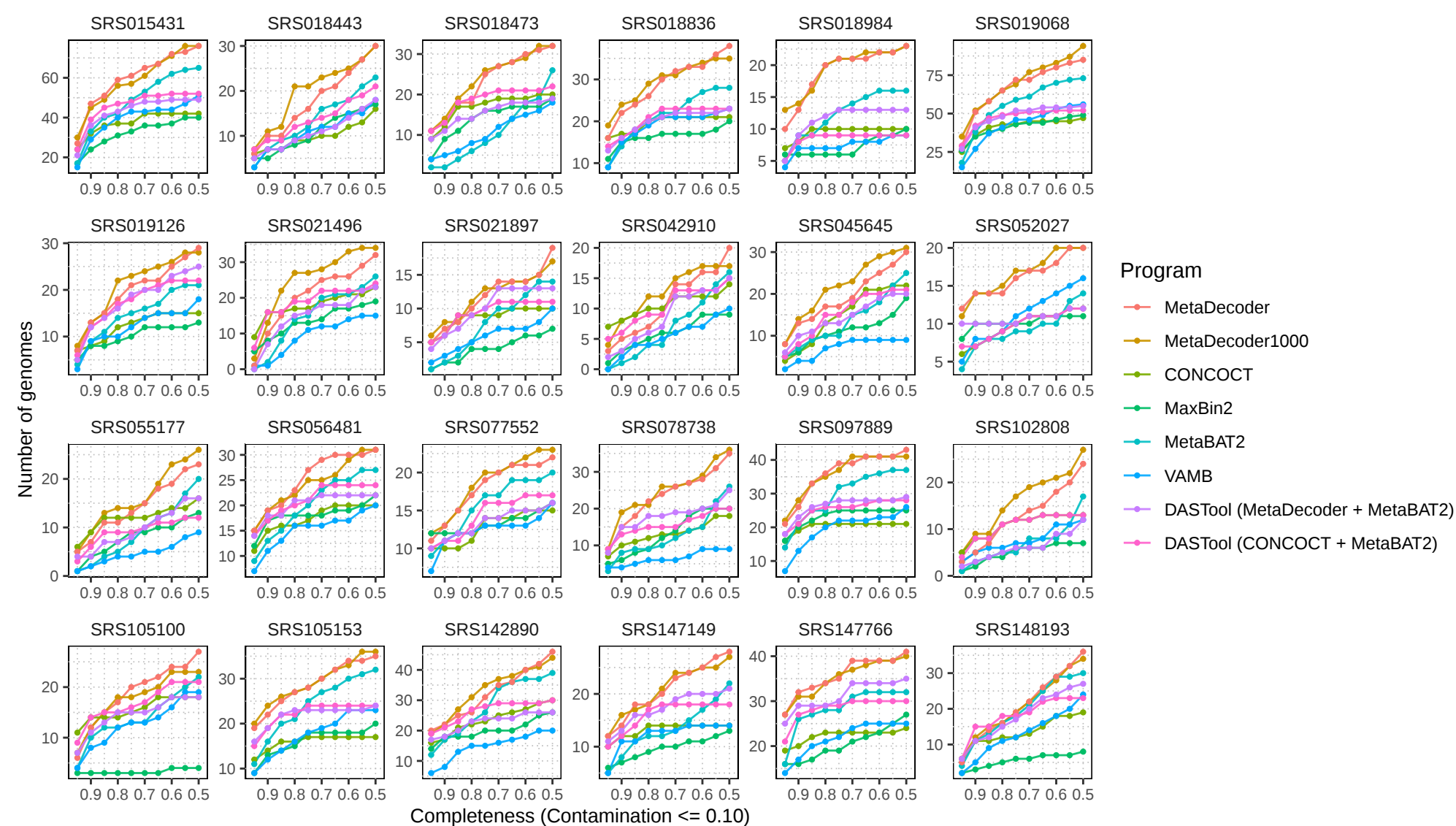




Supplementary Figure S7. Clustering benchmarks on 24 Human Microbiome Project datasets. The number of identified clusters with contamination ≤ 0.10 and different recall levels were shown. All programs were run with their default parameters. MetaDecoder with minimum sequence length setting to 1 Kb (MetaDecoder1000) was also added for benchmarking. Assessments were evaluated using CheckM (version 1.0.13).