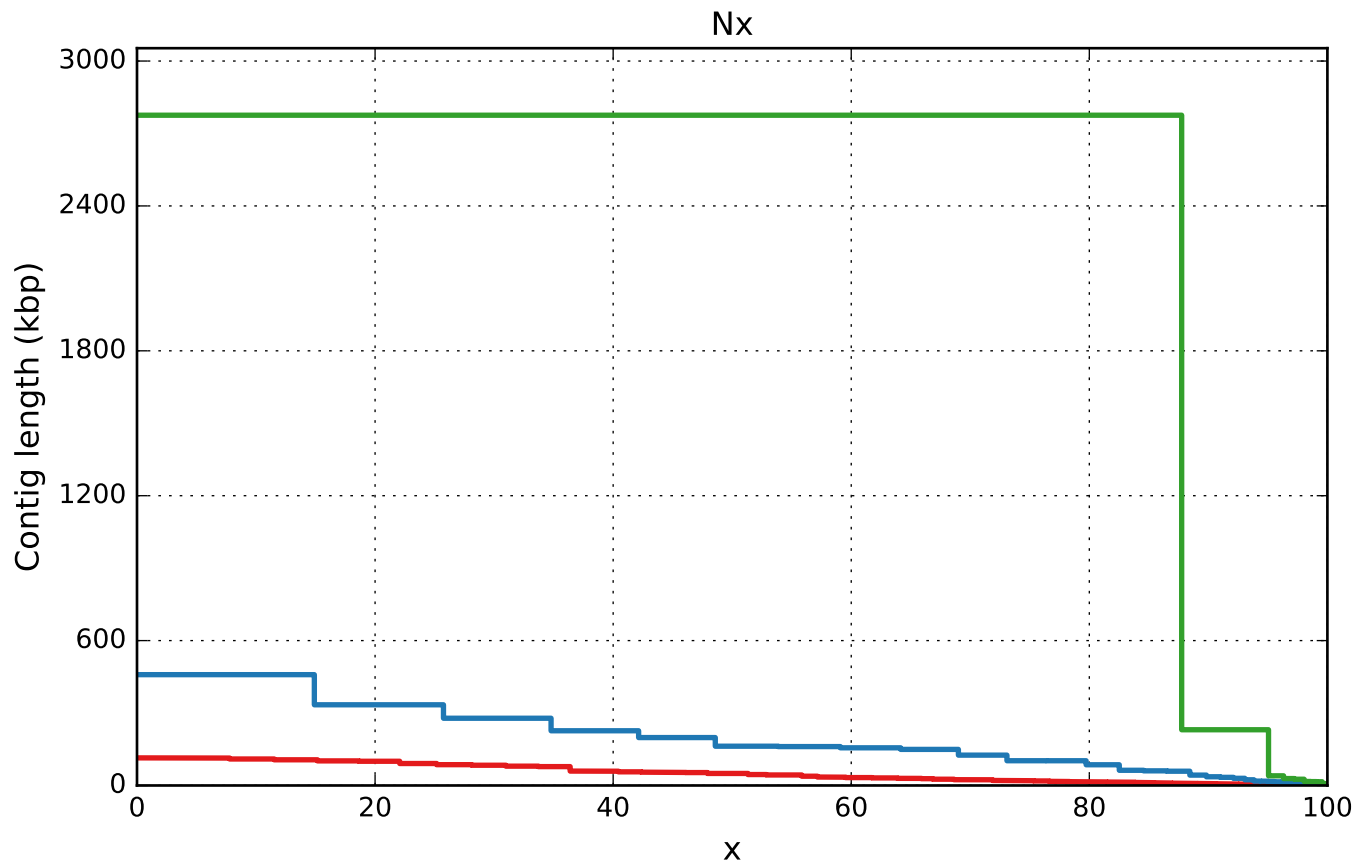


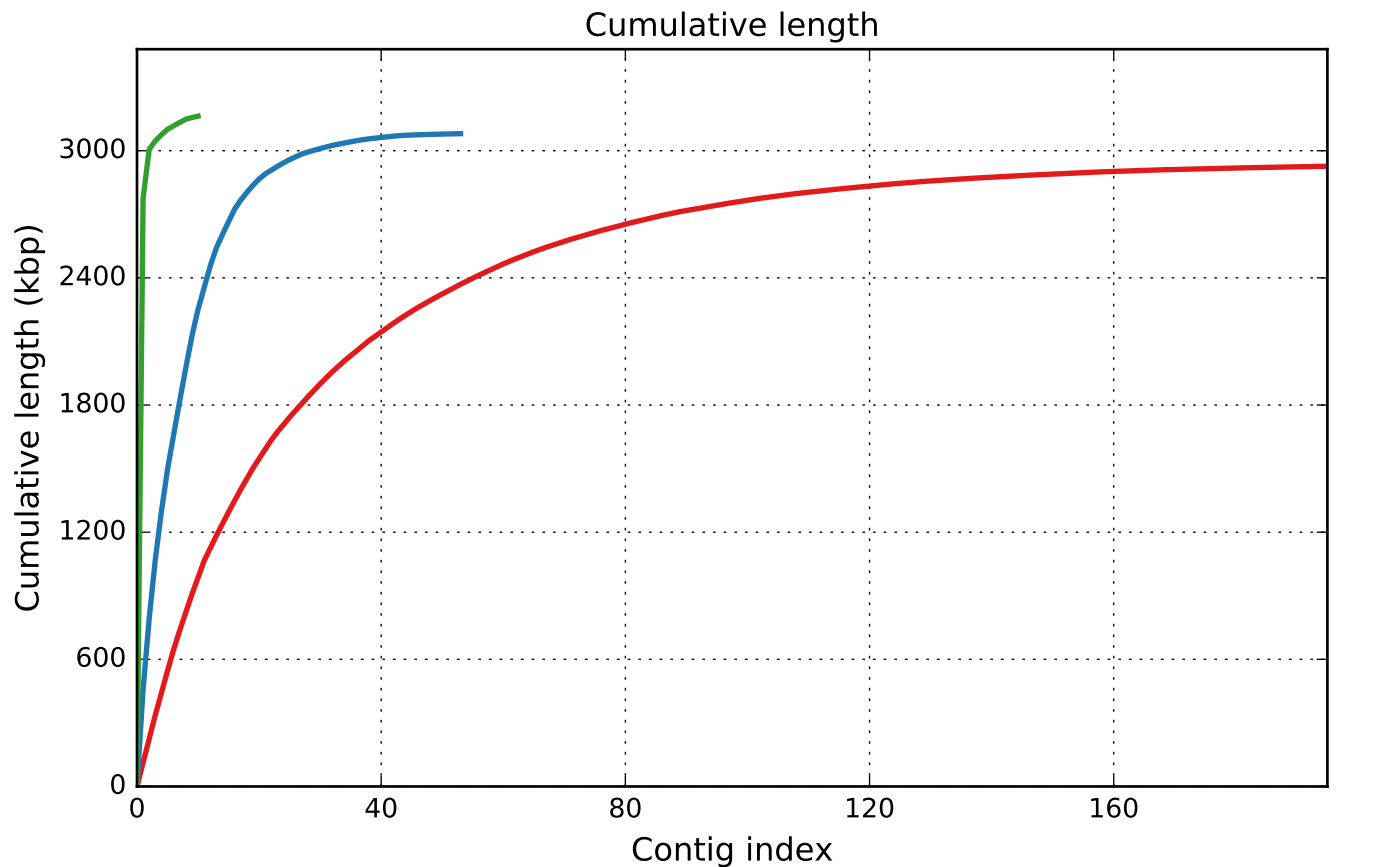
## Report

	illumina.contigs	illumina_Nanopore.contigs	PacBio.contigs
# contigs (>= 0 bp)	488	175	10
# contigs (>= 1000 bp)	163	46	10
# contigs (>= 5000 bp)	92	36	10
# contigs (>= 10000 bp)	67	27	8
# contigs (>= 25000 bp)	34	20	5
# contigs (>= 50000 bp)	17	16	2
Total length (>= 0 bp)	2975893	3101302	3162825
Total length (>= 1000 bp)	2905560	3075336	3162825
Total length (>= 5000 bp)	2728114	3047294	3162825
Total length (>= 10000 bp)	2544168	2984487	3149049
Total length (>= 25000 bp)	2008843	2866708	3100952
Total length (>= 50000 bp)	1402640	2724412	3006416
# contigs	195	53	10
Largest contig	114283	458911	2775639
Total length	2926792	3080395	3162825
GC (%)	37.65	37.64	37.77
N50	49767	162629	2775639
N75	19664	102479	2775639
L50	19	6	1
L75	43	11	1
# N's per 100 kbp	0.00	0.00	0.00

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).

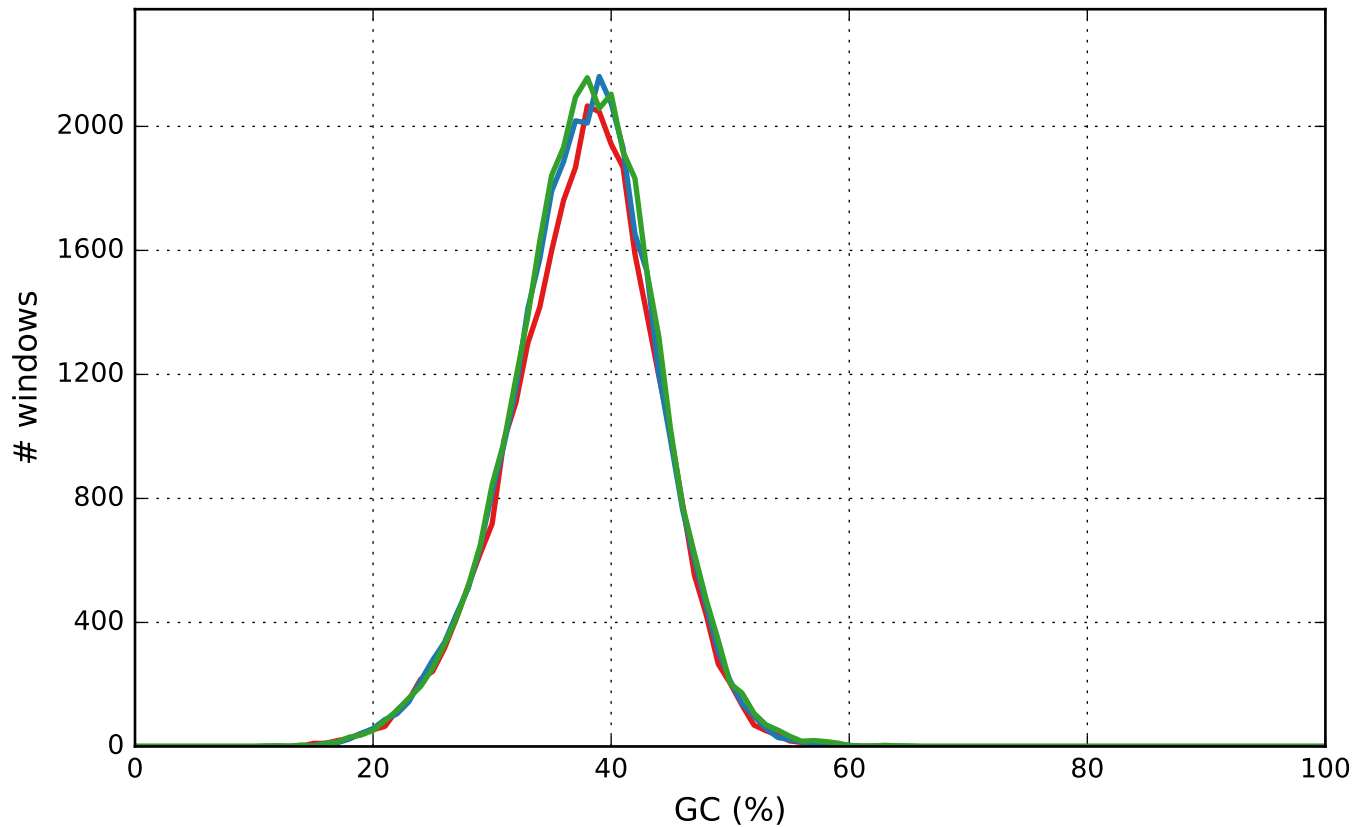


— illumina.contigs — illumina\_Nanopore.contigs — PacBio.contigs



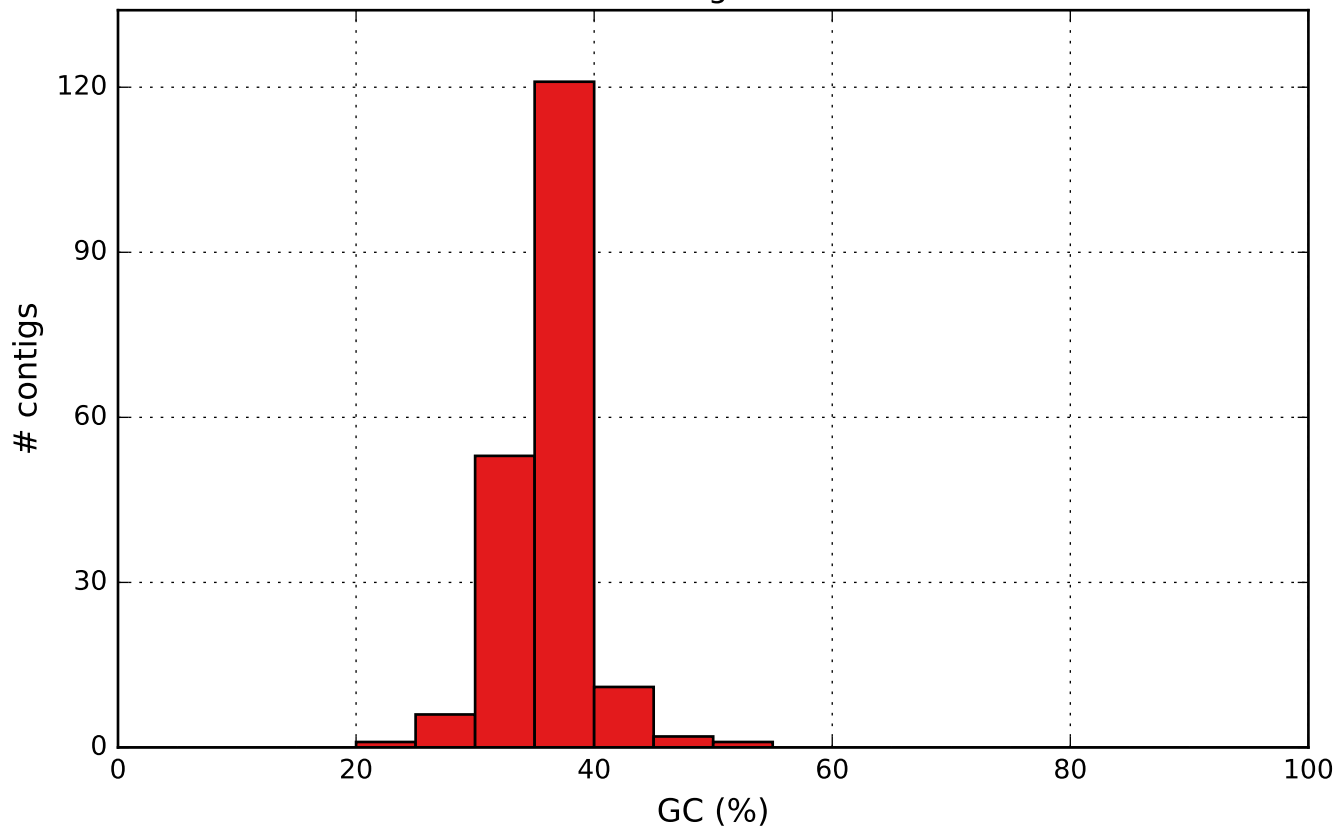
— illumina.contigs    — illumina\_Nanopore.contigs    — PacBio.contigs

GC content

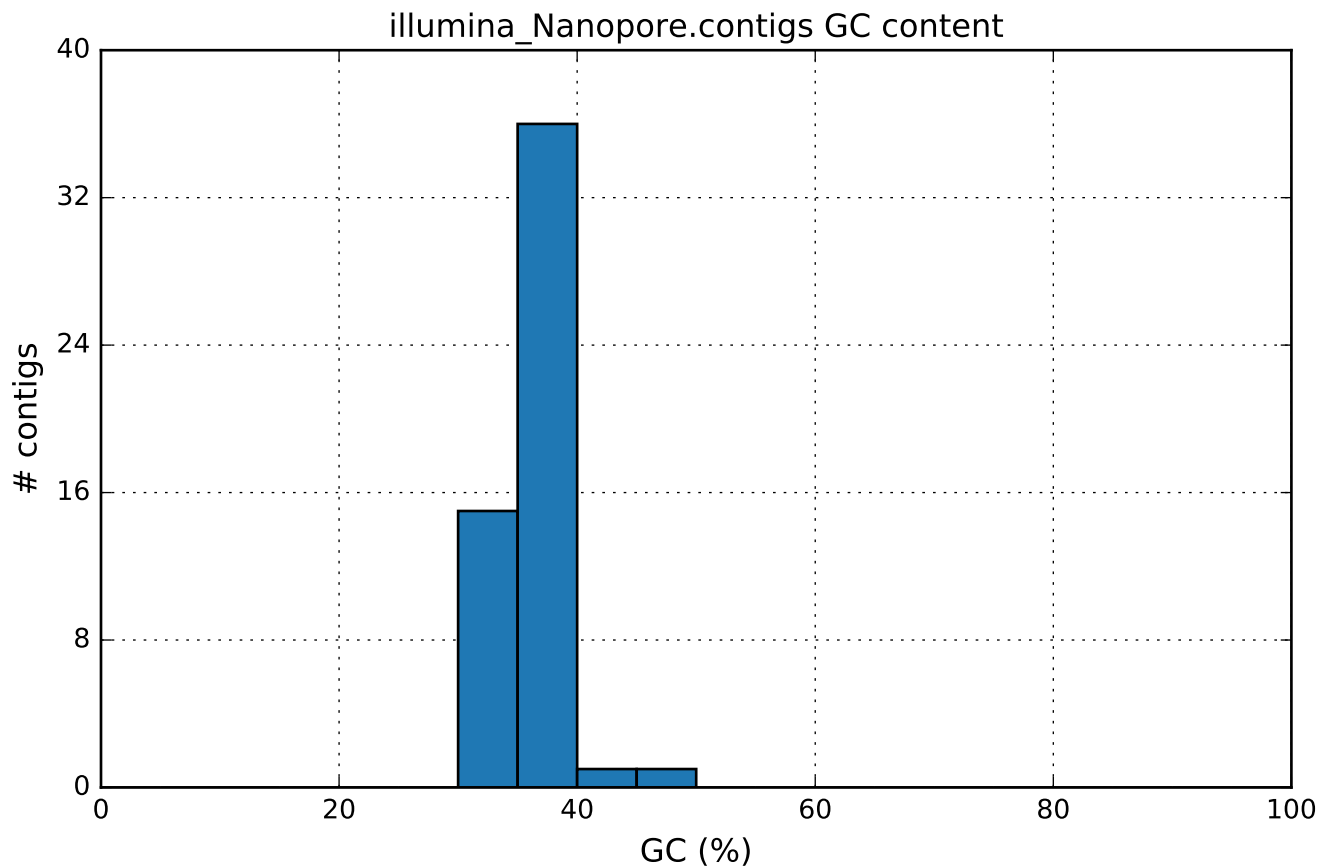


— illumina.contigs — illumina\_Nanopore.contigs — PacBio.contigs

illumina.contigs GC content

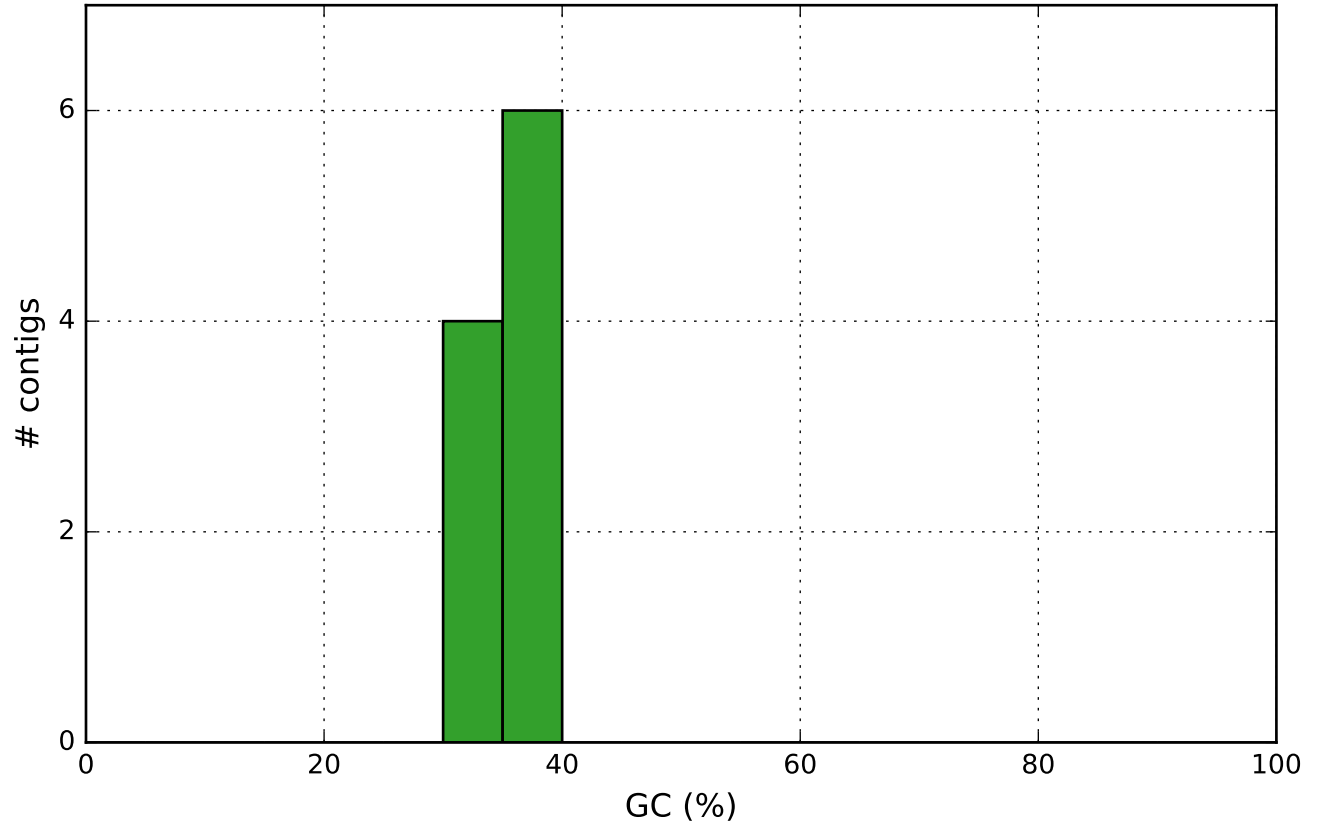


illumina.contigs



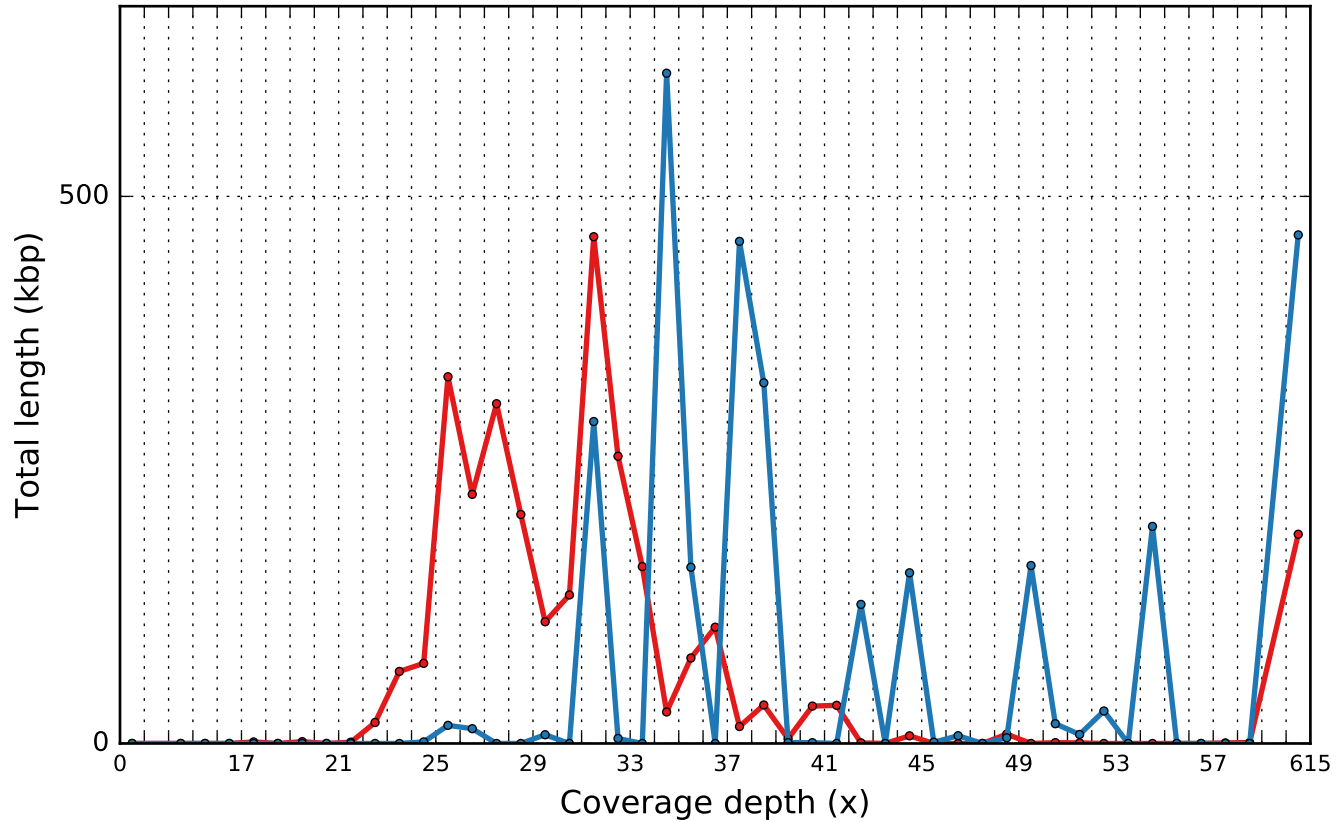
illumina\_Nanopore.contigs

PacBio.contigs GC content



 PacBio.contigs

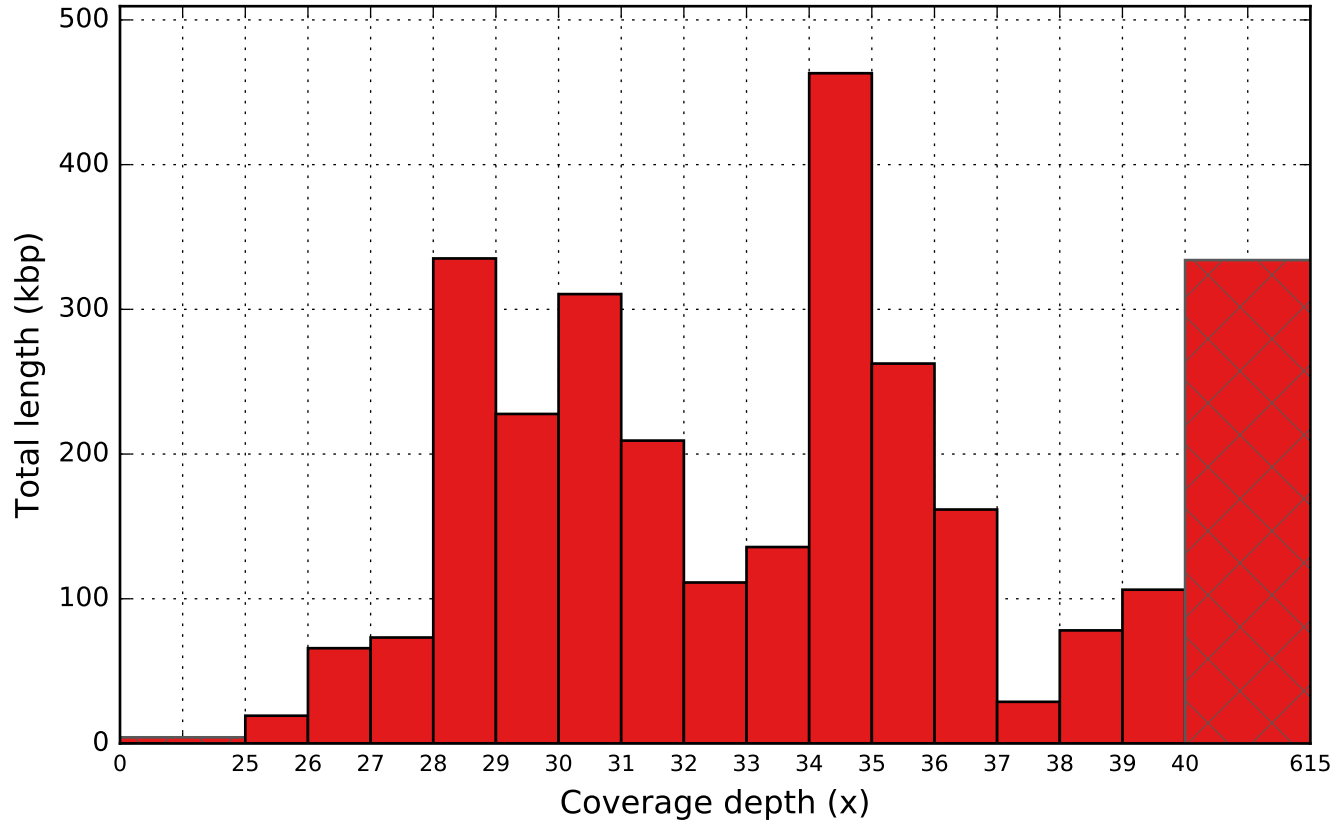
Coverage histogram (bin size: 1x)



illumina.contigs      illumina\_Nanopore.contigs

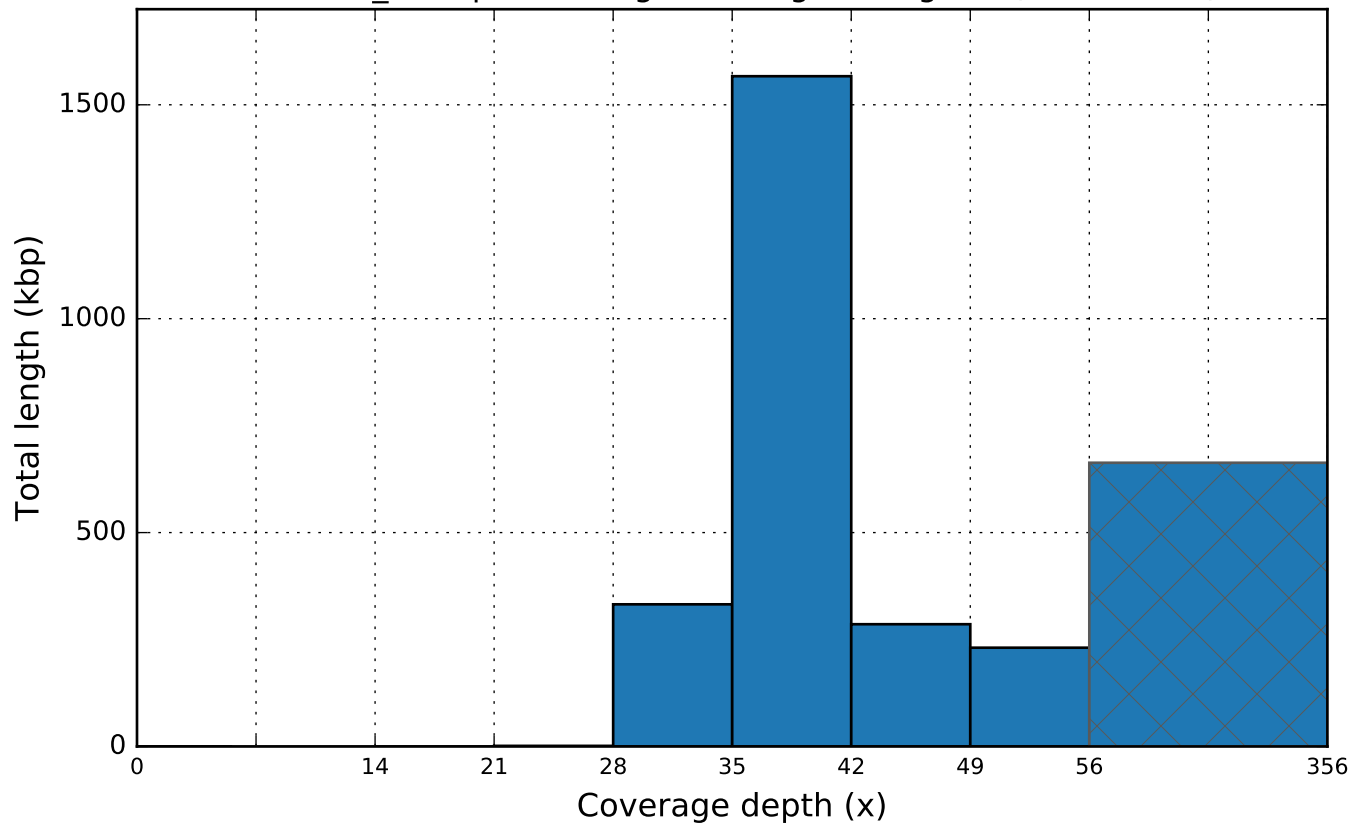


illumina.contigs coverage histogram (bin size: 1x)



illumina.contigs

illumina\_Nanopore.contigs coverage histogram (bin size: 7x)



illumina\_Nanopore.contigs