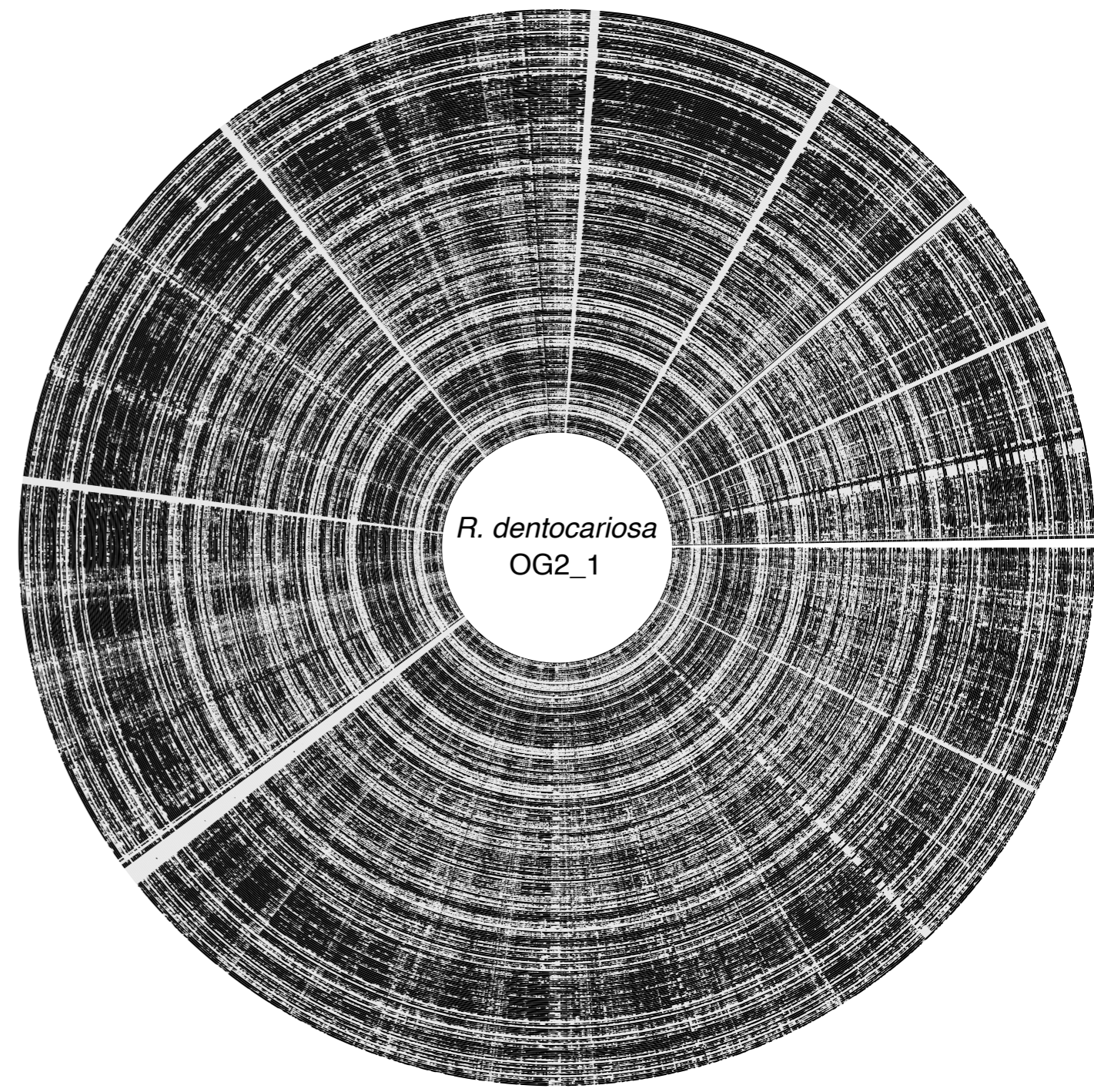
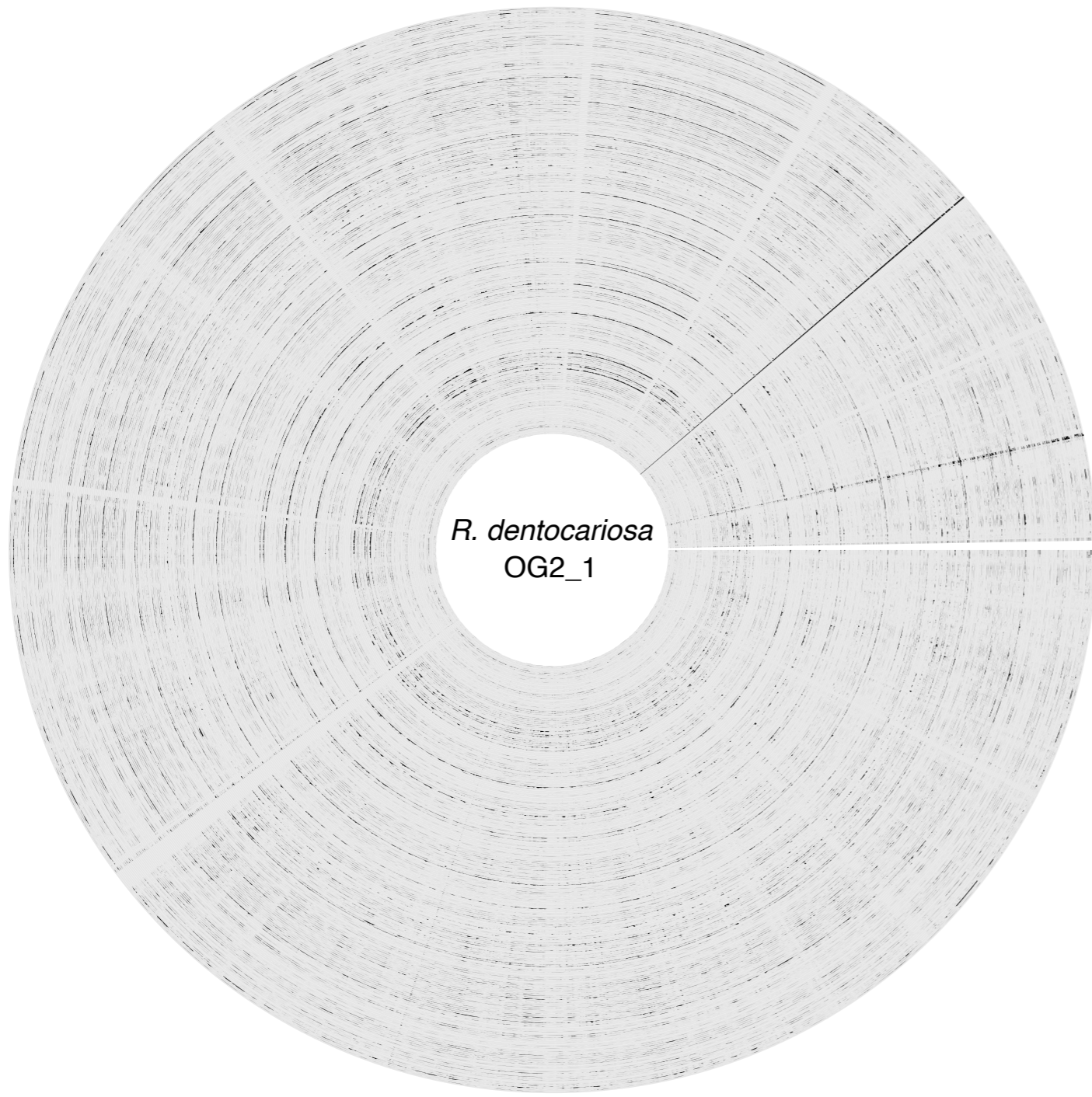


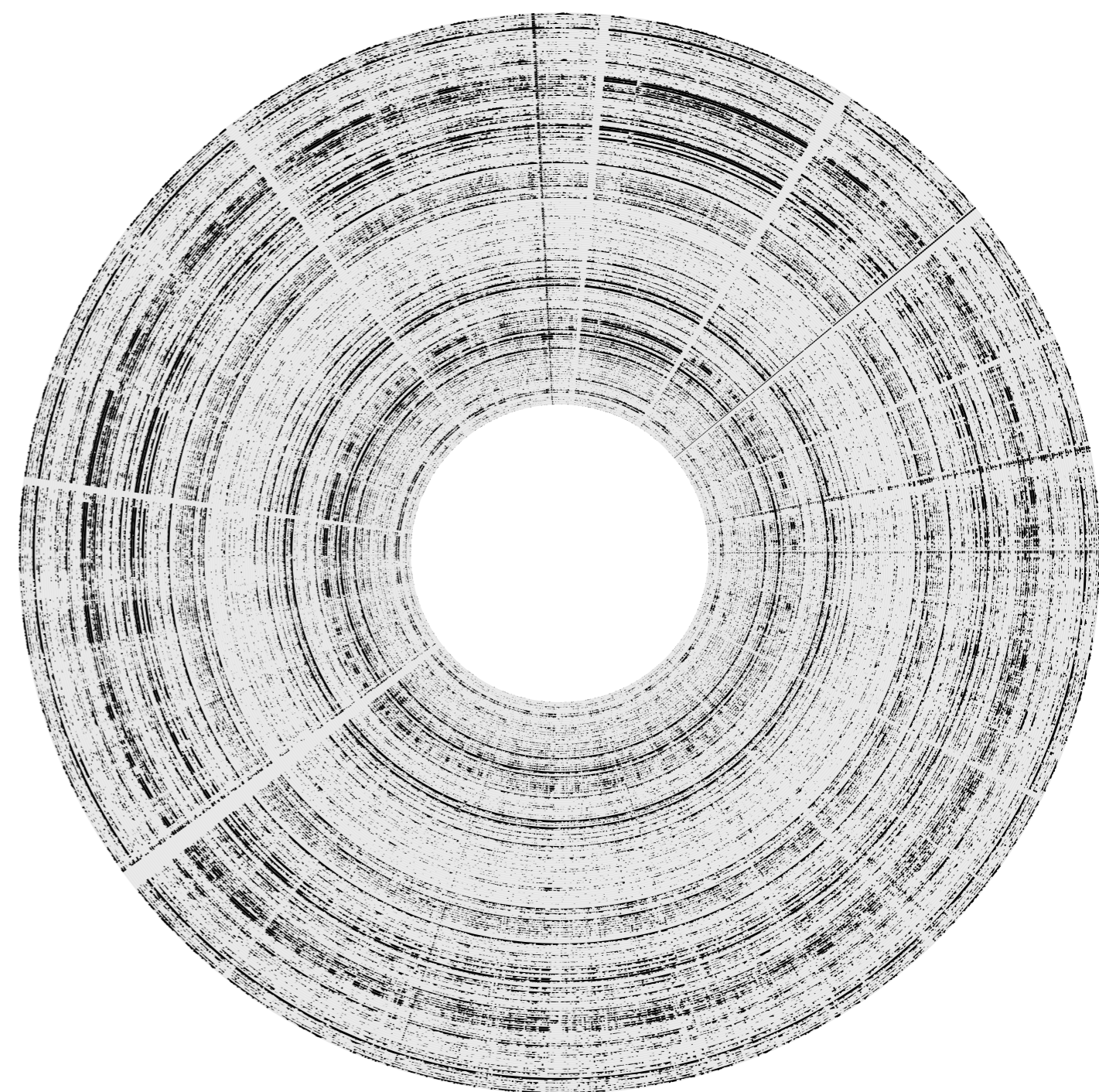
Coverage
(average number of reads per nt)

Detection
(fraction of each gene covered at all)

SUPP



BM



TD

