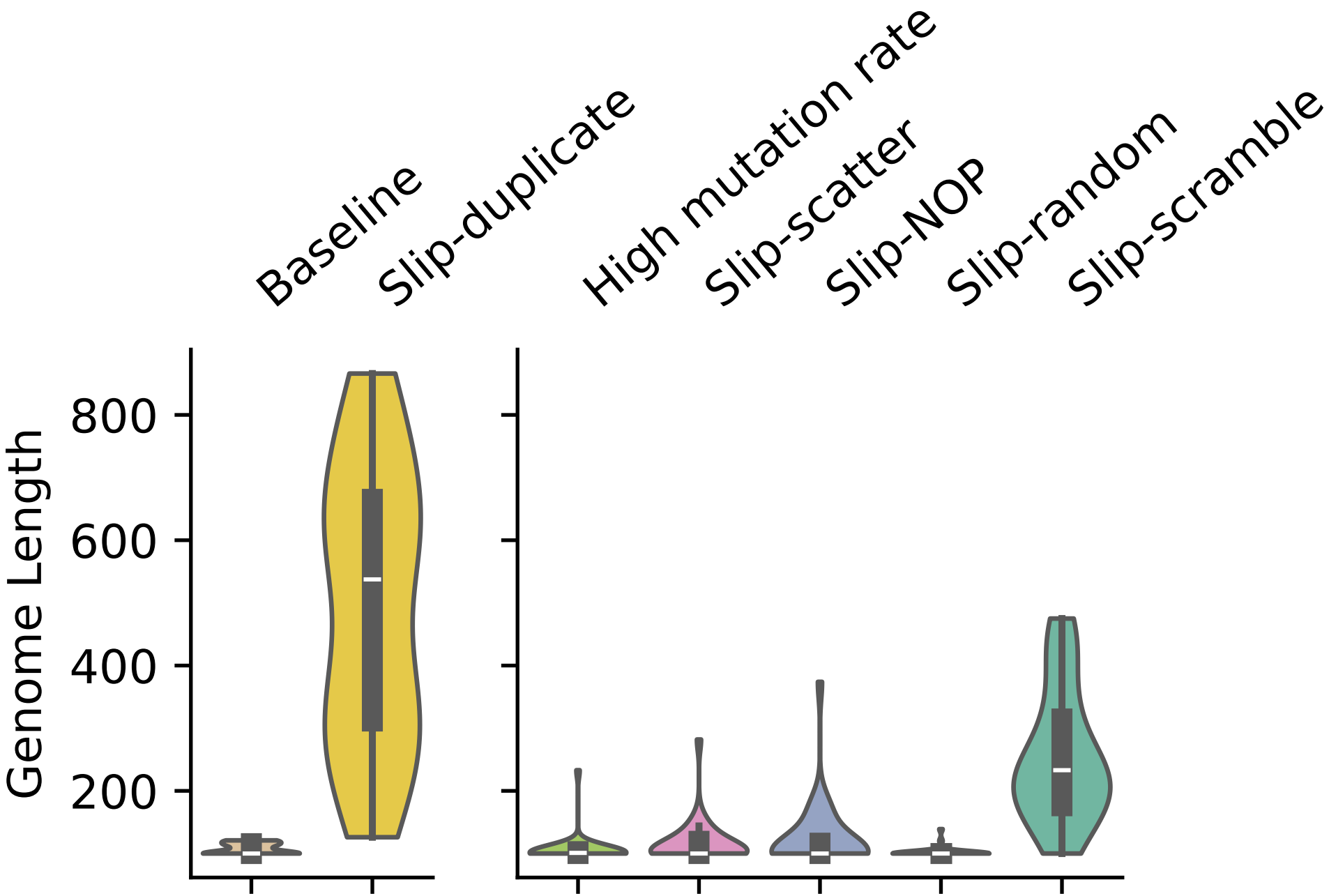
Baseline
Slip-duplicate
High mutation rate
Slip-scatter
Slip-NOP
Slip-random
Slip-scramble
Genome Length
800
600
400
200