

A

Complexity (# genomes)
low = 30 medium = 100 high = 300

DONOR
genomes

RECIPIENT
genomes
before FMT

UNKNOWN
SOURCE
genomes

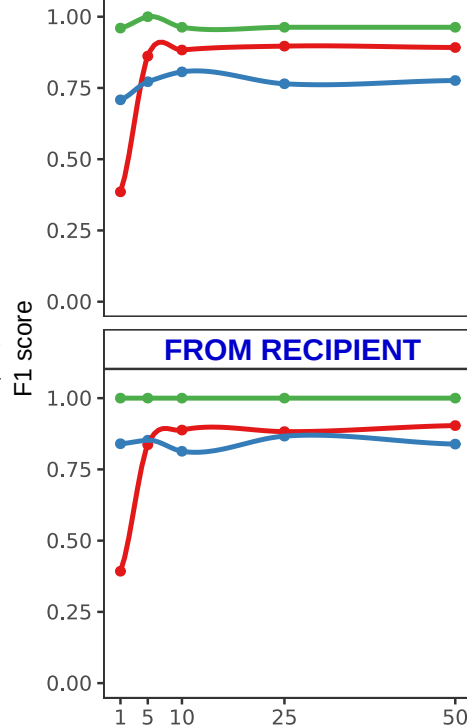
SUBSET

RECIPIENT
genomes
after FMT

B

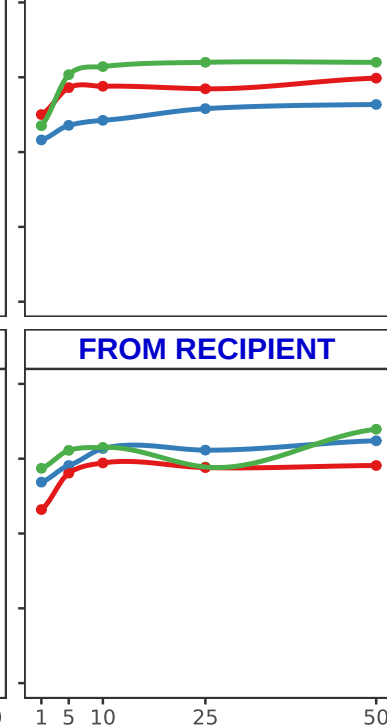
SPECIES

FROM DONOR



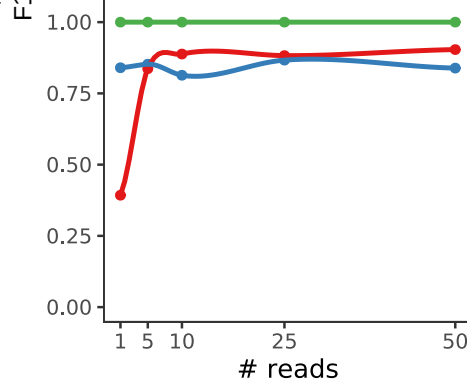
STRAINS

FROM DONOR

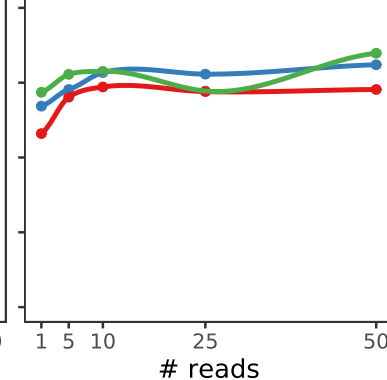


C

FROM RECIPIENT



FROM RECIPIENT



complexity

high

medium

low