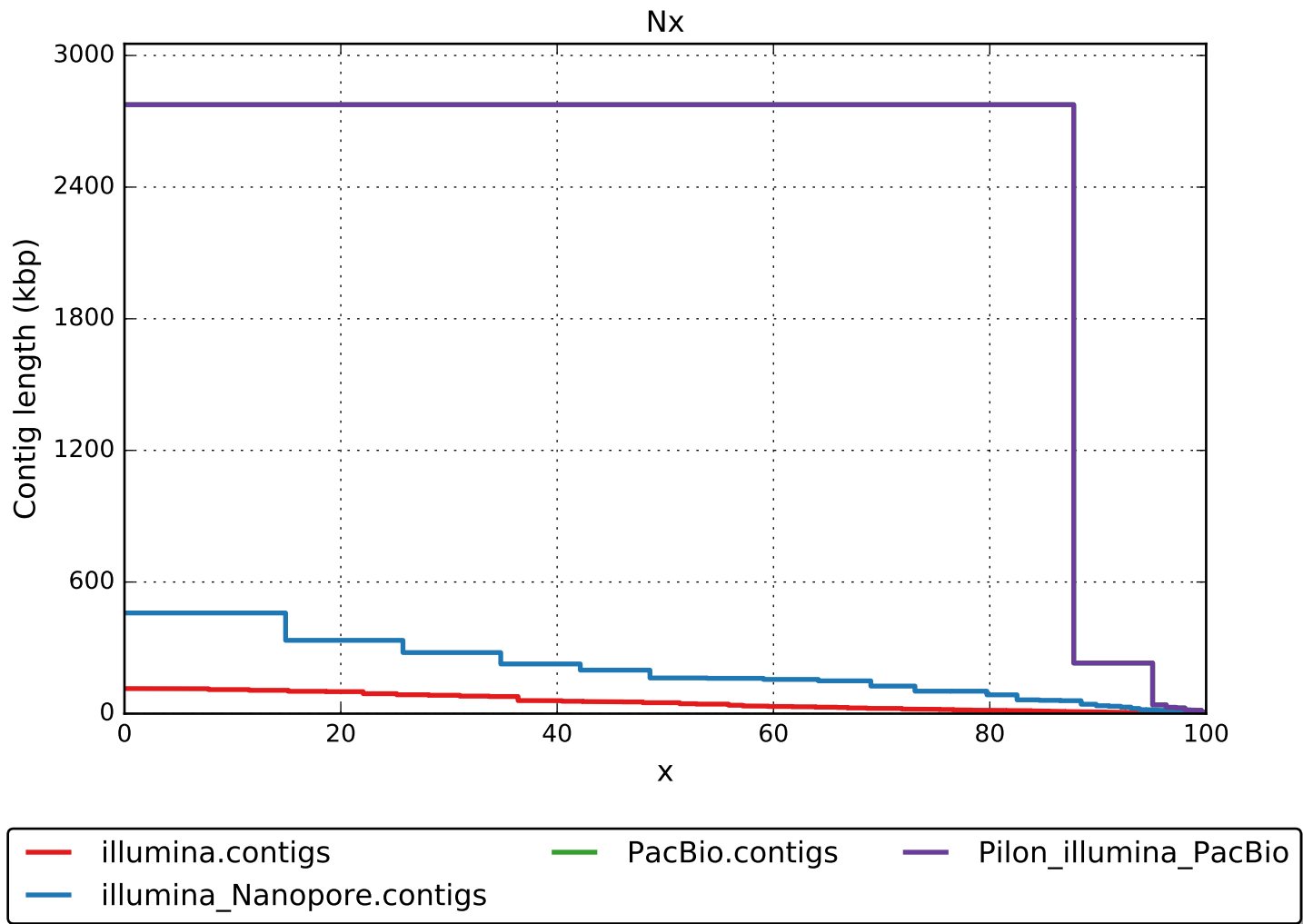
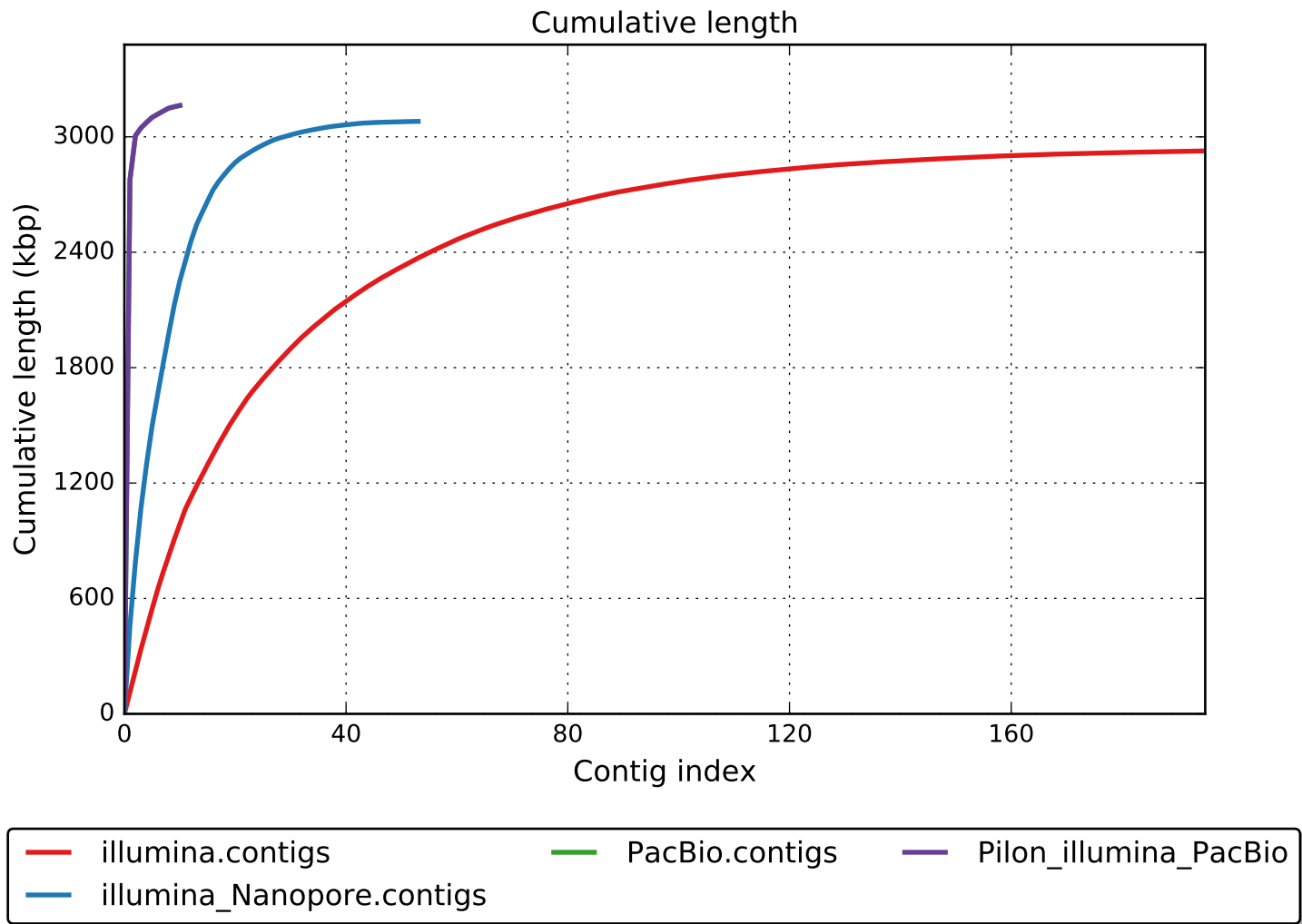


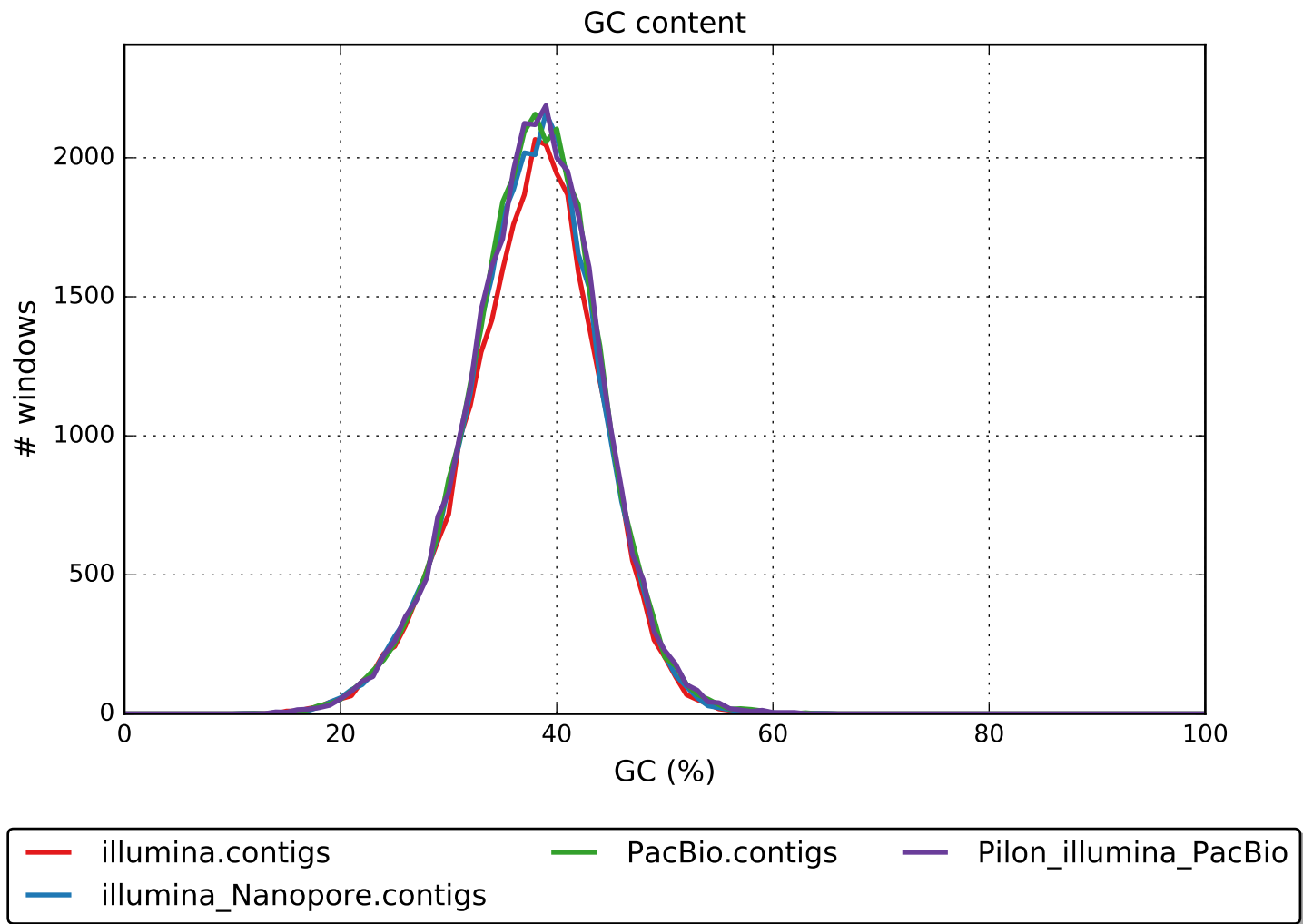
Report

	illumina.contigs	illumina_Nanopore.contigs	PacBio.contigs	Pilon_illumina_PacBio
# contigs (>= 0 bp)	488	175	10	10
# contigs (>= 1000 bp)	163	46	10	10
# contigs (>= 5000 bp)	92	36	10	10
# contigs (>= 10000 bp)	67	27	8	8
# contigs (>= 25000 bp)	34	20	5	5
# contigs (>= 50000 bp)	17	16	2	2
Total length (>= 0 bp)	2975893	3101302	3162825	3162984
Total length (>= 1000 bp)	2905560	3075336	3162825	3162984
Total length (>= 5000 bp)	2728114	3047294	3162825	3162984
Total length (>= 10000 bp)	2544168	2984487	3149049	3149194
Total length (>= 25000 bp)	2008843	2866708	3100952	3101079
Total length (>= 50000 bp)	1402640	2724412	3006416	3006523
# contigs	195	53	10	10
Largest contig	114283	458911	2775639	2775725
Total length	2926792	3080395	3162825	3162984
GC (%)	37.65	37.64	37.77	37.77
N50	49767	162629	2775639	2775725
N75	19664	102479	2775639	2775725
L50	19	6	1	1
L75	43	11	1	1
# N's per 100 kbp	0.00	0.00	0.00	0.00

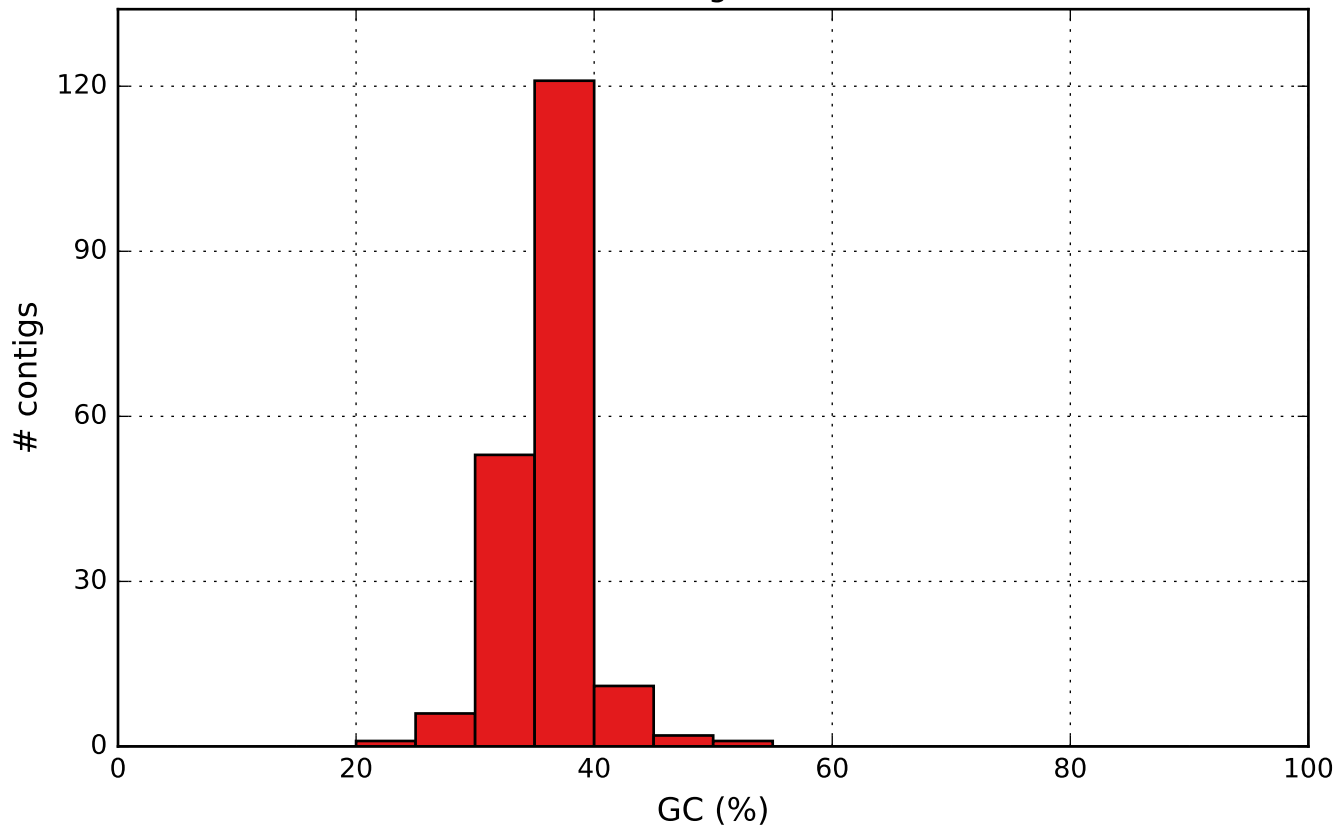
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



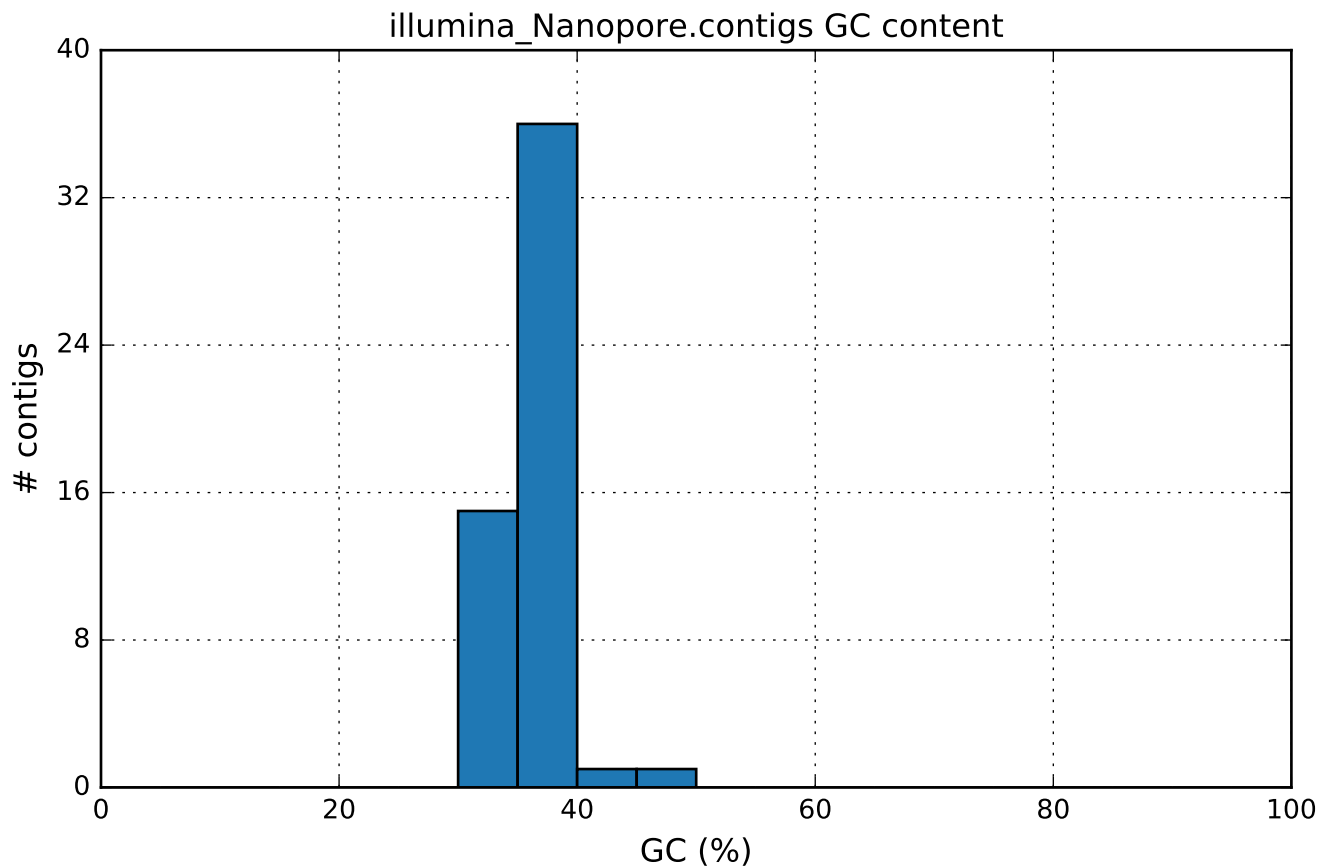




illumina.contigs GC content

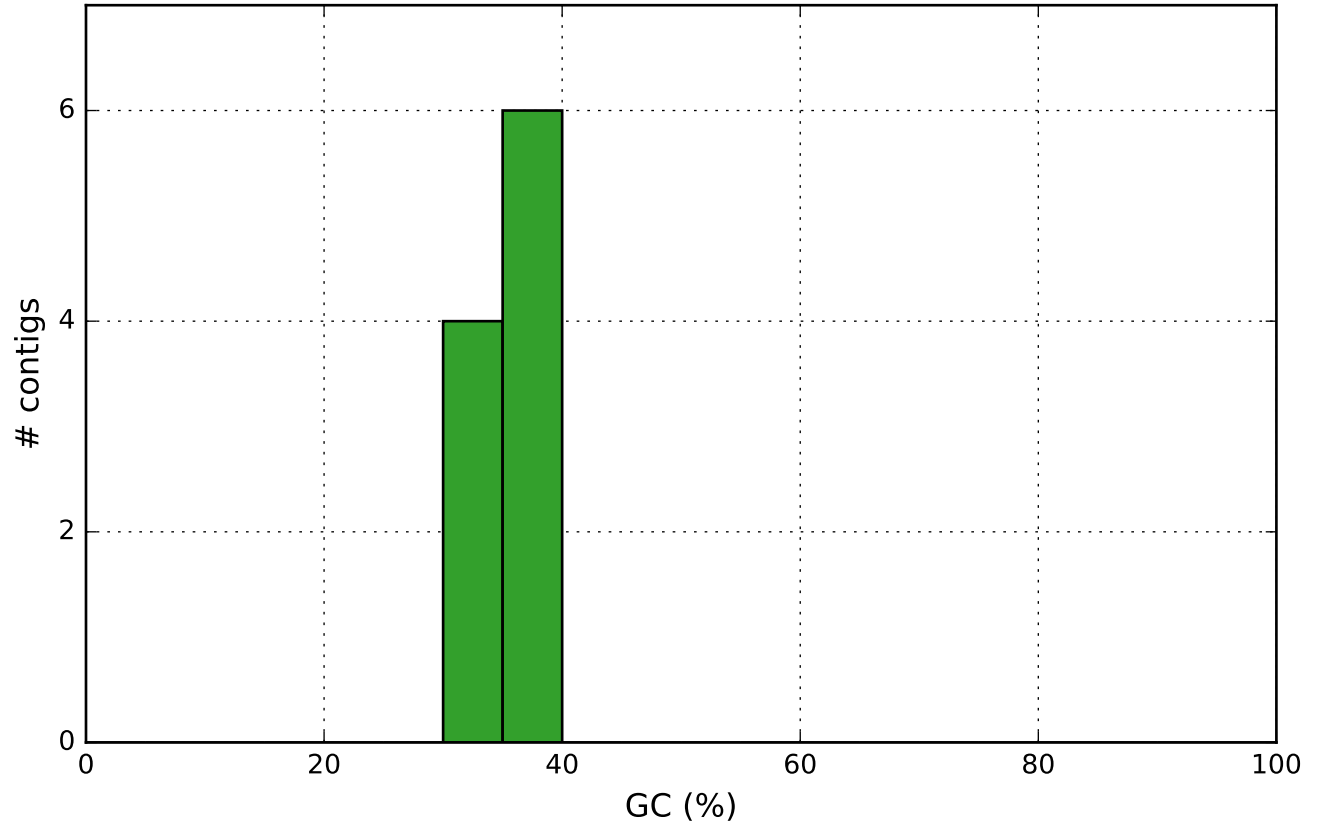


illumina.contigs



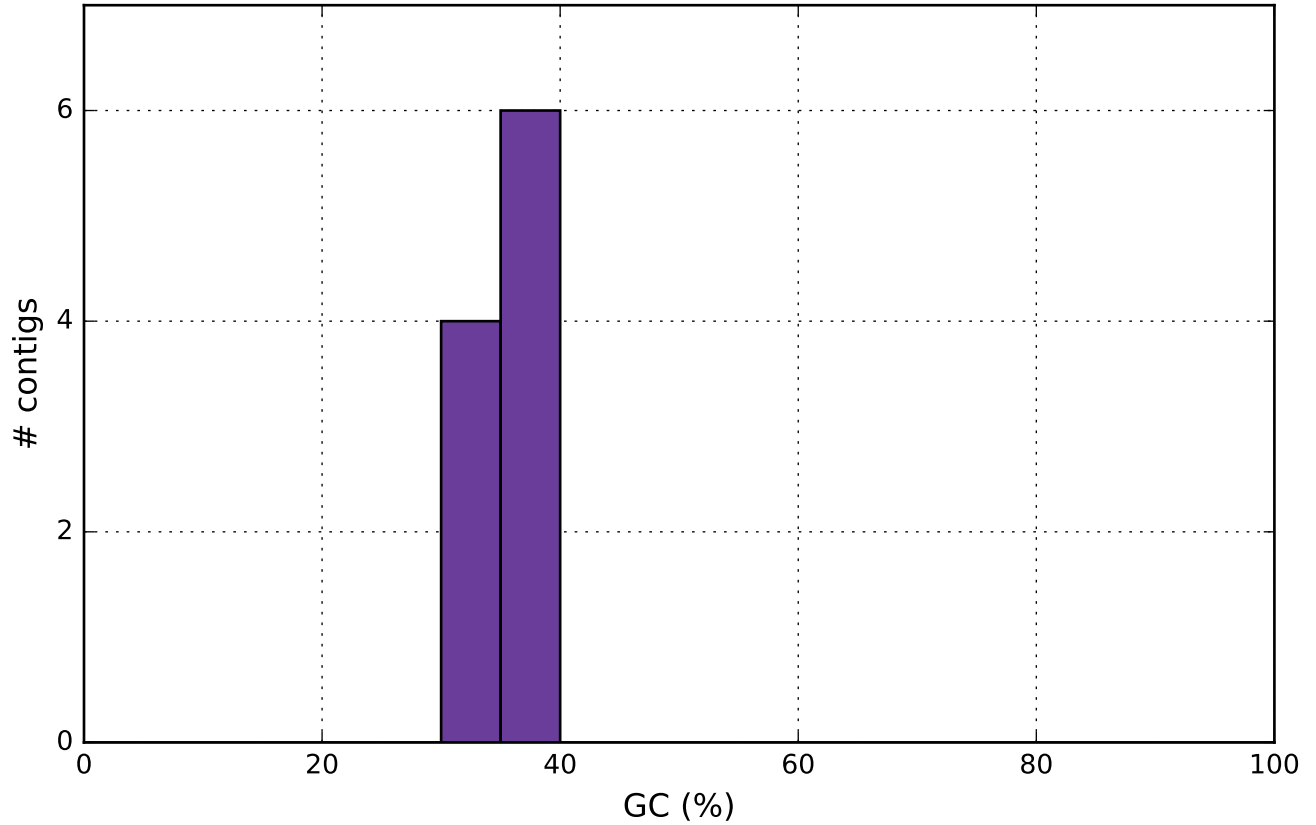
illumina_Nanopore.contigs

PacBio.contigs GC content



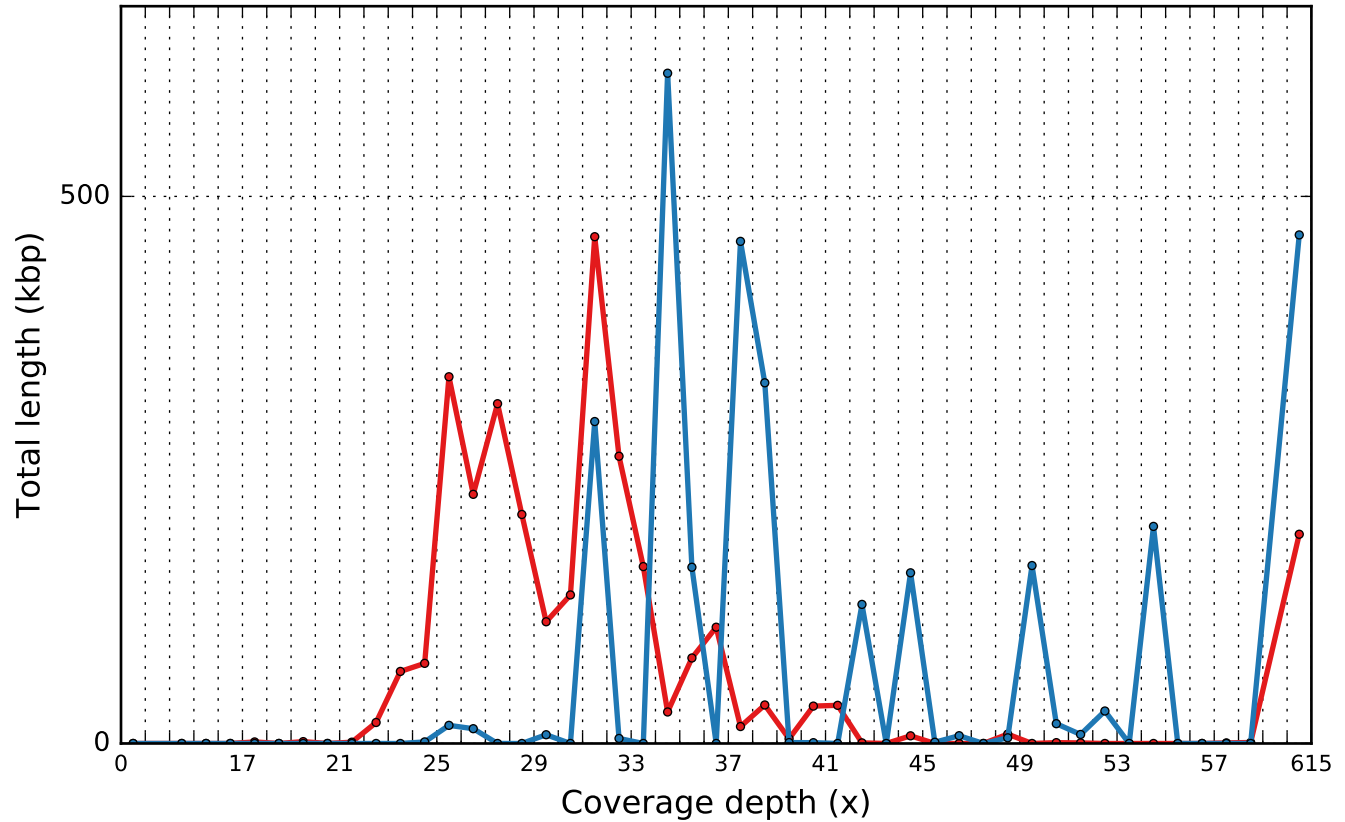
PacBio.contigs

Pilon_illumina_PacBio GC content



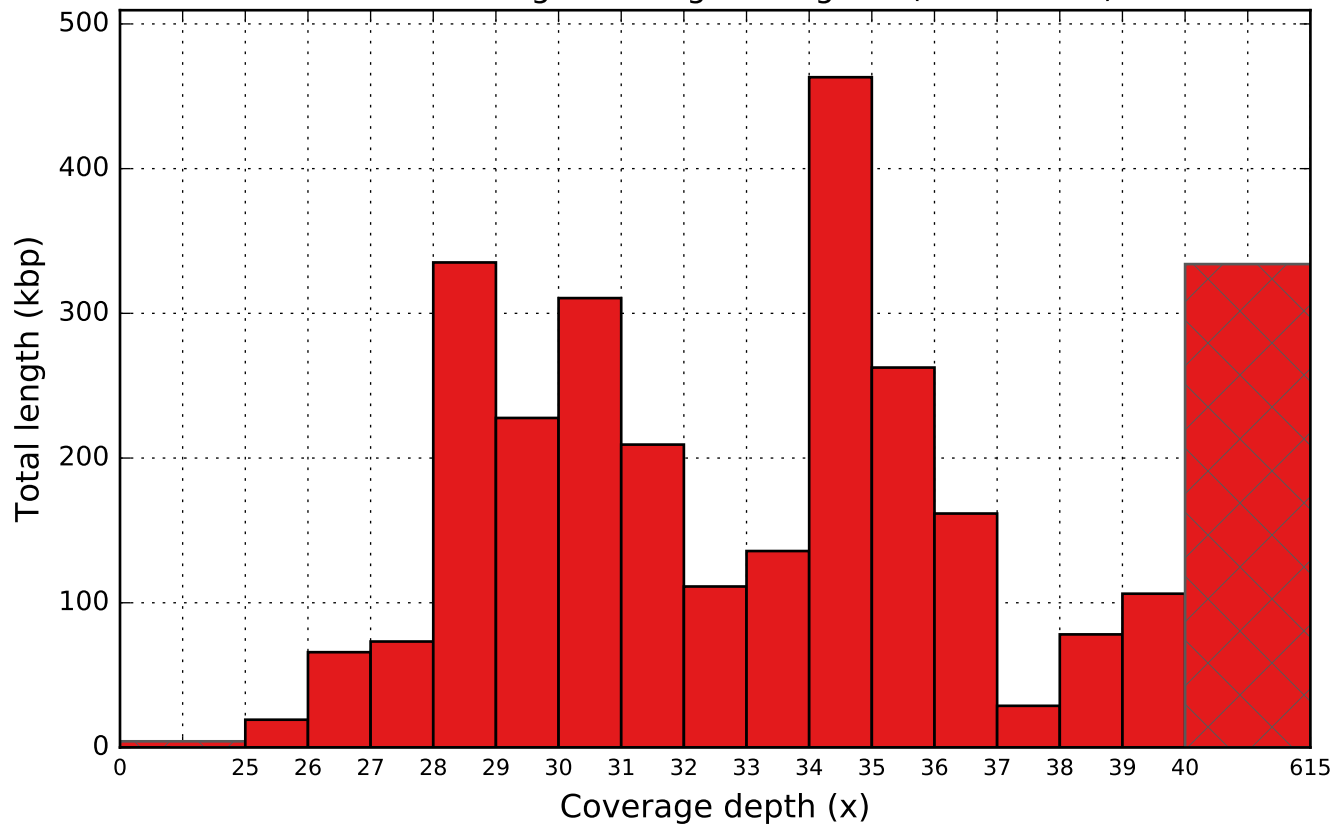
Pilon_illumina_PacBio

Coverage histogram (bin size: 1x)



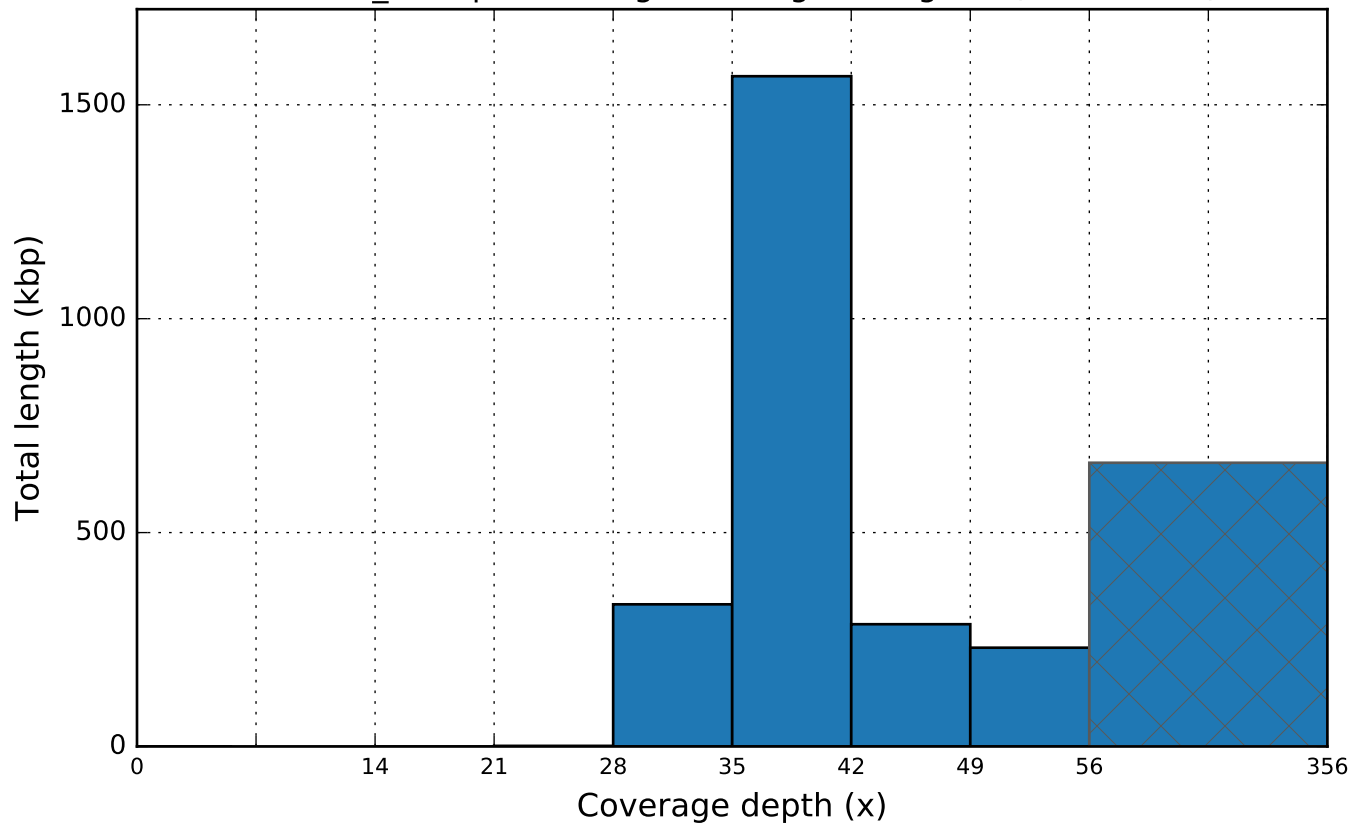
illumina.contigs illumina_Nanopore.contigs

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



illumina.contigs

illumina_Nanopore.contigs coverage histogram (bin size: 7x)



illumina_Nanopore.contigs