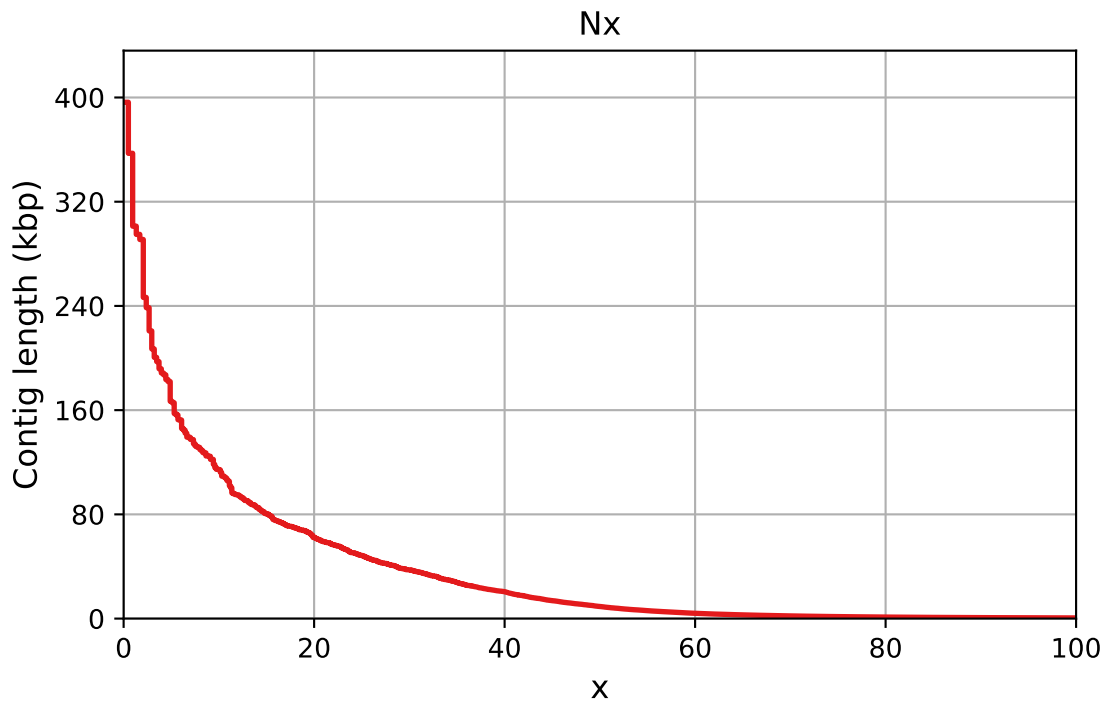


Report

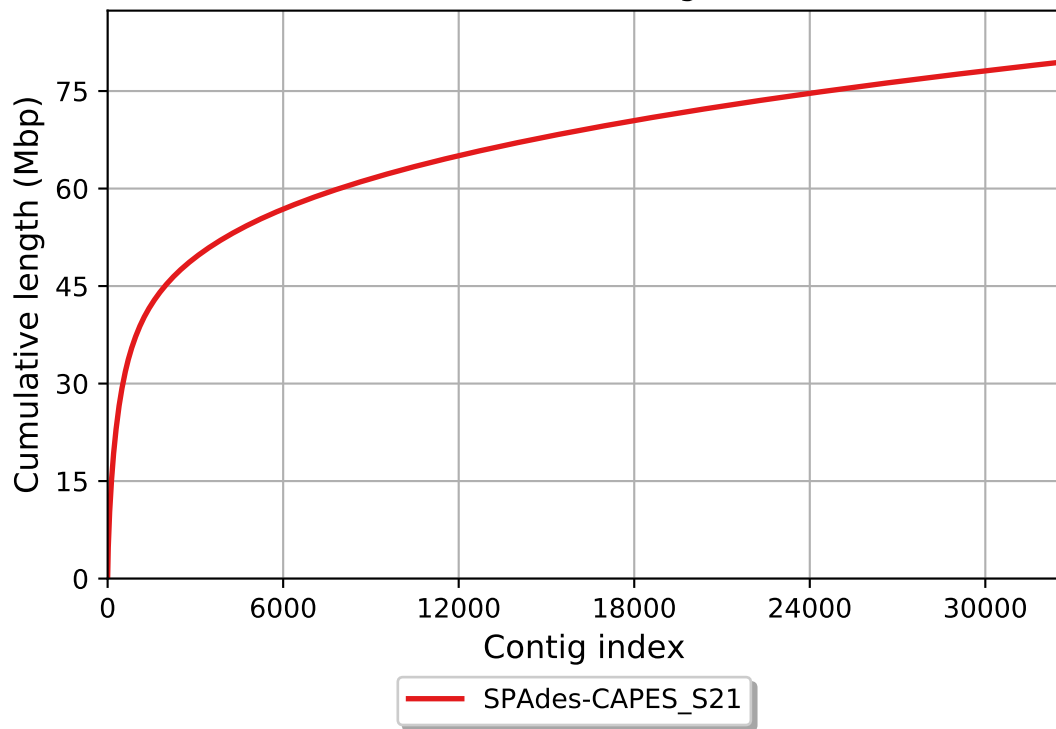
	SPAdes-CAPE_S21
# contigs (>= 0 bp)	111855
# contigs (>= 1000 bp)	12879
# contigs (>= 5000 bp)	2100
# contigs (>= 10000 bp)	1127
# contigs (>= 25000 bp)	481
# contigs (>= 50000 bp)	207
Total length (>= 0 bp)	102929034
Total length (>= 1000 bp)	65962649
Total length (>= 5000 bp)	45726625
Total length (>= 10000 bp)	39086584
Total length (>= 25000 bp)	29043781
Total length (>= 50000 bp)	19377562
# contigs	32556
Largest contig	396331
Total length	79427604
GC (%)	45.84
N50	9283
N75	1518
L50	1193
L75	7629
# N's per 100 kbp	7.72
# predicted rRNA genes	19 + 13 part

All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

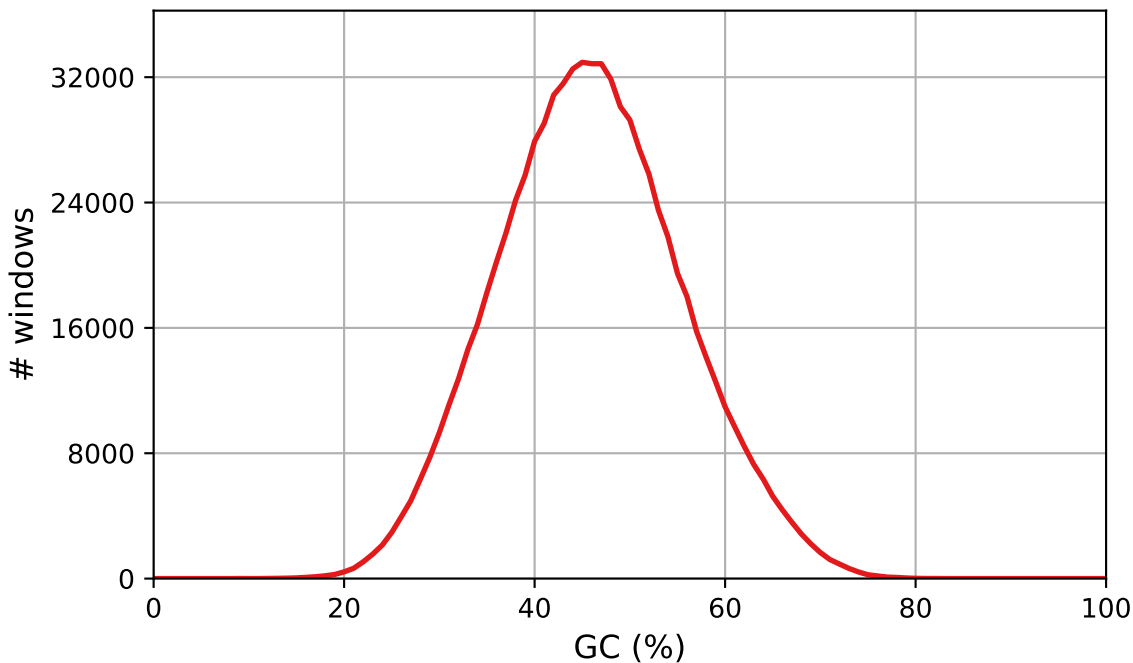


— SPAdes-CAPES_S21

Cumulative length

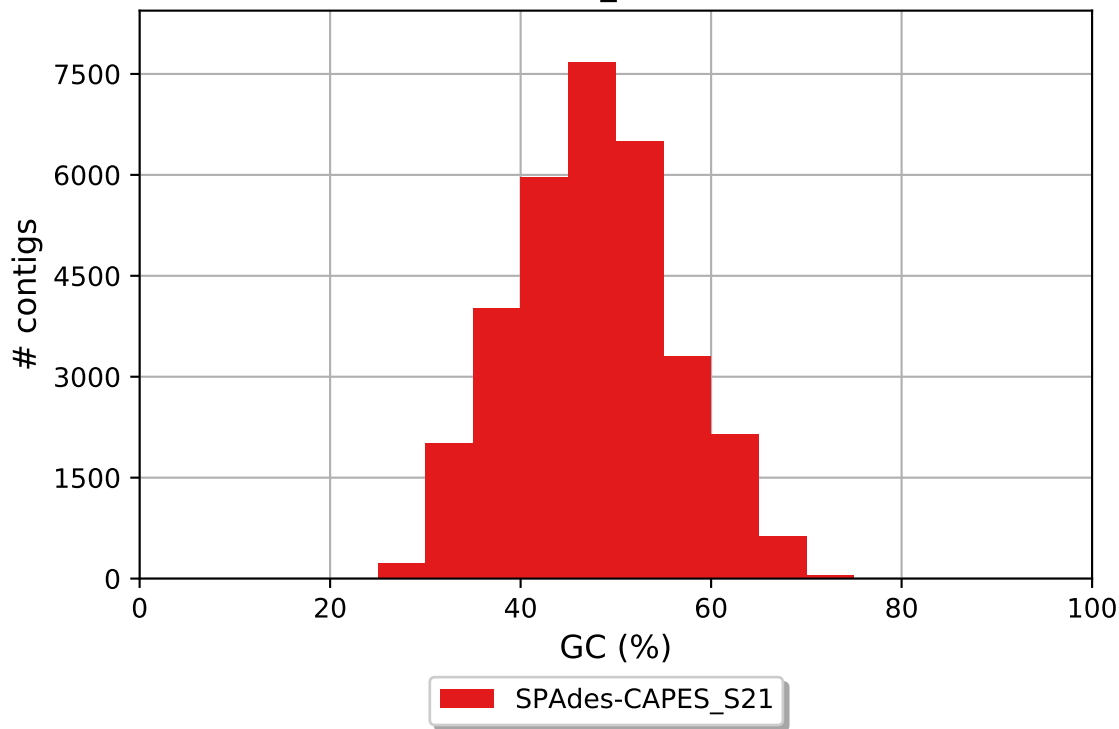


GC content

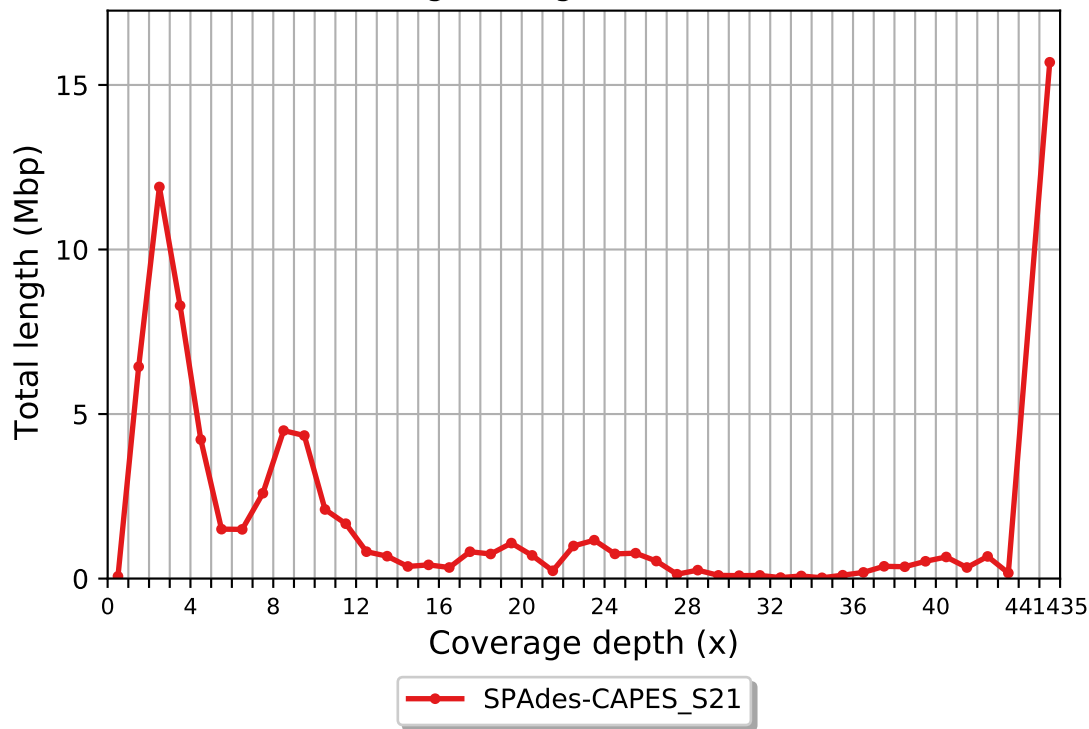


— SPAdes-CAPES_S21

SPAdes-CAPES_S21 GC content



Coverage histogram (bin size: 1x)



SPAdes-CAPES_S21 coverage histogram (bin size: 1x)

