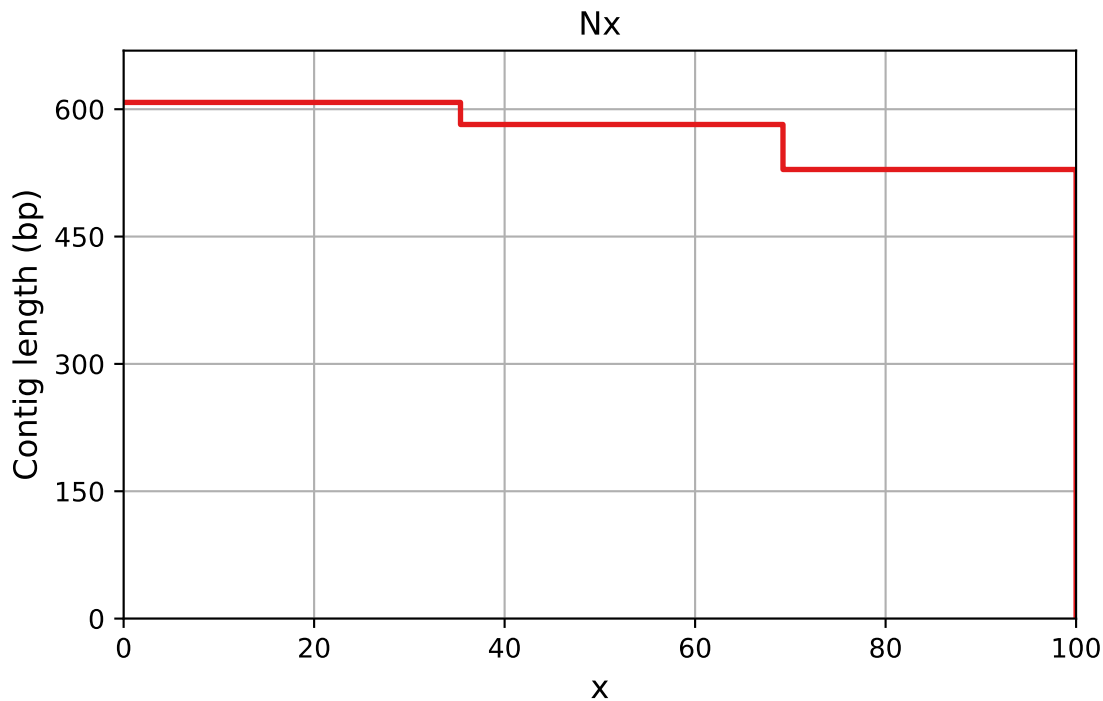


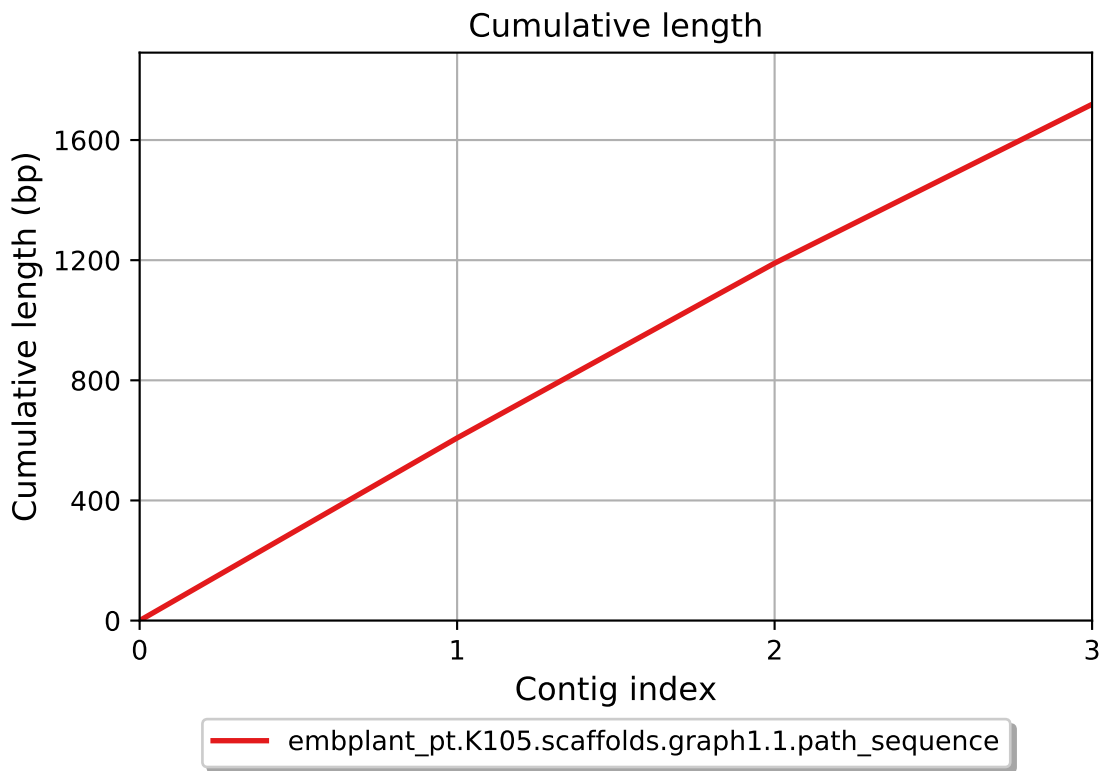
Report

	embplant_pt.K105.scaffolds.graph1.1.path_sequence
# contigs (>= 0 bp)	12
# contigs (>= 1000 bp)	0
# contigs (>= 5000 bp)	0
# contigs (>= 10000 bp)	0
# contigs (>= 25000 bp)	0
# contigs (>= 50000 bp)	0
Total length (>= 0 bp)	4746
Total length (>= 1000 bp)	0
Total length (>= 5000 bp)	0
Total length (>= 10000 bp)	0
Total length (>= 25000 bp)	0
Total length (>= 50000 bp)	0
# contigs	3
Largest contig	608
Total length	1719
GC (%)	24.55
N50	582
N75	529
L50	2
L75	3
# N's per 100 kbp	0.00

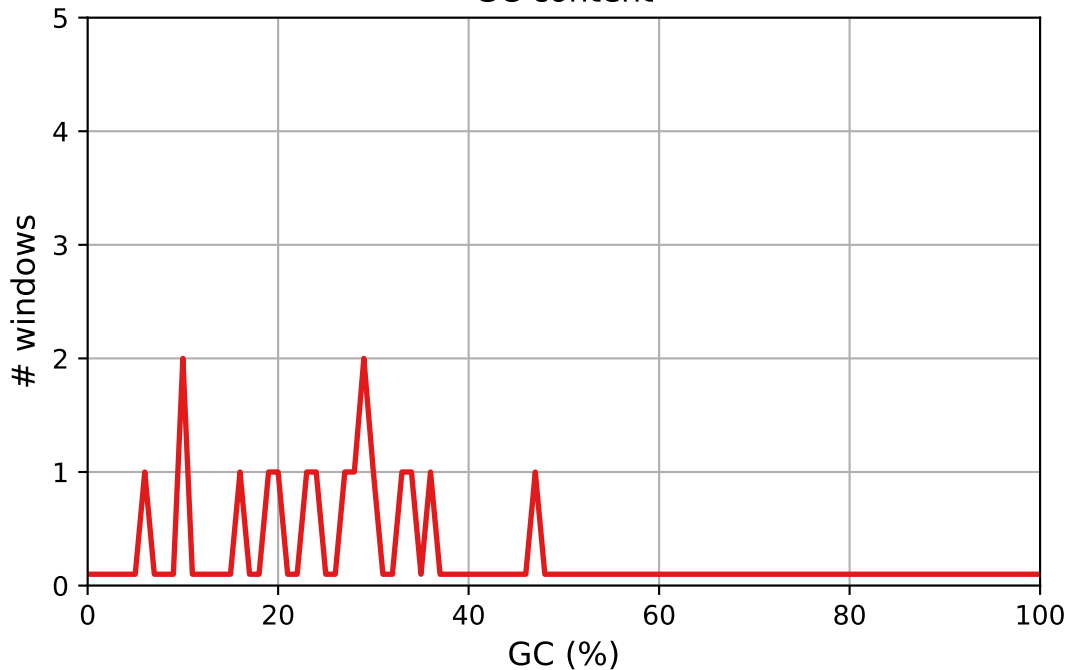
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— embplant_pt.K105.scaffolds.graph1.1.path_sequence

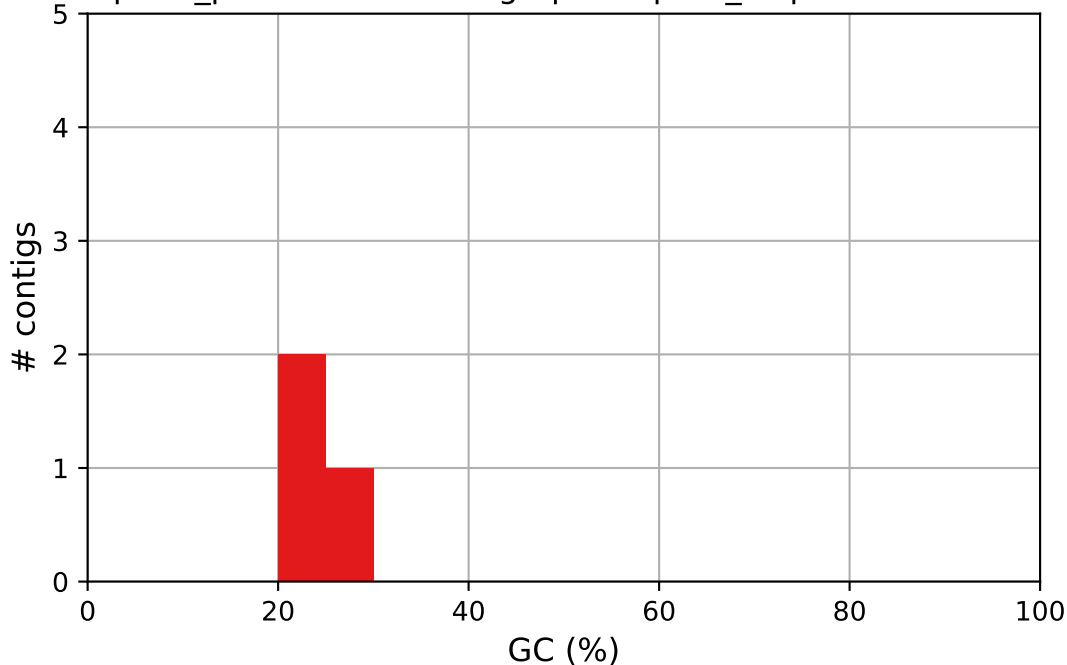


GC content



— embplant_pt.K105.scaffolds.graph1.1.path_sequence

embplant_pt.K105.scaffolds.graph1.1.path_sequence GC content



embplant_pt.K105.scaffolds.graph1.1.path_sequence