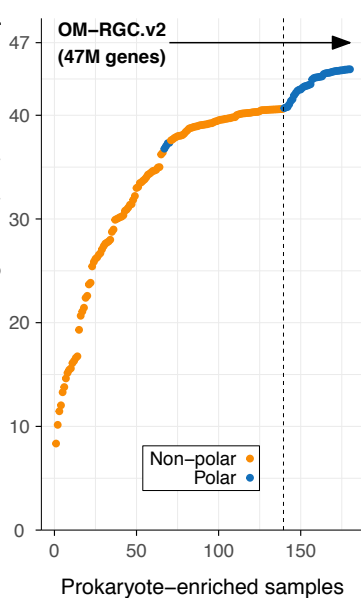


A.

Percentage of reads aligned (%)



B.

Number of genes (10^6)

C.

Percentage of genes

