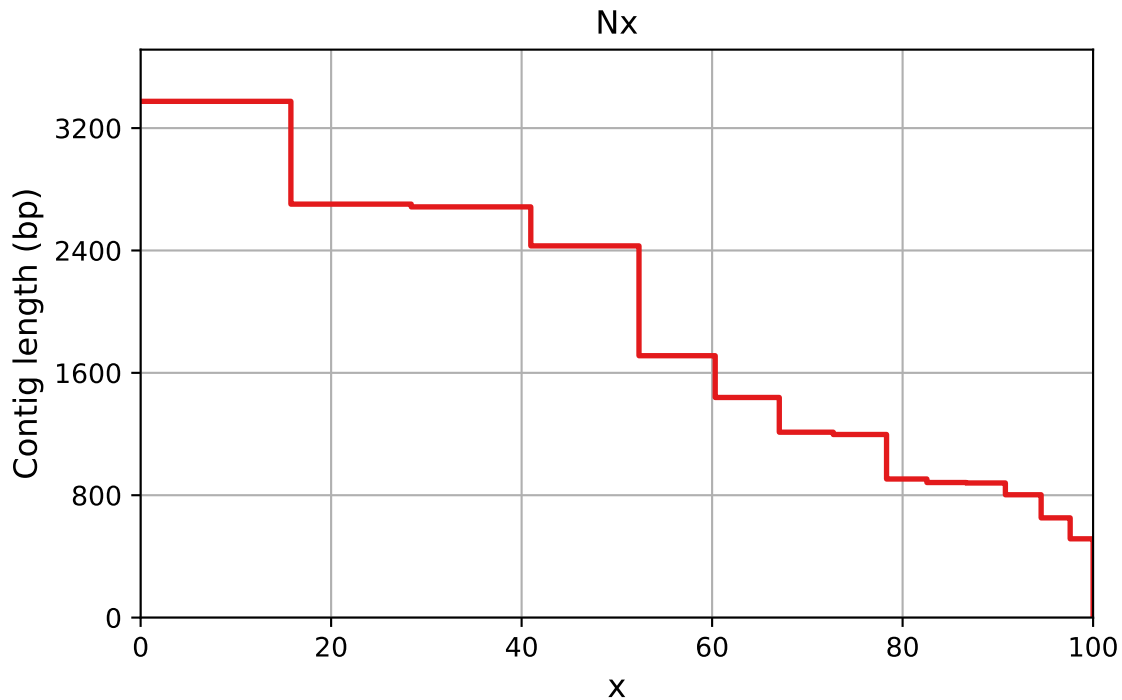


## Report

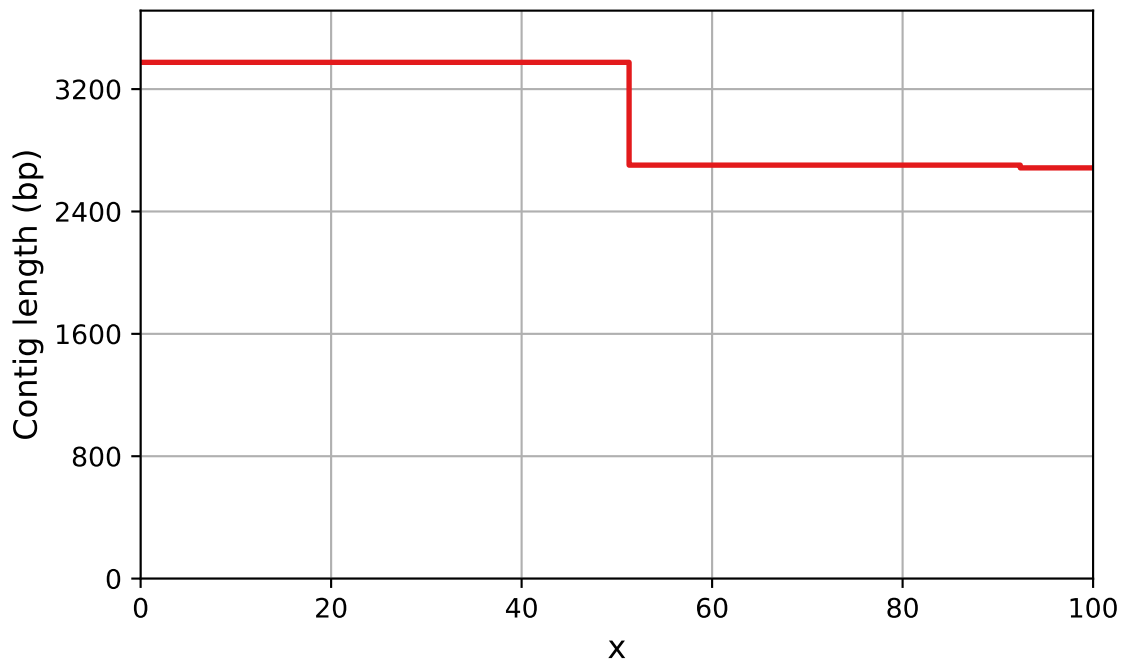
	PTA110.scaffolds
# contigs (>= 0 bp)	28
# contigs (>= 1000 bp)	8
# contigs (>= 5000 bp)	0
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	25401
Total length (>= 1000 bp)	16753
Total length (>= 5000 bp)	0
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	14
Largest contig	3375
Total length	21392
Reference length	6581
GC (%)	34.24
Reference GC (%)	41.79
N50	2430
NG50	3375
N75	1197
NG75	2703
L50	4
LG50	1
L75	8
LG75	2
# unaligned contigs	14 + 0 part
Unaligned length	21392
# N's per 100 kbp	0.00
NGA50	-

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).



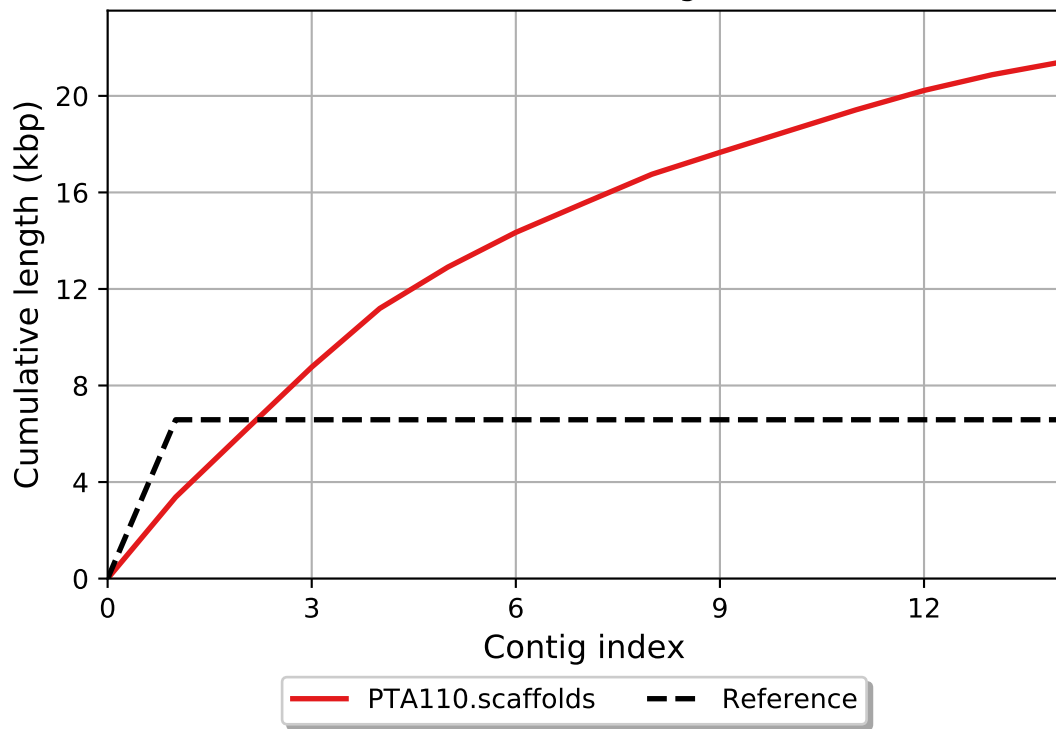
— PTA110.scaffolds

NGx

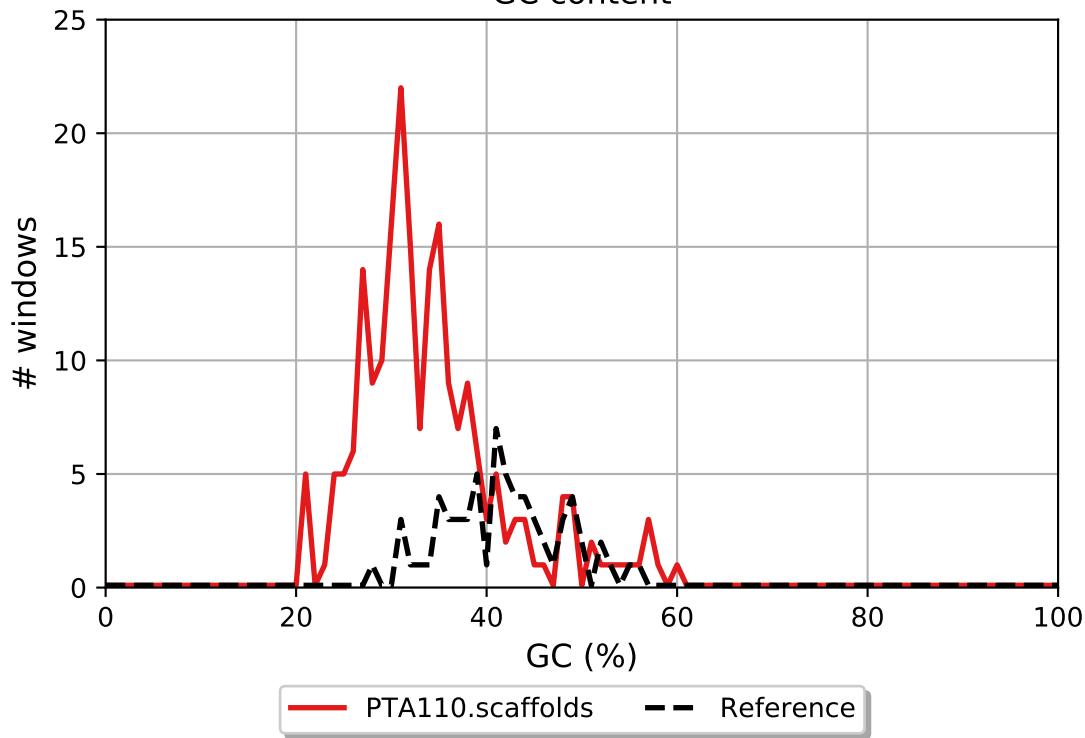


PTA110.scaffolds

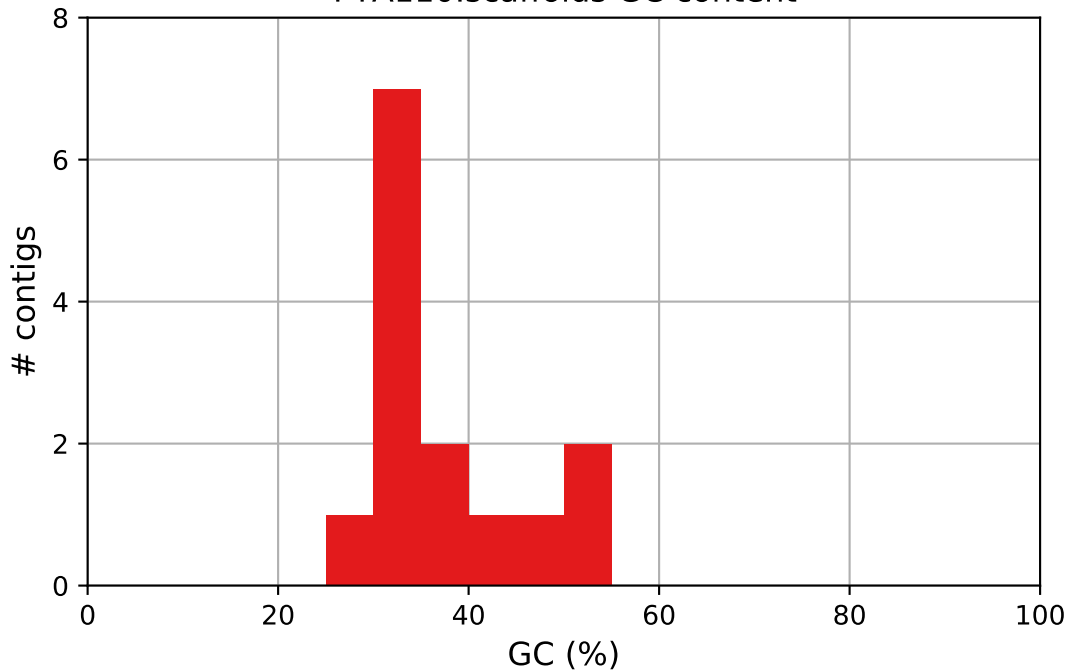
Cumulative length



# GC content

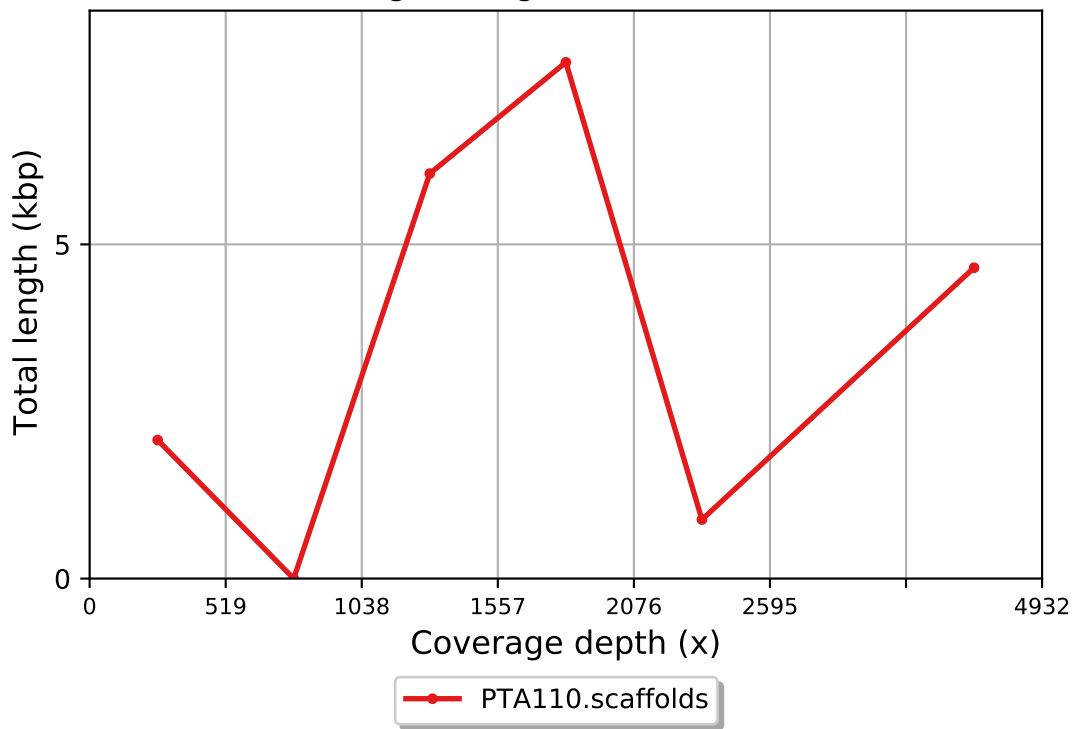


PTA110.scaffolds GC content



PTA110.scaffolds

Coverage histogram (bin size: 519x)



PTA110.scaffolds coverage histogram (bin size: 519x)

