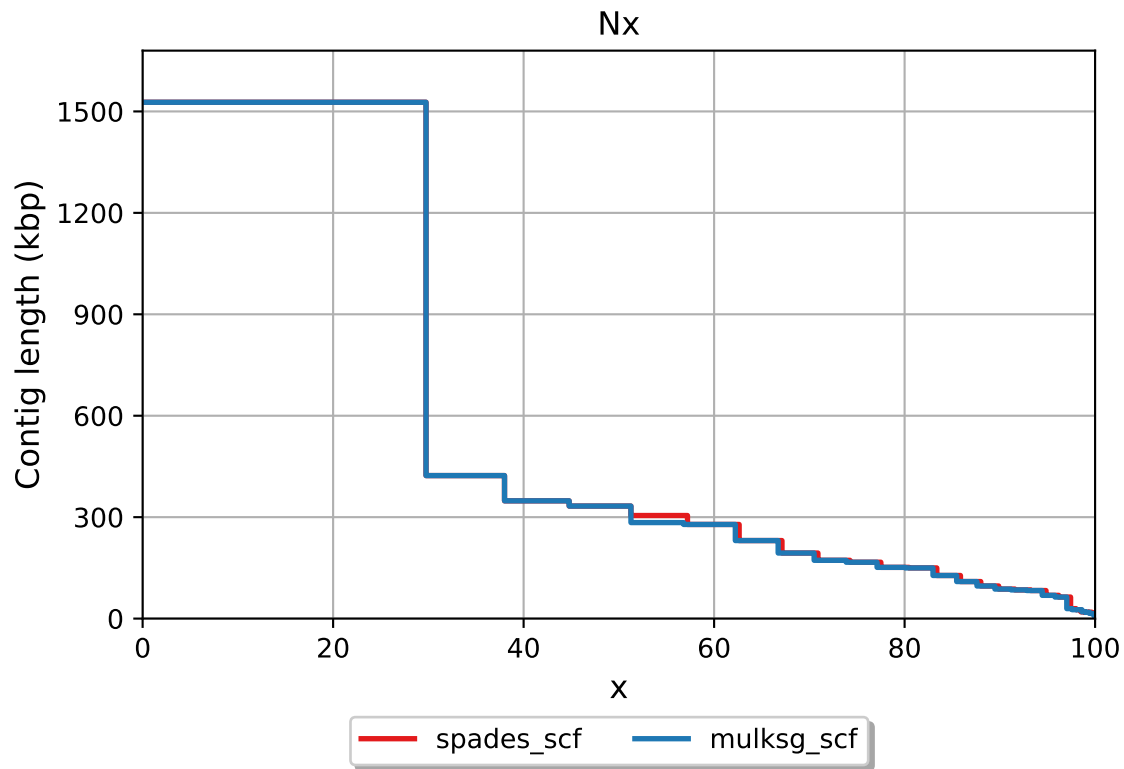


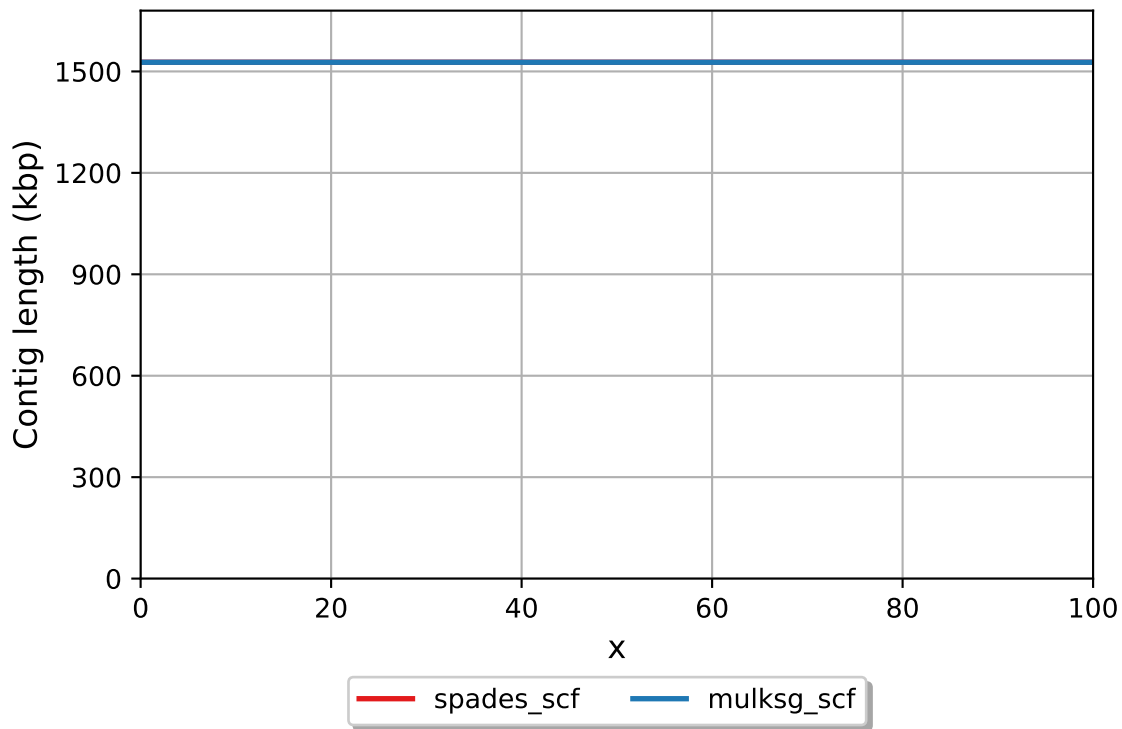
## Report

	spades_scf	mulksq_scf
# contigs (>= 0 bp)	59	62
# contigs (>= 1000 bp)	26	28
# contigs (>= 5000 bp)	26	27
# contigs (>= 10000 bp)	26	27
# contigs (>= 25000 bp)	22	23
# contigs (>= 50000 bp)	20	20
Total length (>= 0 bp)	5141098	5141577
Total length (>= 1000 bp)	5129627	5130798
Total length (>= 5000 bp)	5129627	5129407
Total length (>= 10000 bp)	5129627	5129407
Total length (>= 25000 bp)	5056142	5061454
Total length (>= 50000 bp)	5000464	4979570
# contigs	30	30
Largest contig	1527108	1527108
Total length	5132507	5132188
Reference length	23319	23319
GC (%)	64.11	64.11
Reference GC (%)	67.86	67.86
N50	332824	332816
NG50	1527108	1527108
N75	166687	166687
NG75	1527108	1527108
L50	4	4
LG50	1	1
L75	10	10
LG75	1	1
# unaligned contigs	30 + 0 part	30 + 0 part
Unaligned length	5132507	5132188
# N's per 100 kbp	0.00	0.00
NGA50	-	-

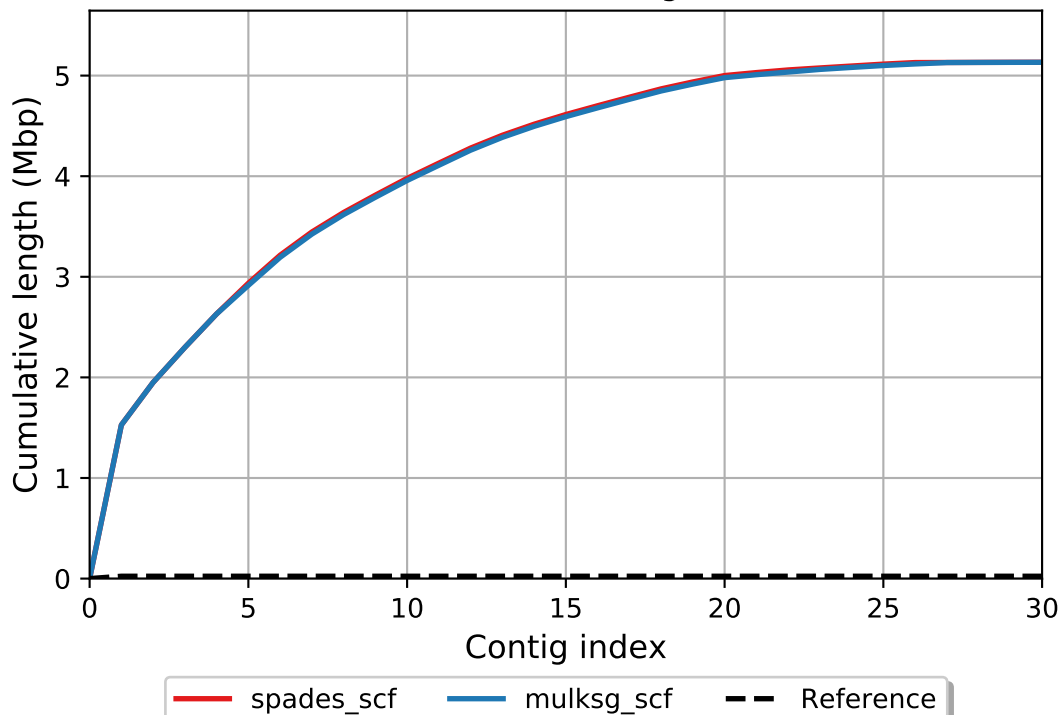
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



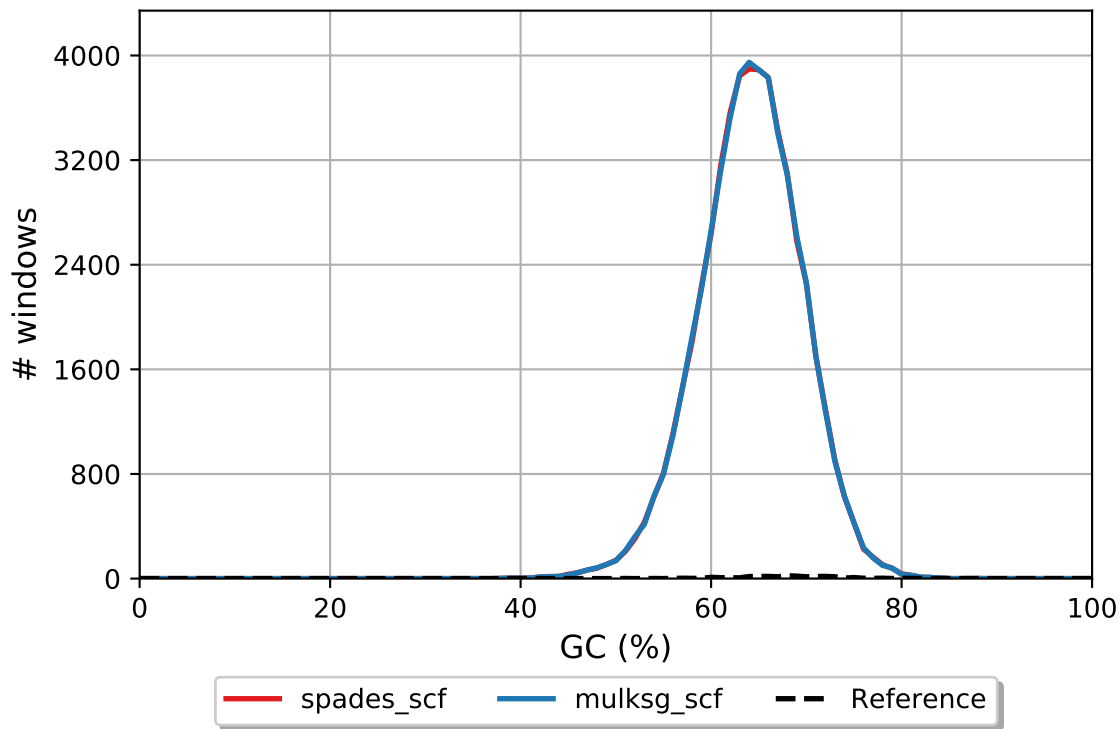
NGx



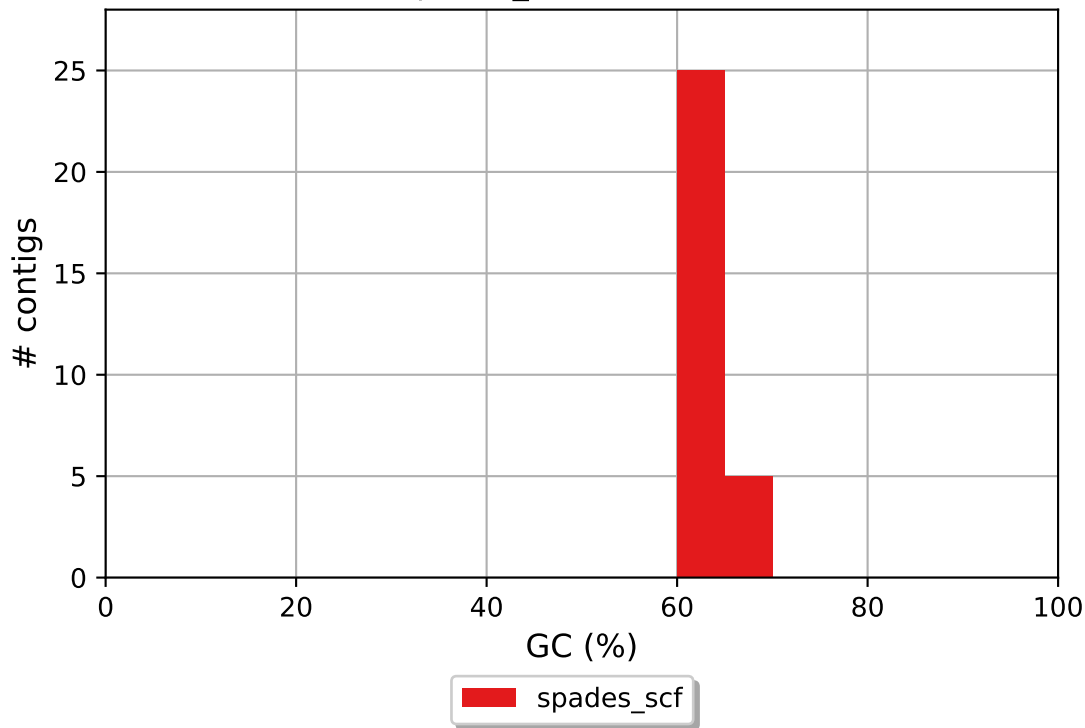
Cumulative length



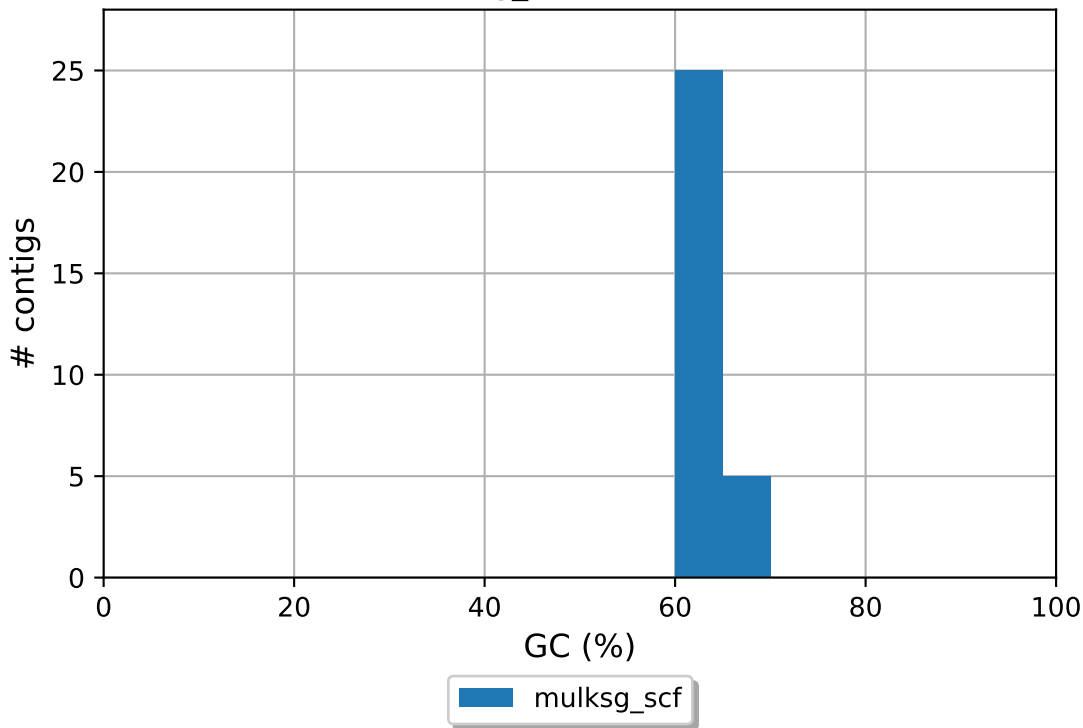
GC content



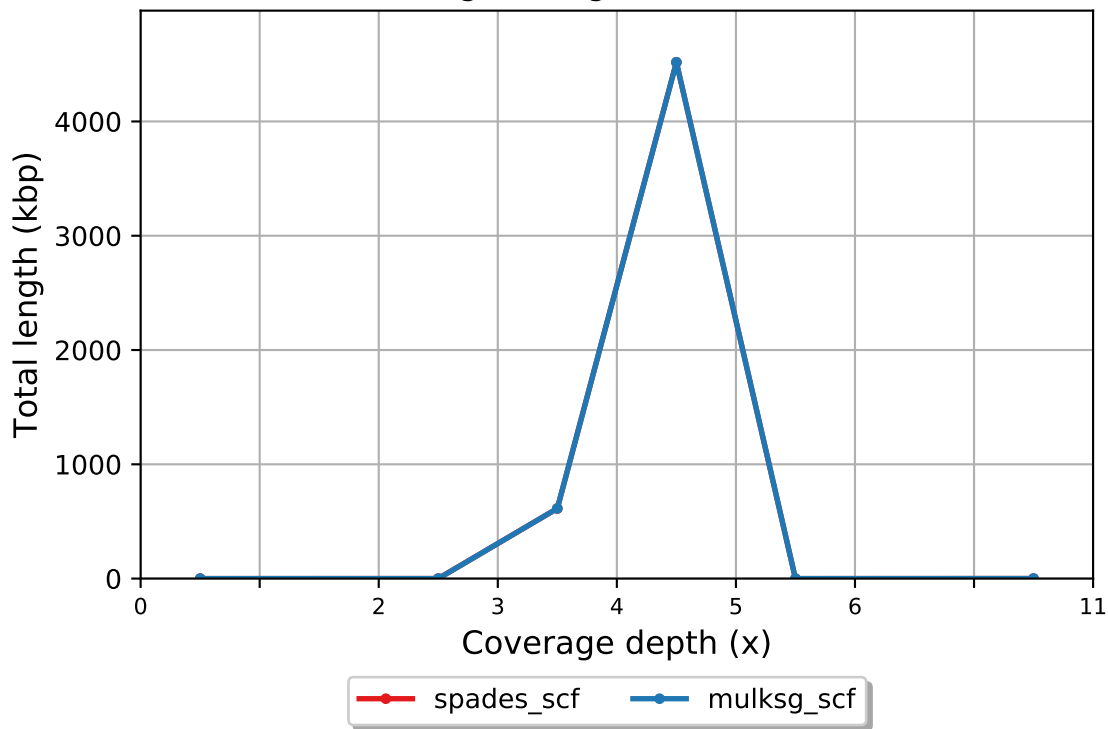
spades\_scf GC content



mulksg\_scf GC content

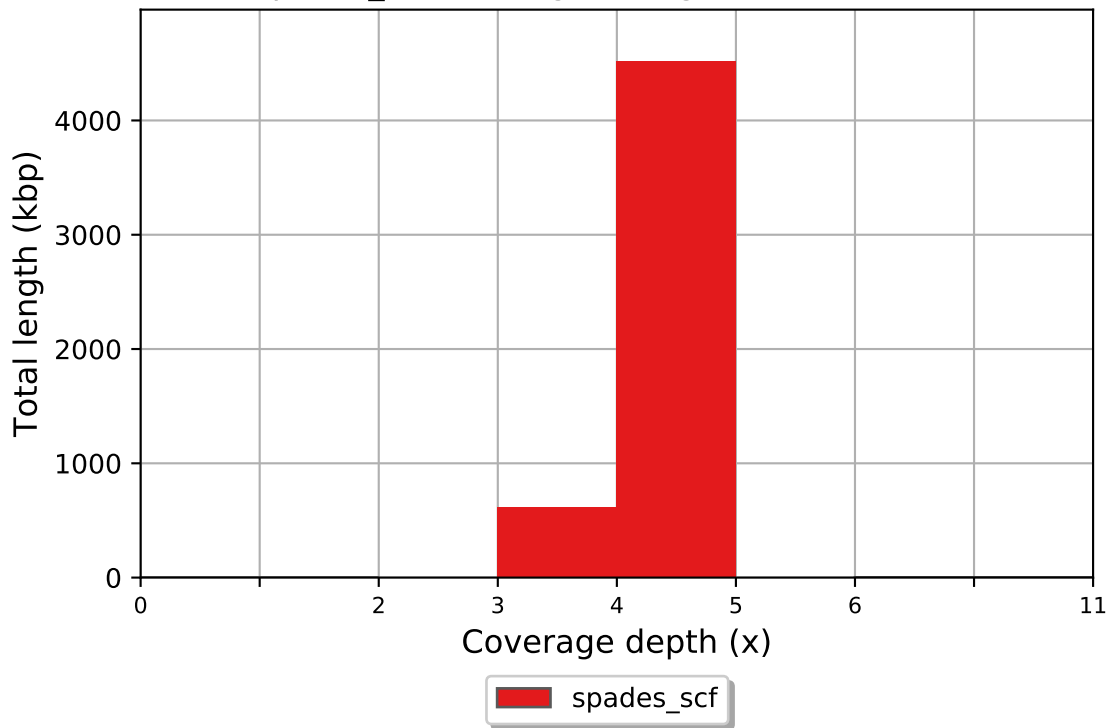


Coverage histogram (bin size: 1x)





spades\_scf coverage histogram (bin size: 1x)



mulksg\_scf coverage histogram (bin size: 1x)

