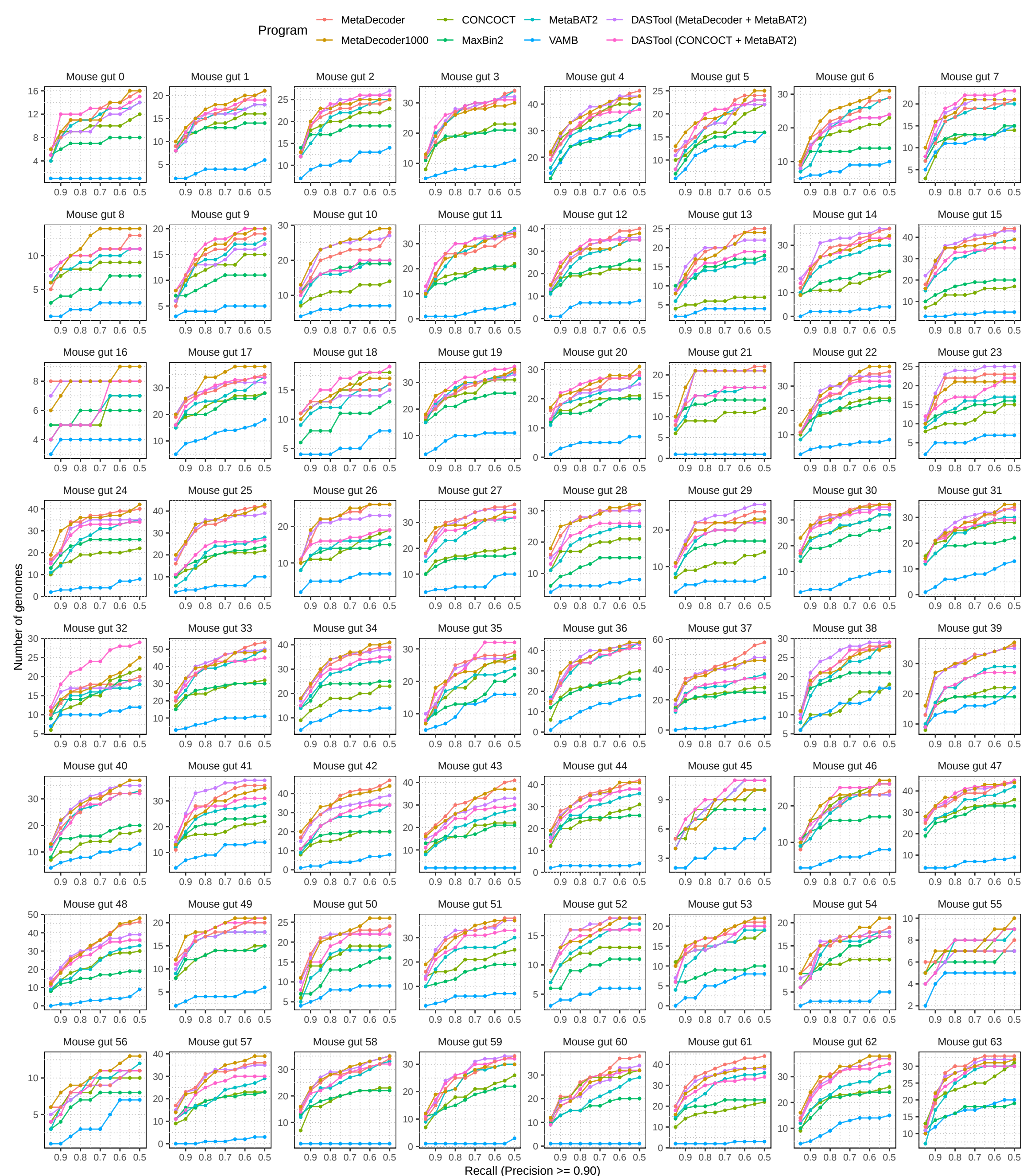




Supplementary Figure S5. Clustering benchmarks on 64 two CAMI II Mouse gut datasets. The number of identified clusters with precision ≥ 0.90 and different recall levels were shown. All programs were run with their default parameters. MetaDecoder with minimum sequence length setting to 1 Kb (MetaDecoder1000) was also added for benchmarking. Assessments were evaluated using AMBER (version 2.0.2).