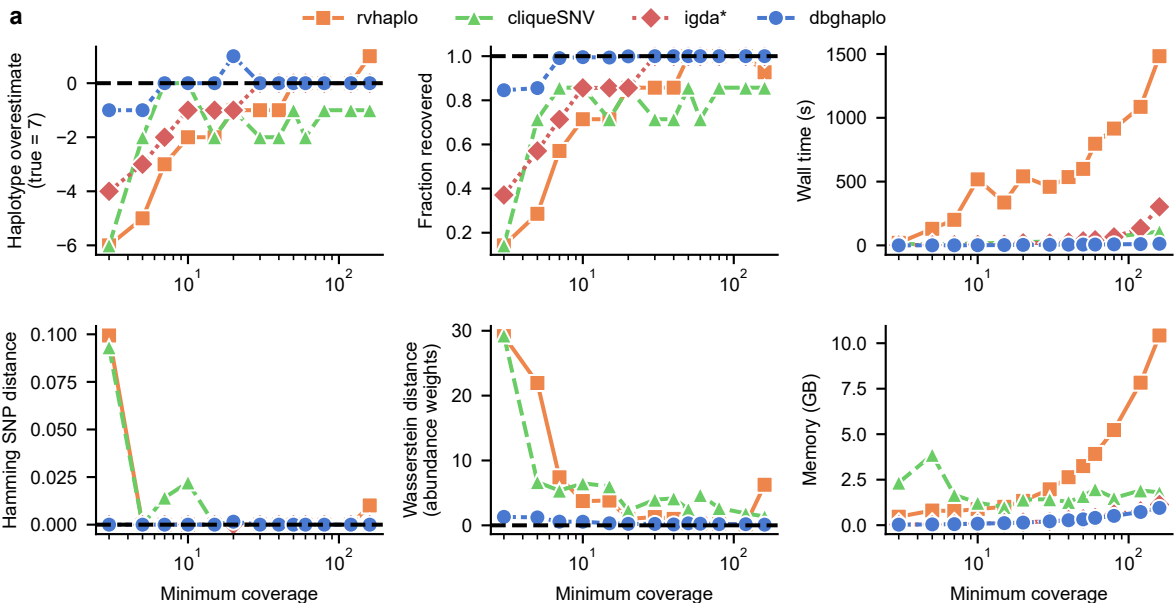


7 HIV genomes (1:3:5:7:9:20 ratio)



2-30 HIV genomes (15-145x coverage)

