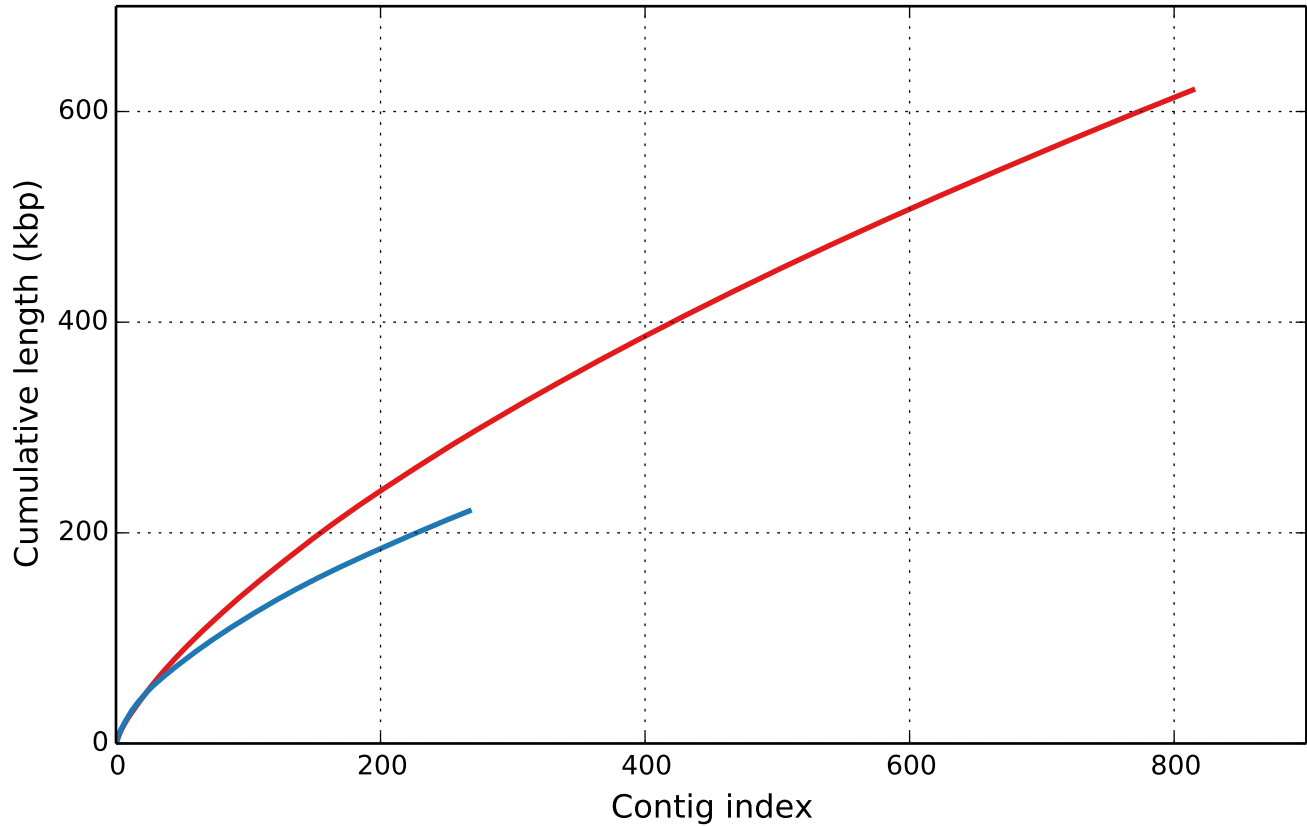


Report

	20150707_DNASeq_PE_P_kings-unitigs	20150721_DNASeq_PE_P_kings-unitigs
# contigs (≥ 0 bp)	304688	61713
# contigs (≥ 1000 bp)	119	45
Total length (≥ 0 bp)	30858010	6215246
Total length (≥ 1000 bp)	165188	72680
# contigs	814	267
Largest contig	3652	5024
Total length	620428	220758
GC (%)	43.71	44.38
N50	737	800
N75	589	600
L50	290	88
L75	527	167
# N's per 100 kbp	0.00	0.00
# predicted genes (unique)	488	145
# predicted genes (≥ 0 bp)	488	145
# predicted genes (≥ 300 bp)	250	69
# predicted genes (≥ 1500 bp)	7	5
# predicted genes (≥ 3000 bp)	2	1

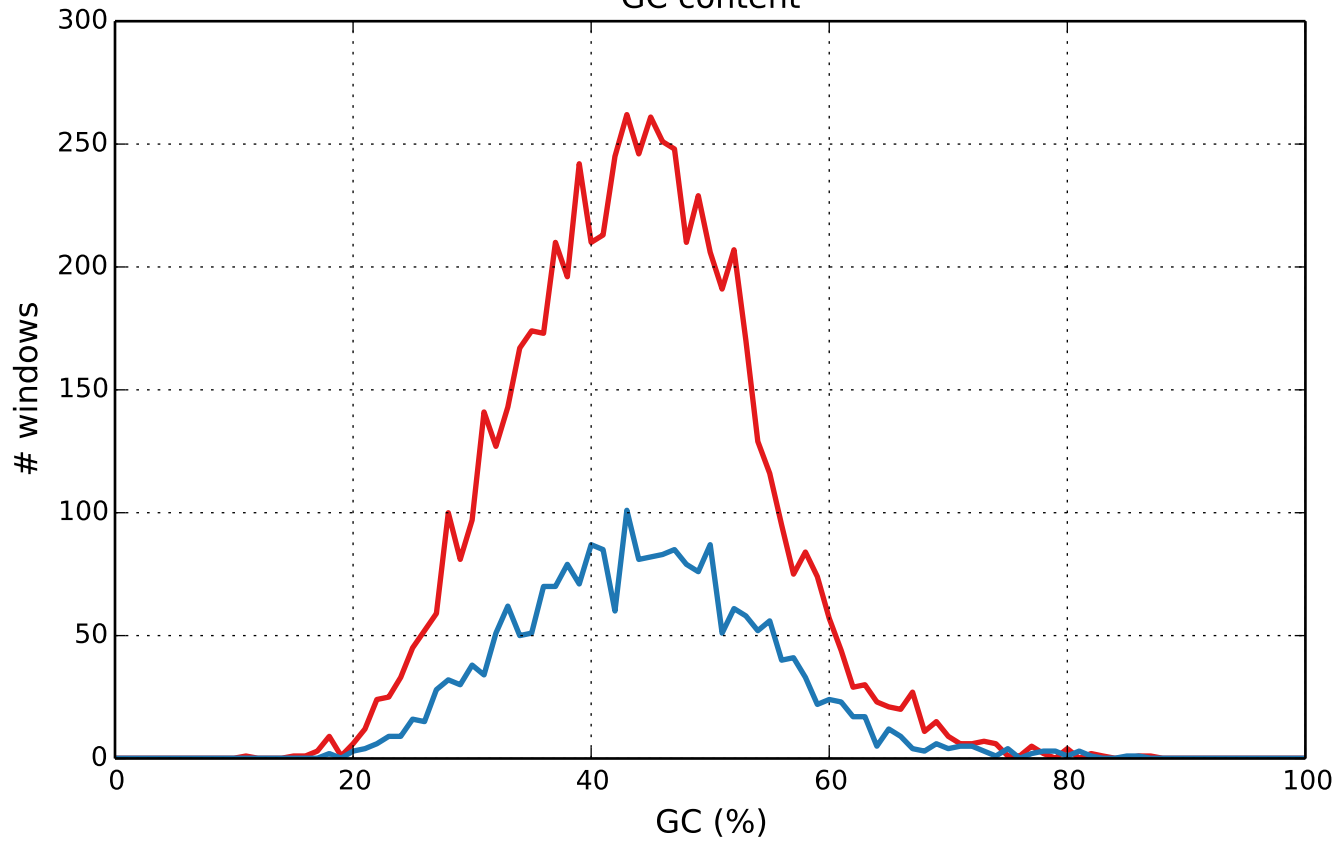
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Cumulative length

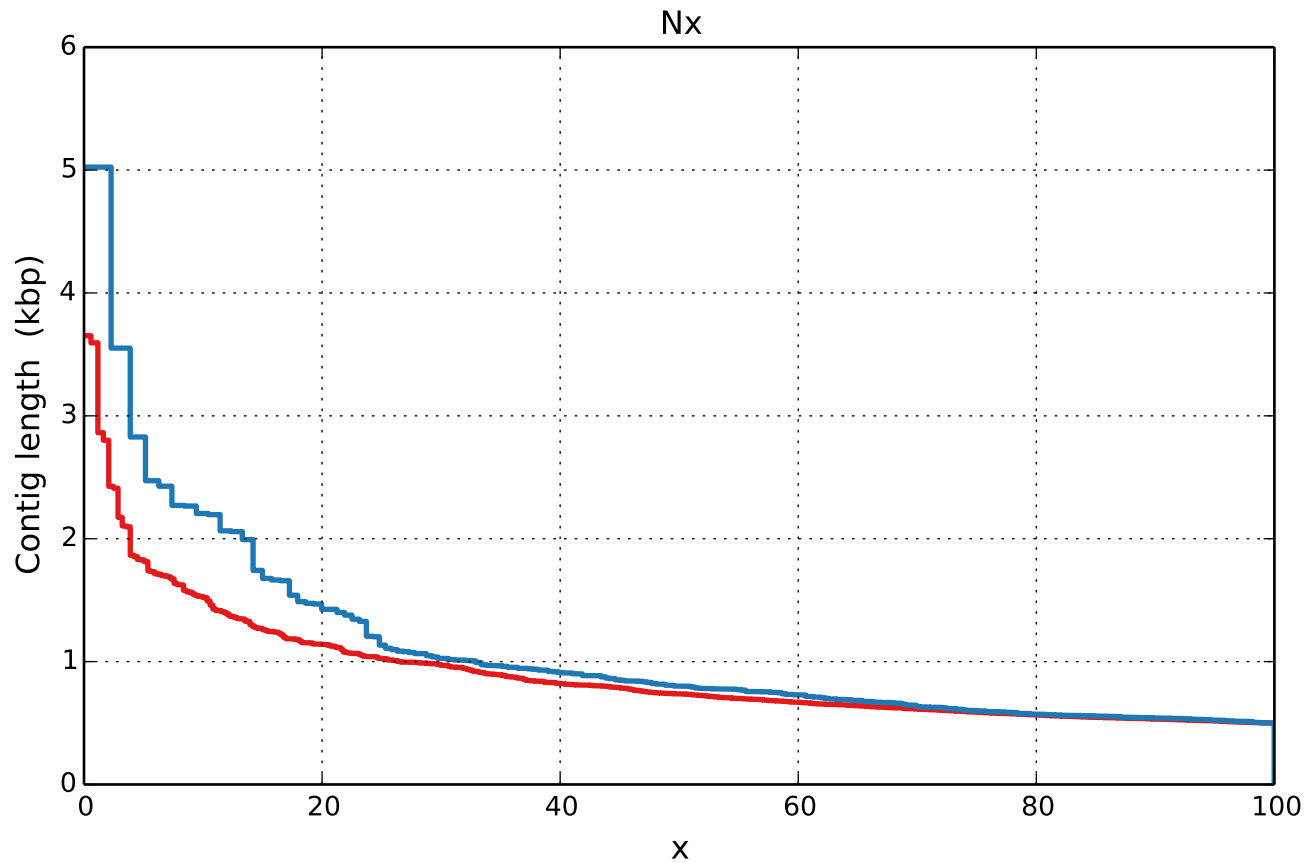


— 20150707_DNASeq_PE_P_kings-unitigs — 20150721_DNASeq_PE_P_kings-unitigs

GC content



— 20150707_DNASeq_PE_P_kings-unitigs — 20150721_DNASeq_PE_P_kings-unitigs



— 20150707_DNASeq_PE_P_kings-unitigs — 20150721_DNASeq_PE_P_kings-unitigs