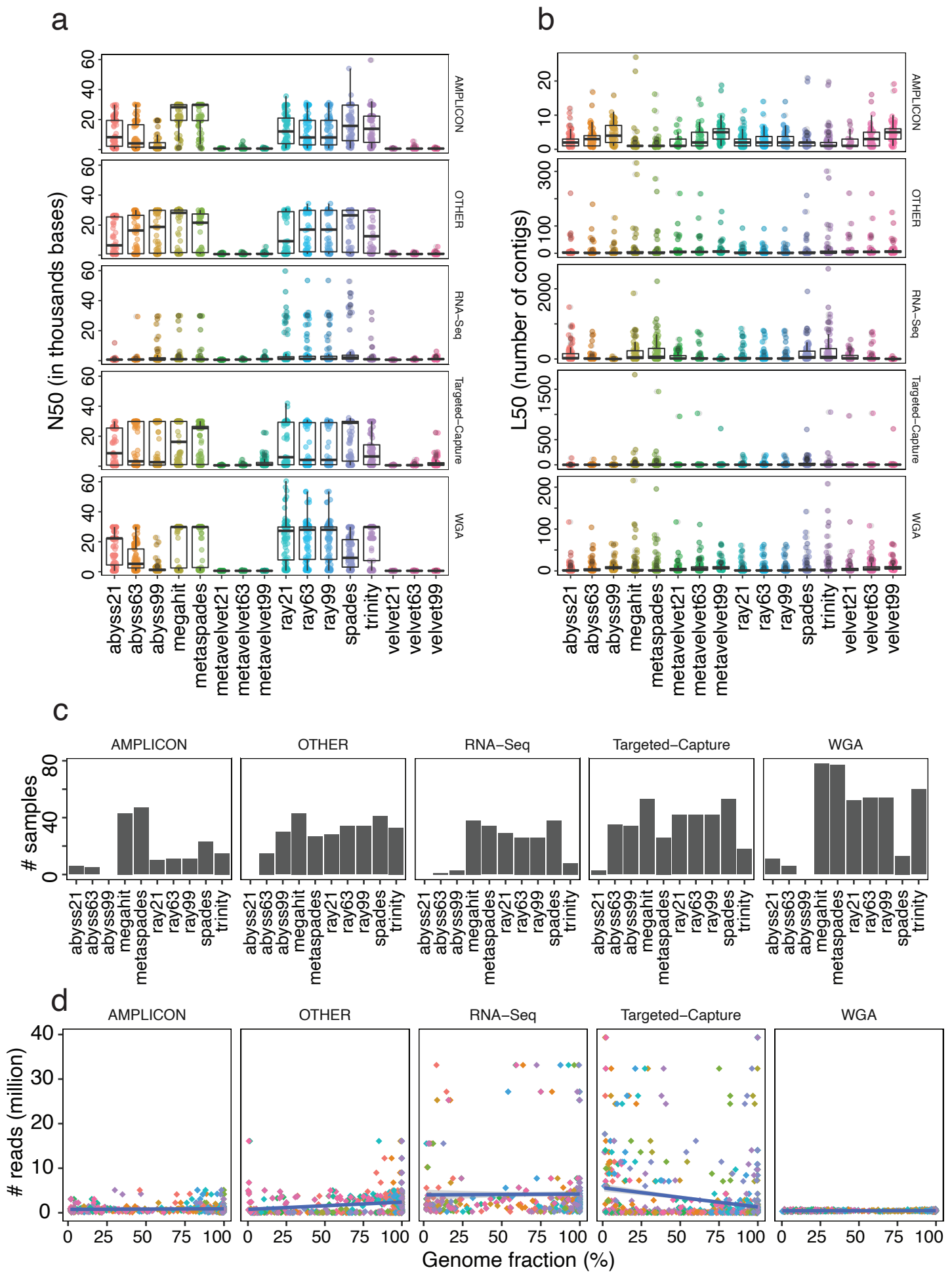
a
b
c
d
N50 (in thousands bases)
L50 (number of contigs)
AMPLICON
OTHER
RNA−Seq
Targeted−Capture
WGA
abyss21
abyss63
abyss99
megahit
metaspades
metavelvet21
metavelvet63
metavelvet99
ray21
ray63
ray99
spades
trinity
velvet21
velvet63
velvet99
samples
reads (million)
Genome fraction (%)