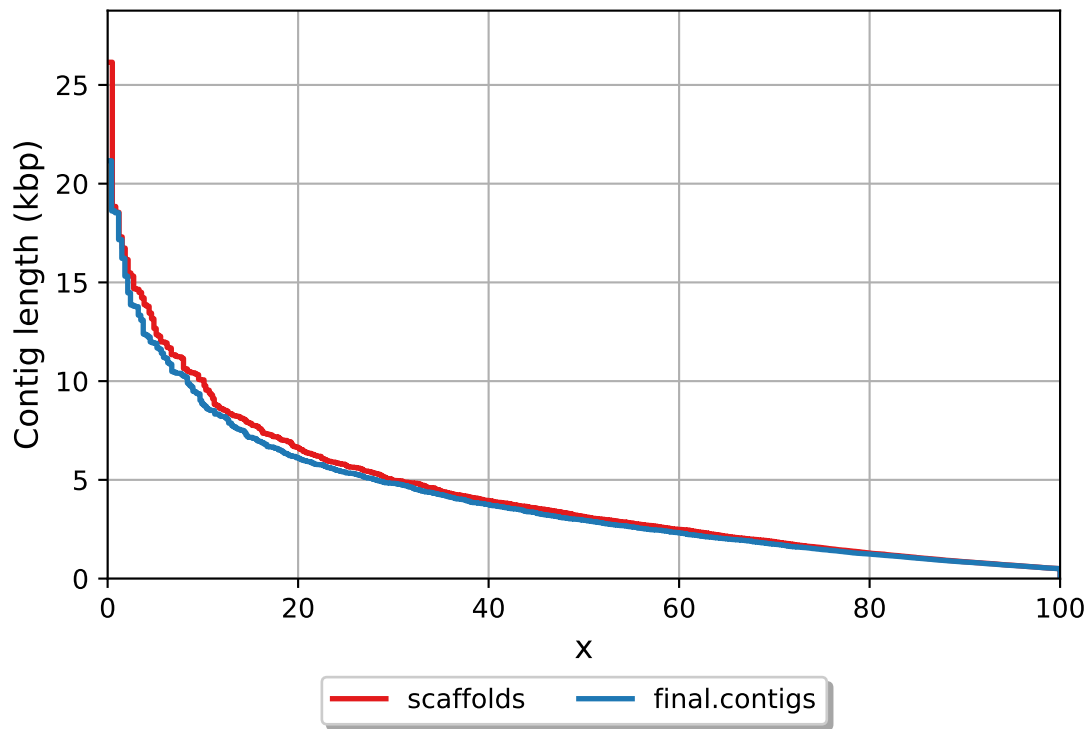


# Report

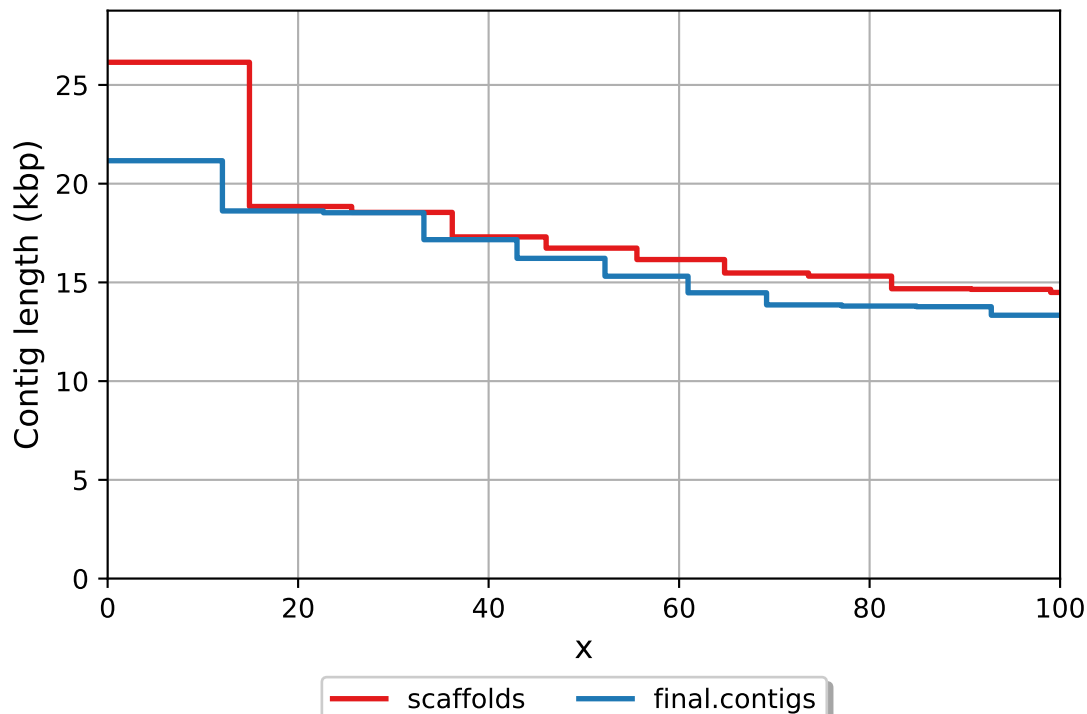
	scaffolds	final.contigs
# contigs (>= 0 bp)	5021	3486
# contigs (>= 1000 bp)	1543	1535
# contigs (>= 5000 bp)	198	186
# contigs (>= 10000 bp)	41	33
# contigs (>= 25000 bp)	1	0
# contigs (>= 50000 bp)	0	0
Total length (>= 0 bp)	5925121	5427303
Total length (>= 1000 bp)	4569336	4344915
Total length (>= 5000 bp)	1574924	1421602
Total length (>= 10000 bp)	534932	423843
Total length (>= 25000 bp)	26148	0
Total length (>= 50000 bp)	0	0
# contigs	2549	2543
Largest contig	26148	21162
Total length	5284622	5061808
Reference length	175592	175592
GC (%)	42.27	42.39
Reference GC (%)	65.98	65.98
N50	3149	2954
NG50	16735	16218
N75	1565	1472
NG75	15318	13860
L50	468	479
LG50	5	5
L75	1058	1083
LG75	8	8
# unaligned contigs	2549 + 0 part	2543 + 0 part
Unaligned length	5284622	5061808
# N's per 100 kbp	11.83	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).

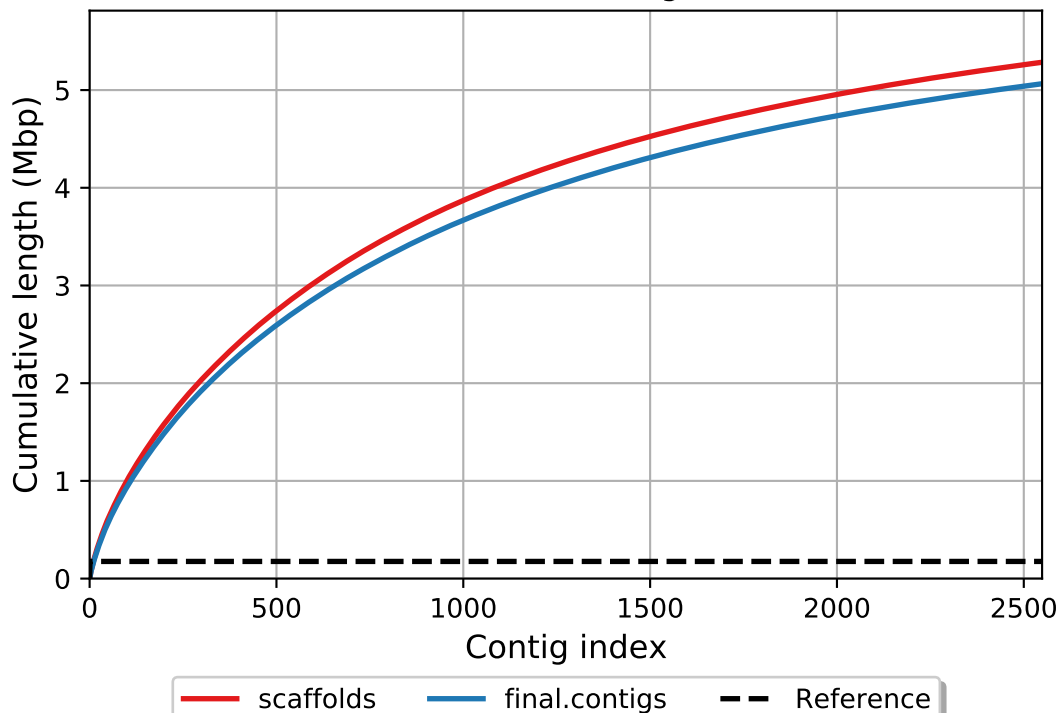
Nx



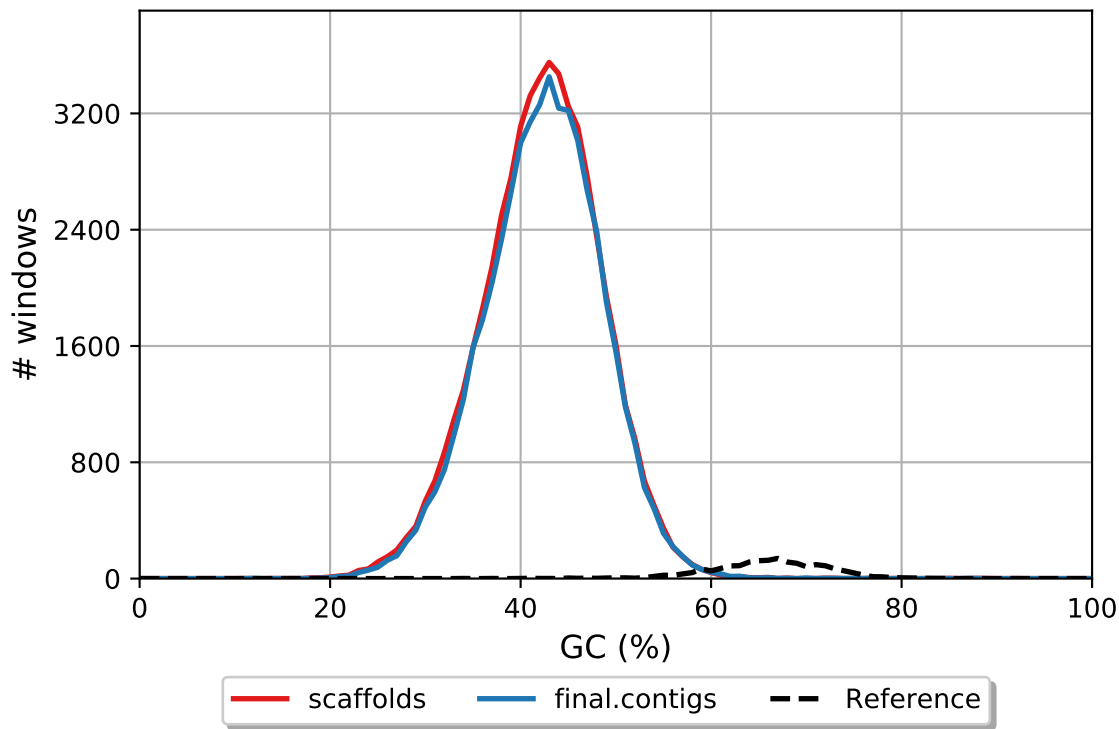
# NGx



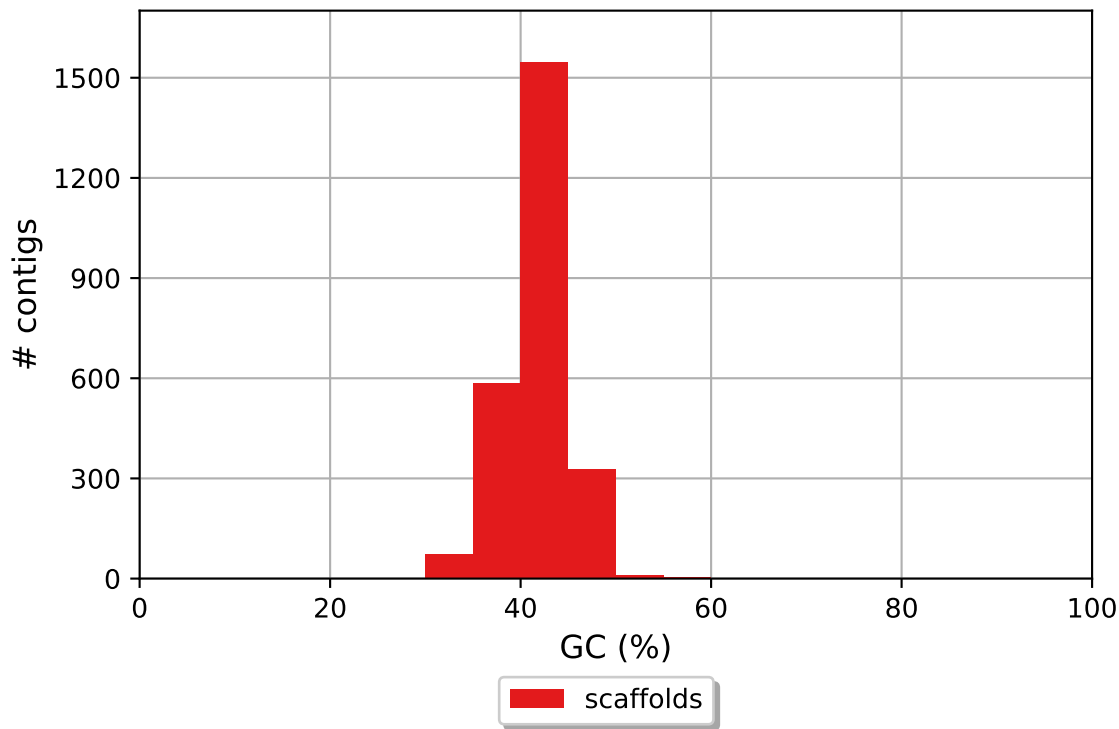
Cumulative length



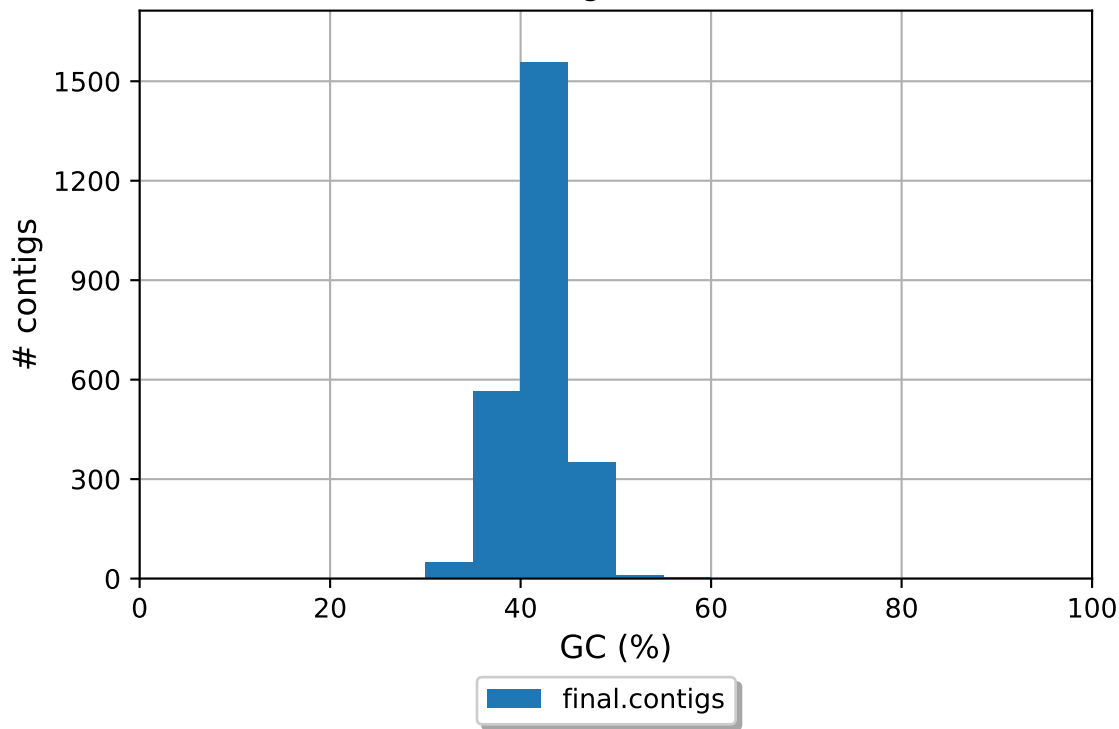
## GC content



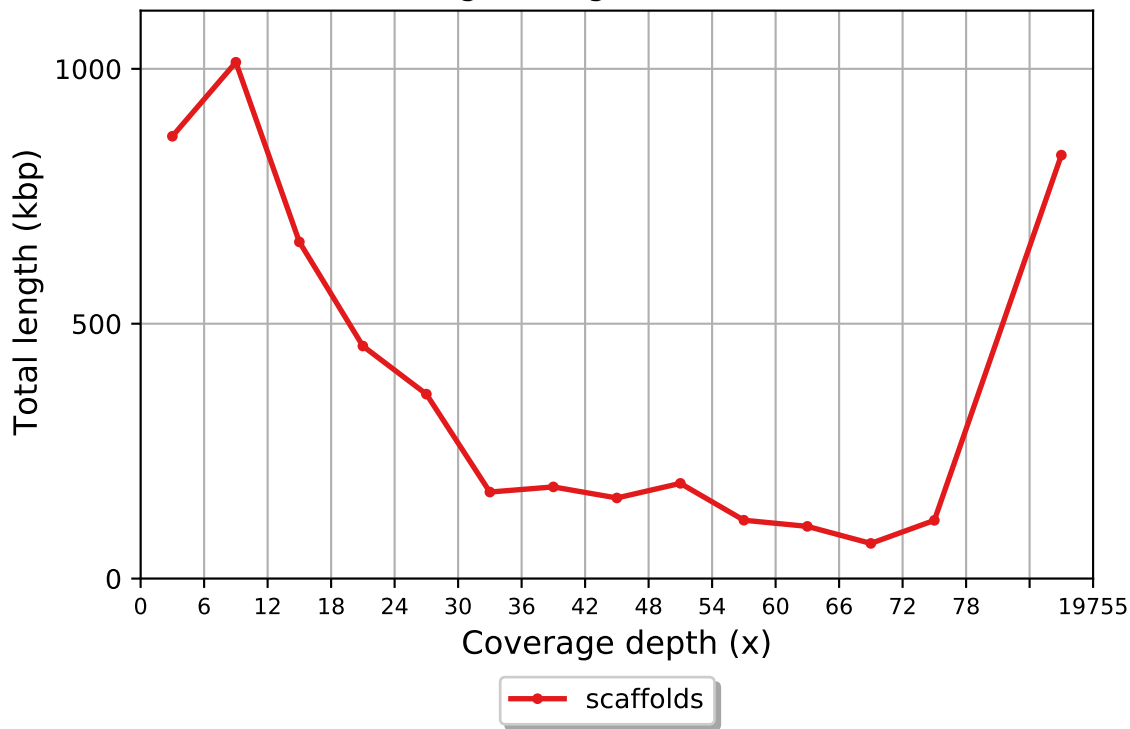
scaffolds GC content



final.contigs GC content



Coverage histogram (bin size: 6x)





scaffolds coverage histogram (bin size: 6x)

