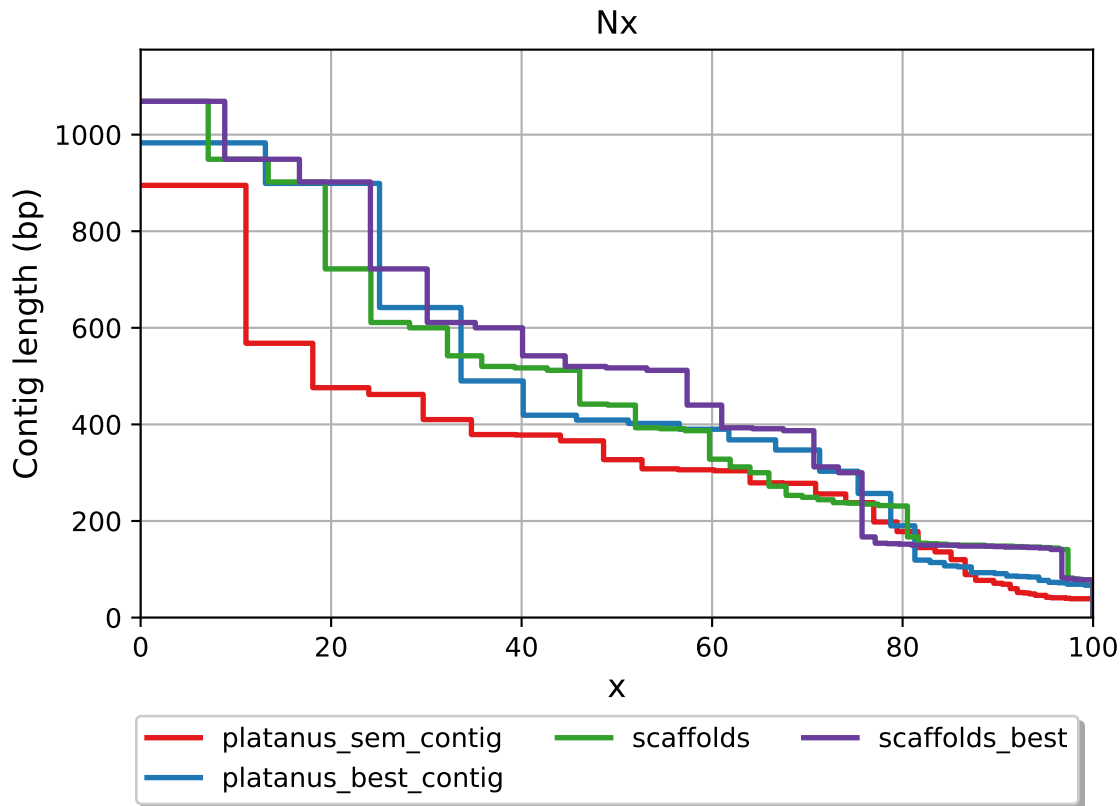
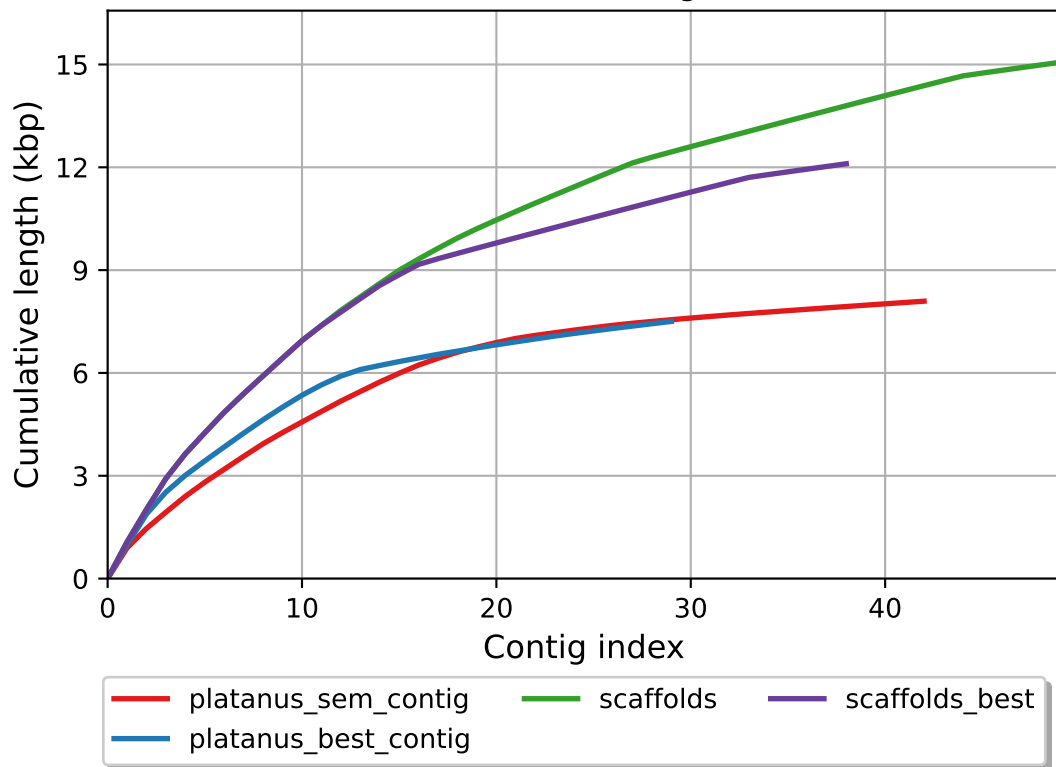


# Report

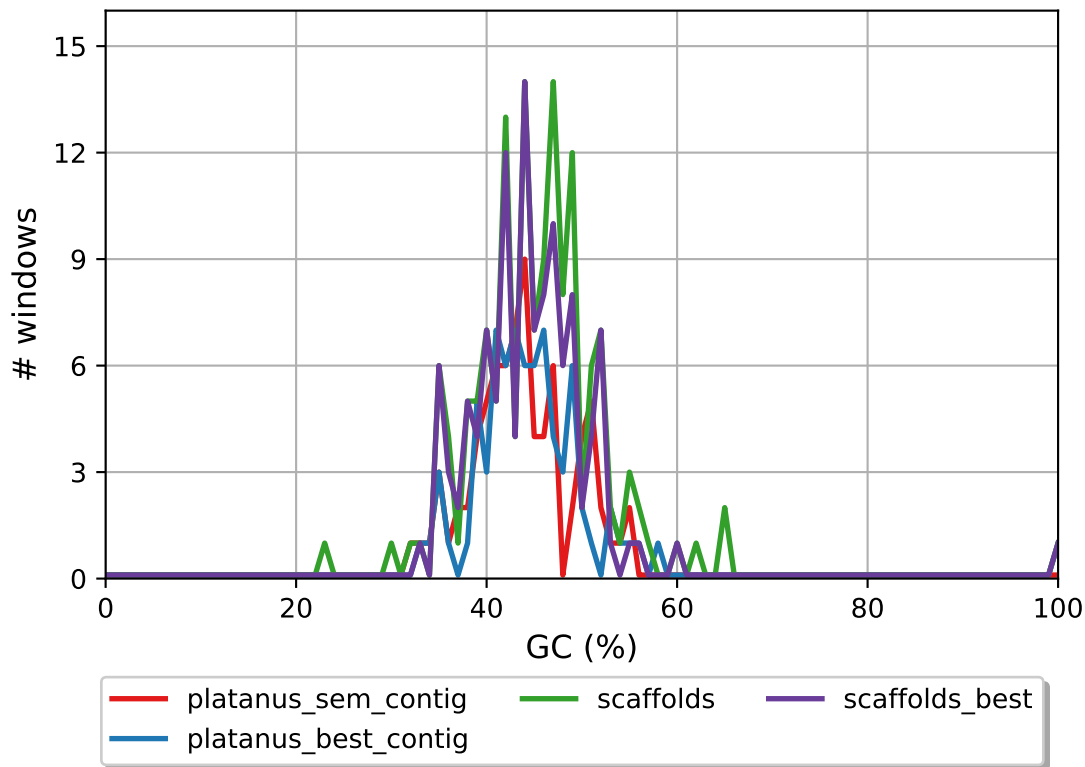
	platanus_sem_contig	platanus_best_contig	scaffolds	scaffolds_best
# contigs (>= 0 bp)	42	29	49	38
# contigs (>= 1000 bp)	0	0	1	1
# contigs (>= 5000 bp)	0	0	0	0
# contigs (>= 10000 bp)	0	0	0	0
# contigs (>= 25000 bp)	0	0	0	0
# contigs (>= 50000 bp)	0	0	0	0
Total length (>= 0 bp)	8094	7503	15064	12103
Total length (>= 1000 bp)	0	0	1069	1069
Total length (>= 5000 bp)	0	0	0	0
Total length (>= 10000 bp)	0	0	0	0
Total length (>= 25000 bp)	0	0	0	0
Total length (>= 50000 bp)	0	0	0	0
# contigs	42	29	49	38
Largest contig	895	983	1069	1069
Total length	8094	7503	15064	12103
GC (%)	44.22	44.76	45.47	44.64
N50	327	409	440	517
N90	71	91	148	147
auN	371.1	487.9	484.1	533.3
L50	9	6	12	9
L90	25	20	37	28
# N's per 100 kbp	0.00	0.00	0.00	0.00



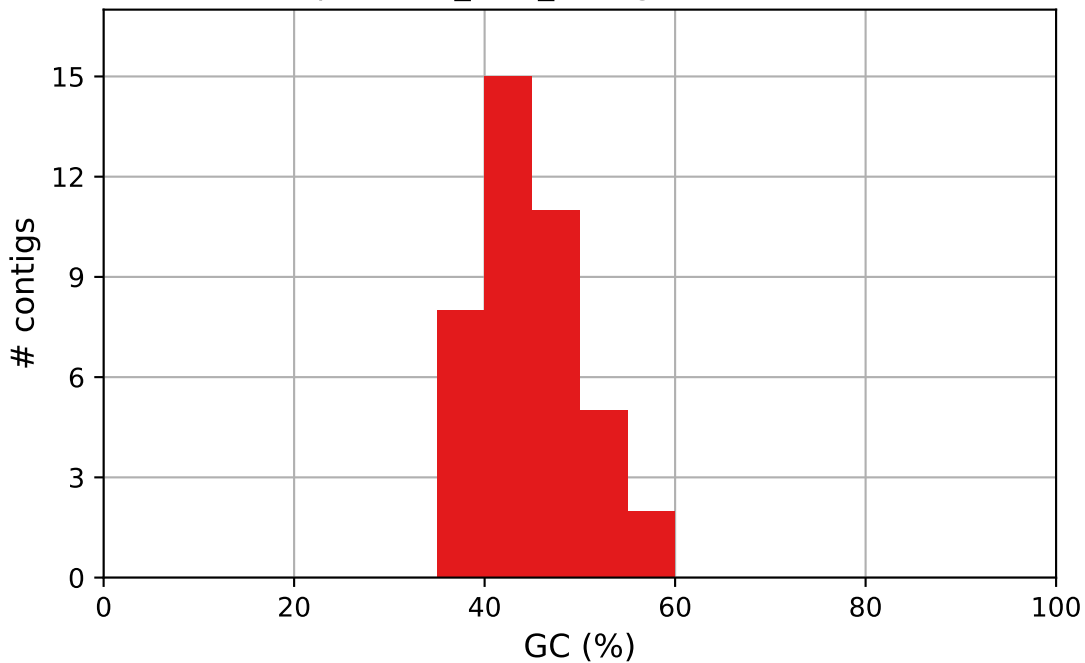
Cumulative length



GC content

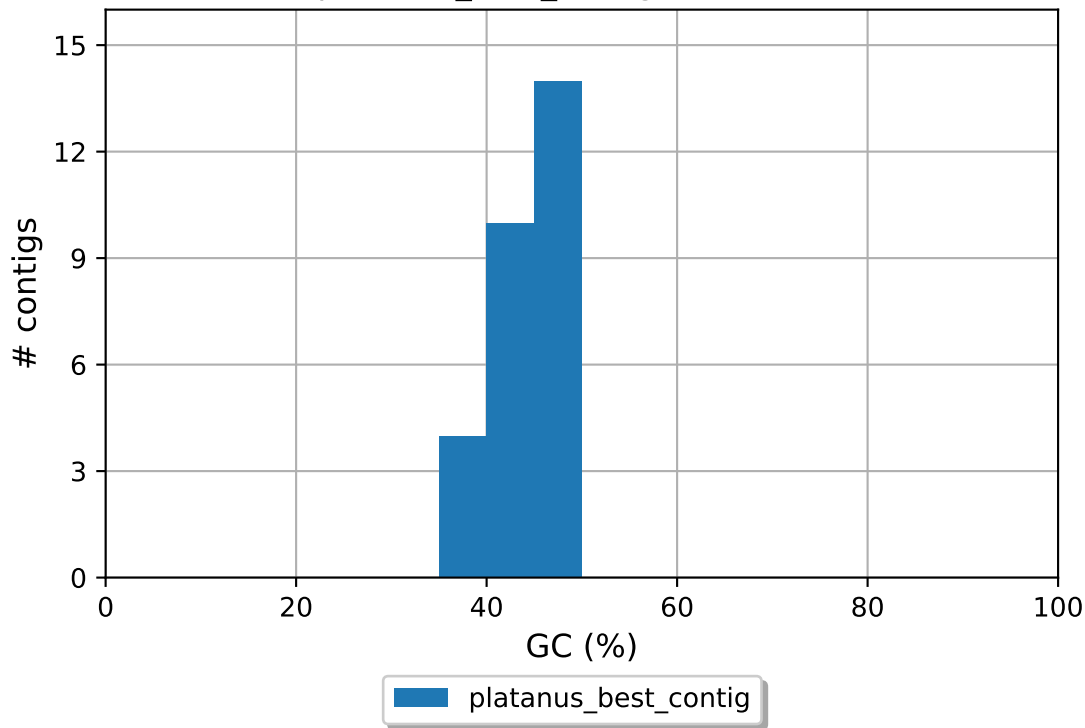


platanus\_sem\_contig GC content

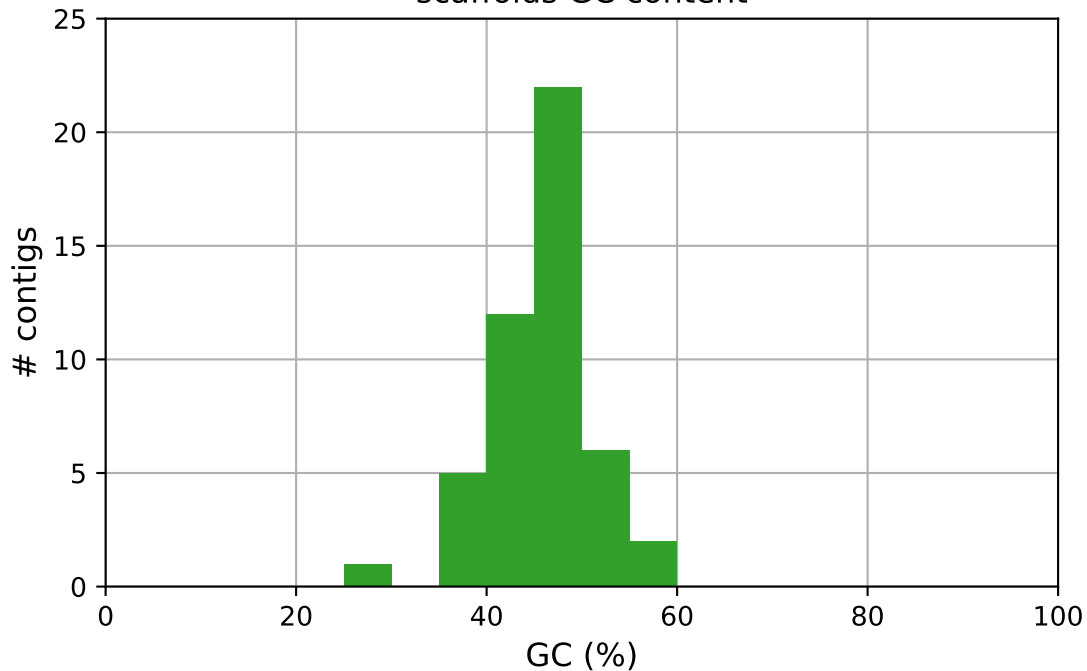


platanus\_sem\_contig

platanus\_best\_contig GC content

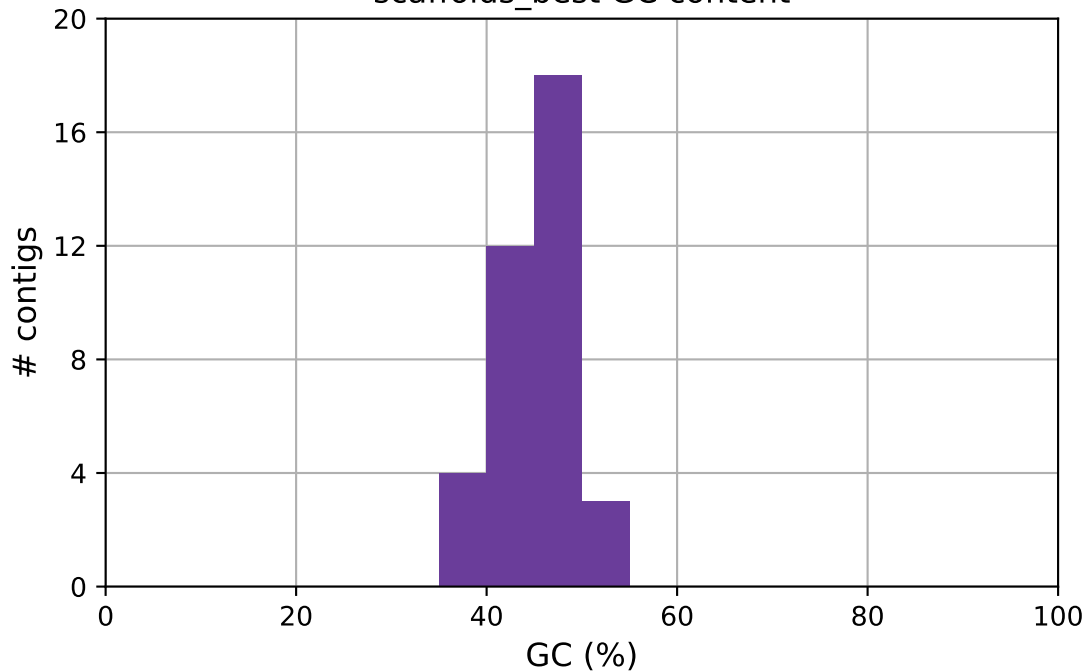


scaffolds GC content



scaffolds

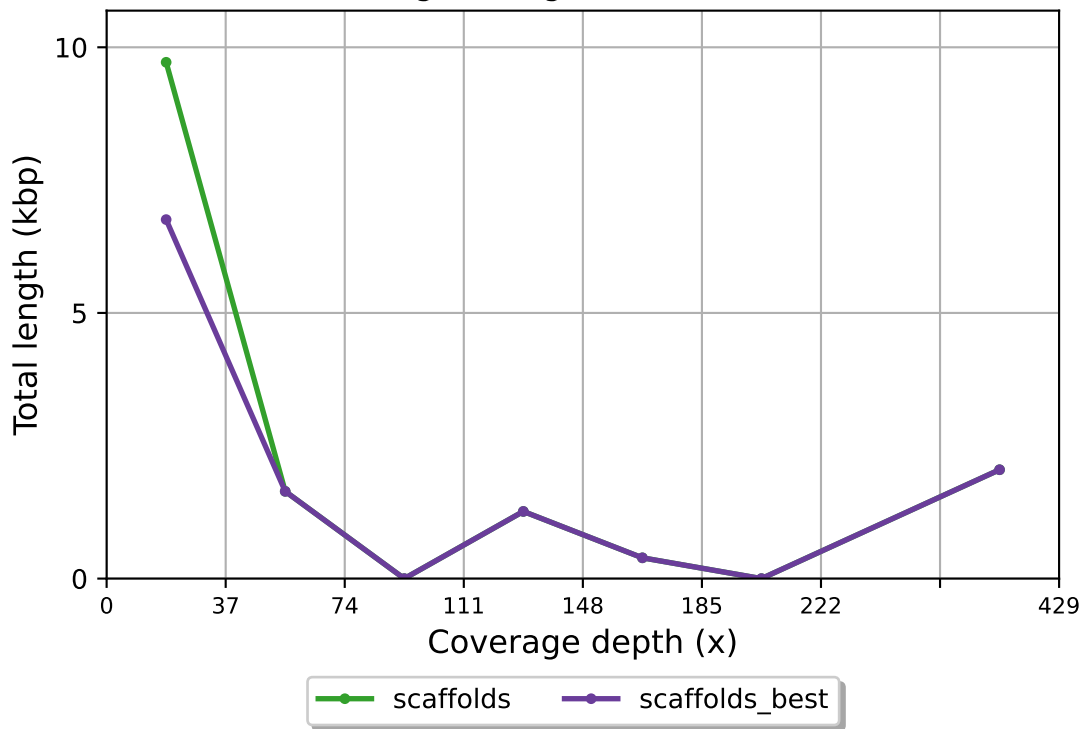
scaffolds\_best GC content



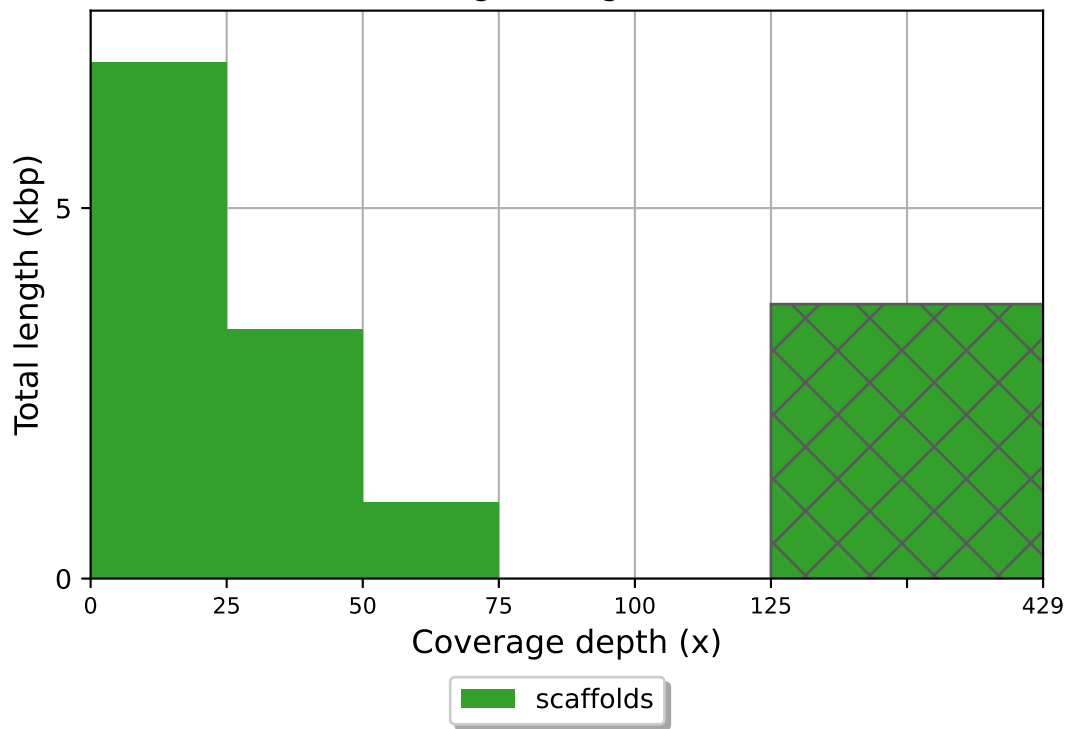
scaffolds\_best



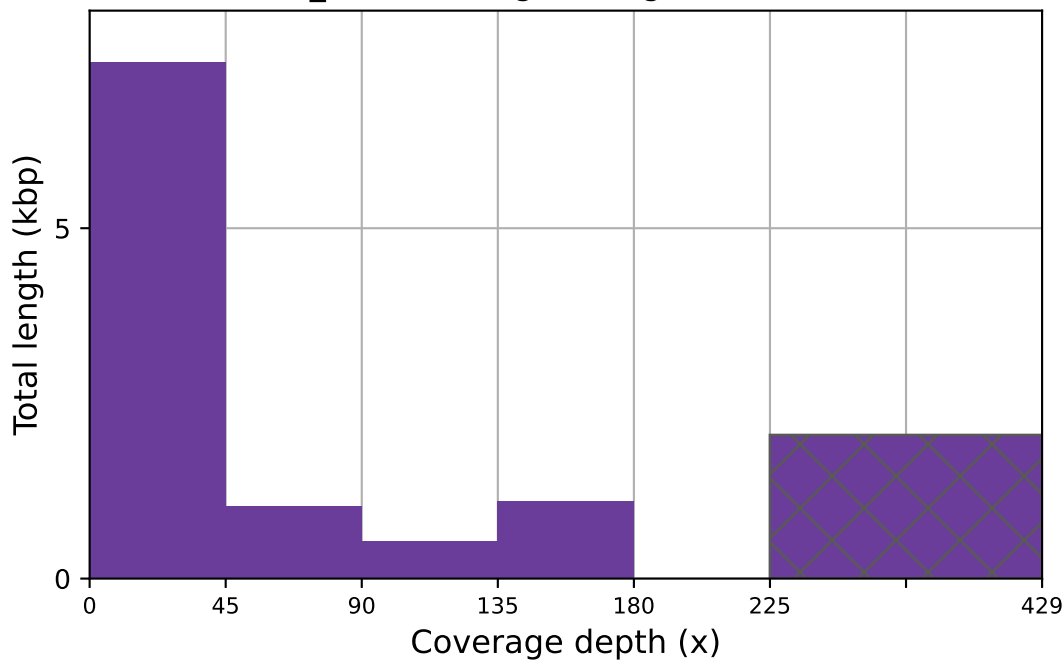
Coverage histogram (bin size: 37x)



scaffolds coverage histogram (bin size: 25x)



scaffolds\_best coverage histogram (bin size: 45x)



scaffolds\_best