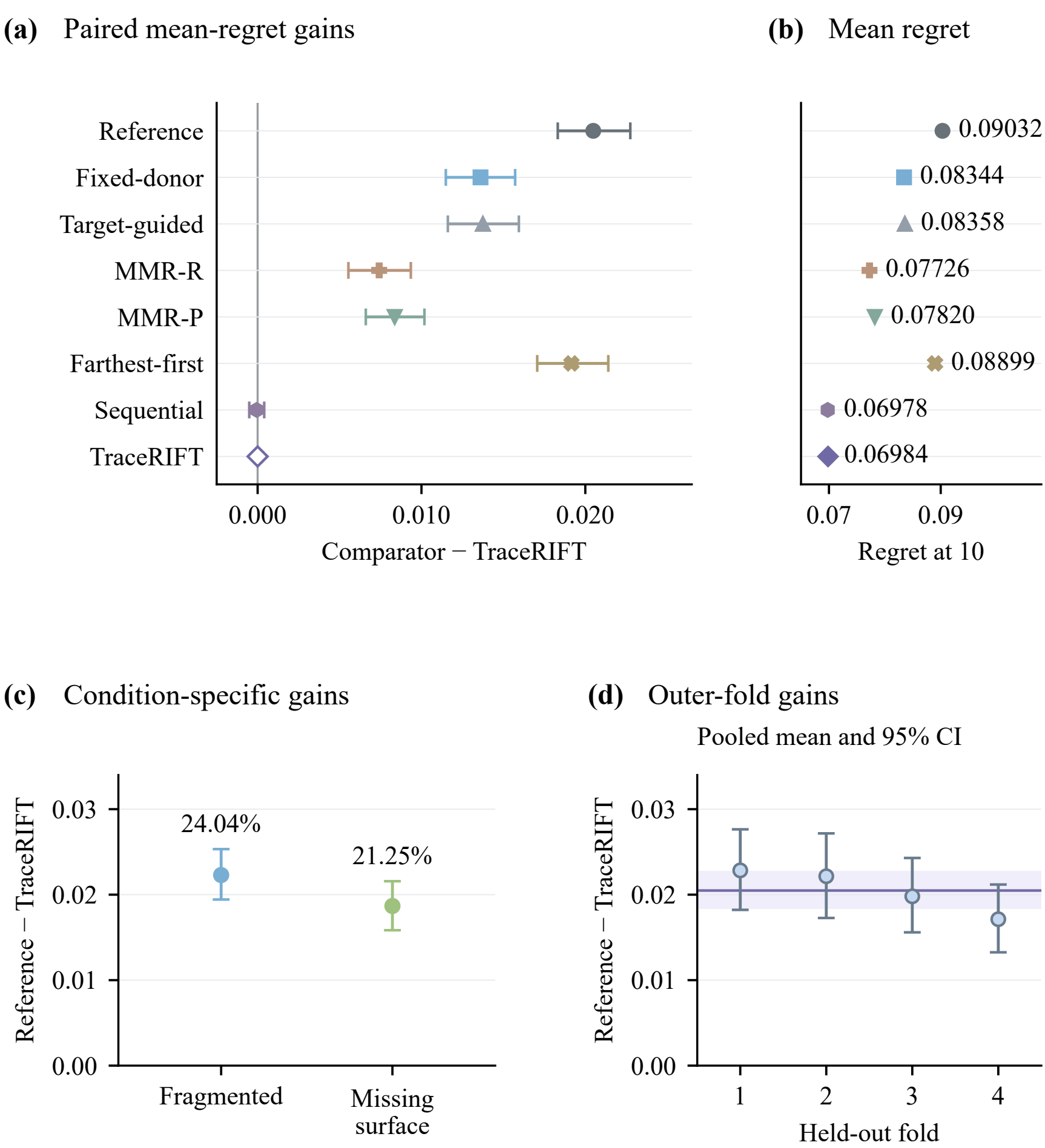

(a) Paired mean-regret gains
Reference
Fixed-donor
Target-guided
MMR-R
MMR-P
Farthest-first
Sequential
TraceRIFT
0.000
0.010
0.020
Comparator − TraceRIFT
(b) Mean regret
0.09032
0.08344
0.08358
0.07726
0.07820
0.08899
0.06978
0.06984
0.07
0.09
Regret at 10
(c) Condition-specific gains
Reference − TraceRIFT
0.03
0.02
0.01
0.00
24.04%
21.25%
Fragmented
Missing surface
(d) Outer-fold gains
Pooled mean and 95% CI
Reference − TraceRIFT
0.03
0.02
0.01
0.00
1
2
3
4
Held-out fold