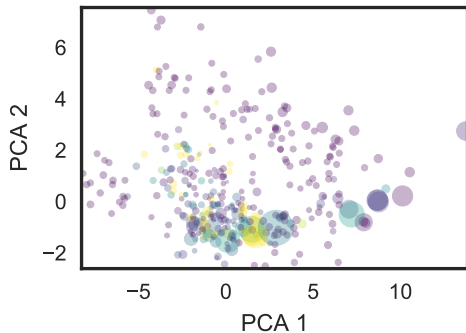
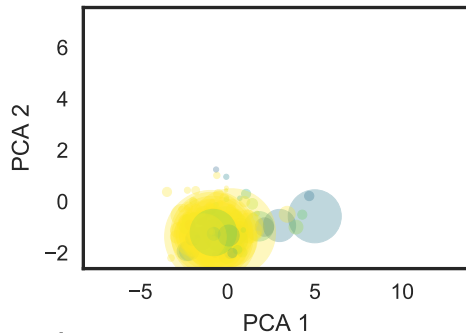


BACL7 - A11

a SAG Contigs with < 5% aligned to MAG



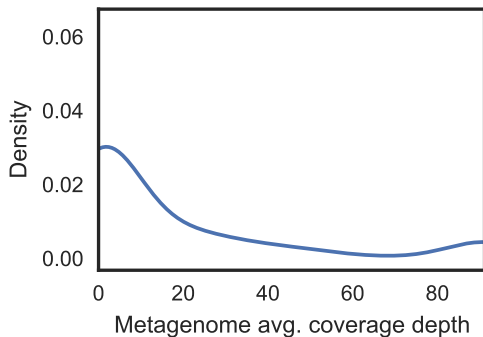
b SAG Contigs with > 5% aligned to MAG



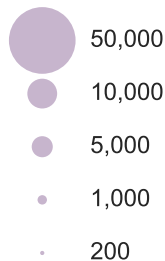
Metagenome avg. coverage depth



c



Contig lengths



d

