

File Name: Supplementary Data 1

Description: Long read metagenomic sequencing samples on human feces

File Name: Supplementary Data 2

Description: List of redundant genomes

File Name: Supplementary Data 3

Description: List of circular contigs by filtering process

File Name: Supplementary Data 4

Description: Final 102 HiFi cMAGs and their information

File Name: Supplementary Data 5

Description: List of taxa without isolated genome

File Name: Supplementary Data 6

Description: List of taxa that are assembled only from KR001 sample

File Name: Supplementary Data 7

Description: Quality assessment of 102 HiFi cMAGs by CheckM v1 and v2