

Report

	lferriphilum_1contig
# contigs (>= 0 bp)	1
# contigs (>= 1000 bp)	1
# contigs (>= 5000 bp)	1
# contigs (>= 10000 bp)	1
# contigs (>= 25000 bp)	1
# contigs (>= 50000 bp)	1
Total length (>= 0 bp)	2577733
Total length (>= 1000 bp)	2577733
Total length (>= 5000 bp)	2577733
Total length (>= 10000 bp)	2577733
Total length (>= 25000 bp)	2577733
Total length (>= 50000 bp)	2577733
# contigs	1
Largest contig	2577733
Total length	2577733
Reference length	2610531
GC (%)	54.22
Reference GC (%)	54.14
N50	2577733
NG50	2577733
N75	2577733
NG75	2577733
L50	1
LG50	1
L75	1
LG75	1
# misassemblies	2
# misassembled contigs	1
Misassembled contigs length	2577733
# local misassemblies	0
# unaligned mis. contigs	0
# unaligned contigs	0 + 0 part
Unaligned length	0
Genome fraction (%)	97.566
Duplication ratio	1.012
# N's per 100 kbp	0.00
# mismatches per 100 kbp	0.12
# indels per 100 kbp	3.57
Largest alignment	1921378
Total aligned length	2577692
NA50	1921378
NGA50	1921378
NA75	644322
NGA75	644322
LA50	1
LGA50	1
LA75	2
LGA75	2

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).

Misassemblies report

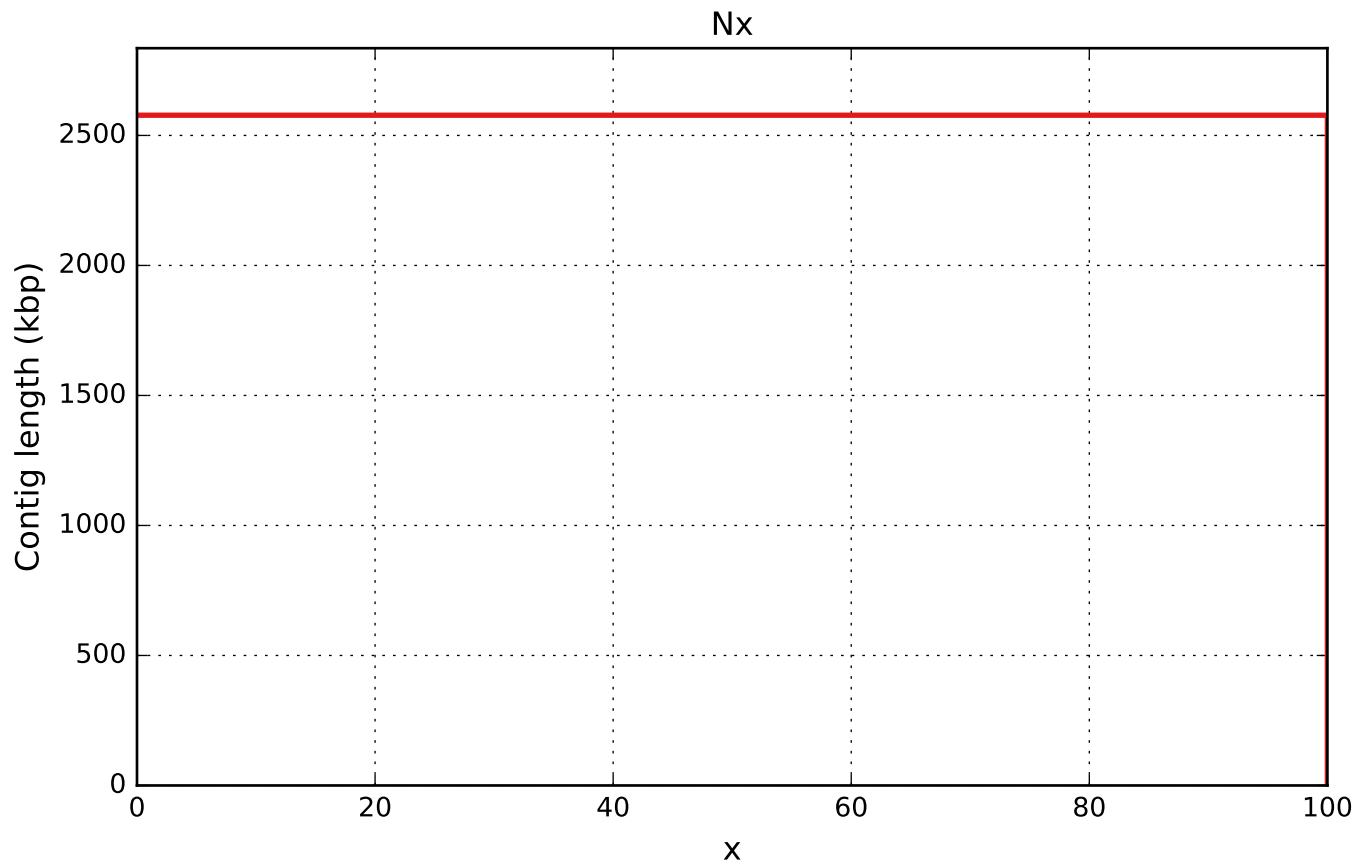
	lferriphilum_1contig
# misassemblies	2
# relocations	2
# translocations	0
# inversions	0
# misassembled contigs	1
Misassembled contigs length	2577733
# local misassemblies	0
# unaligned mis. contigs	0
# mismatches	3
# indels	91
# indels (<= 5 bp)	91
# indels (> 5 bp)	0
Indels length	93

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).

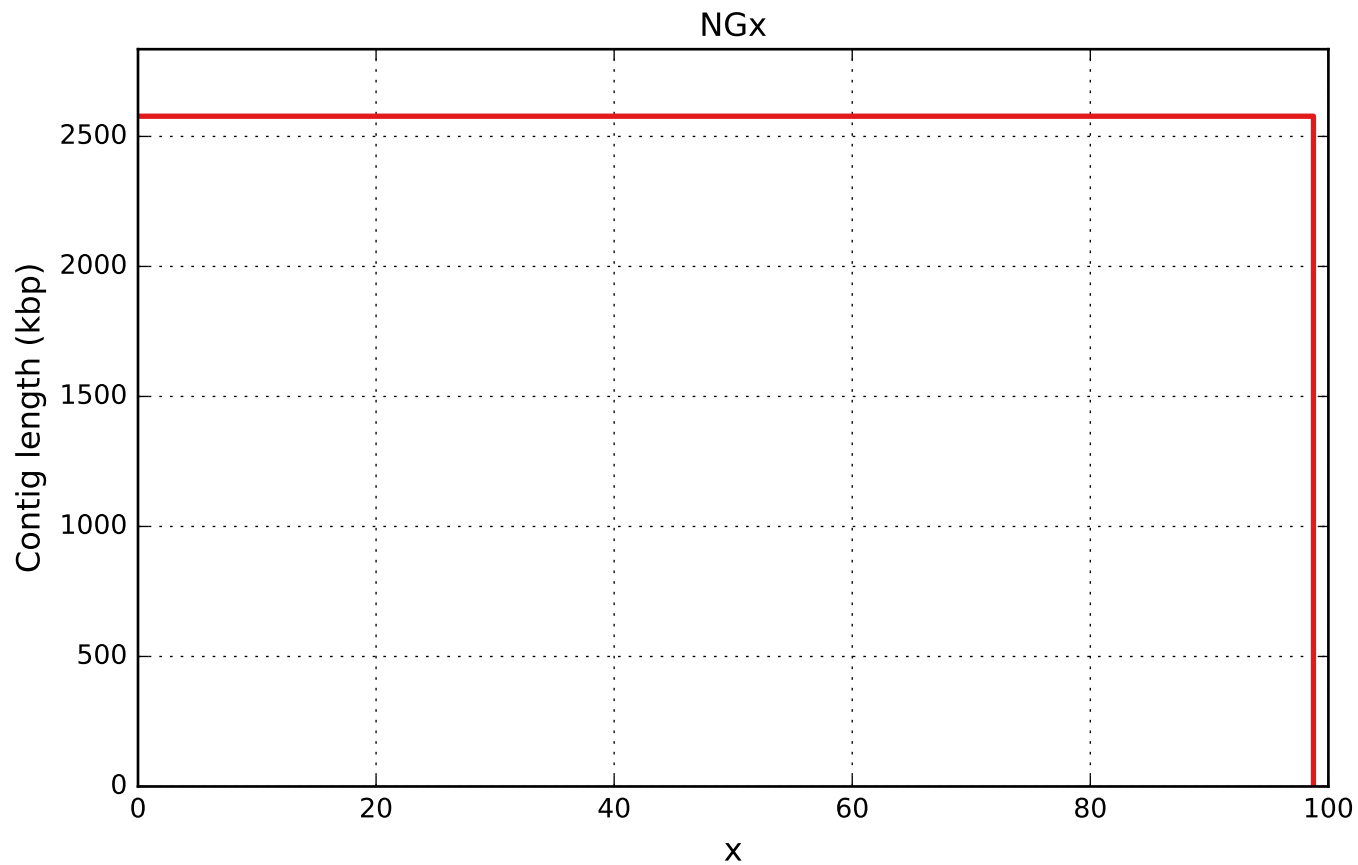
Unaligned report

	lferriphilum_1contig
# fully unaligned contigs	0
Fully unaligned length	0
# partially unaligned contigs	0
Partially unaligned length	0
# N's	0

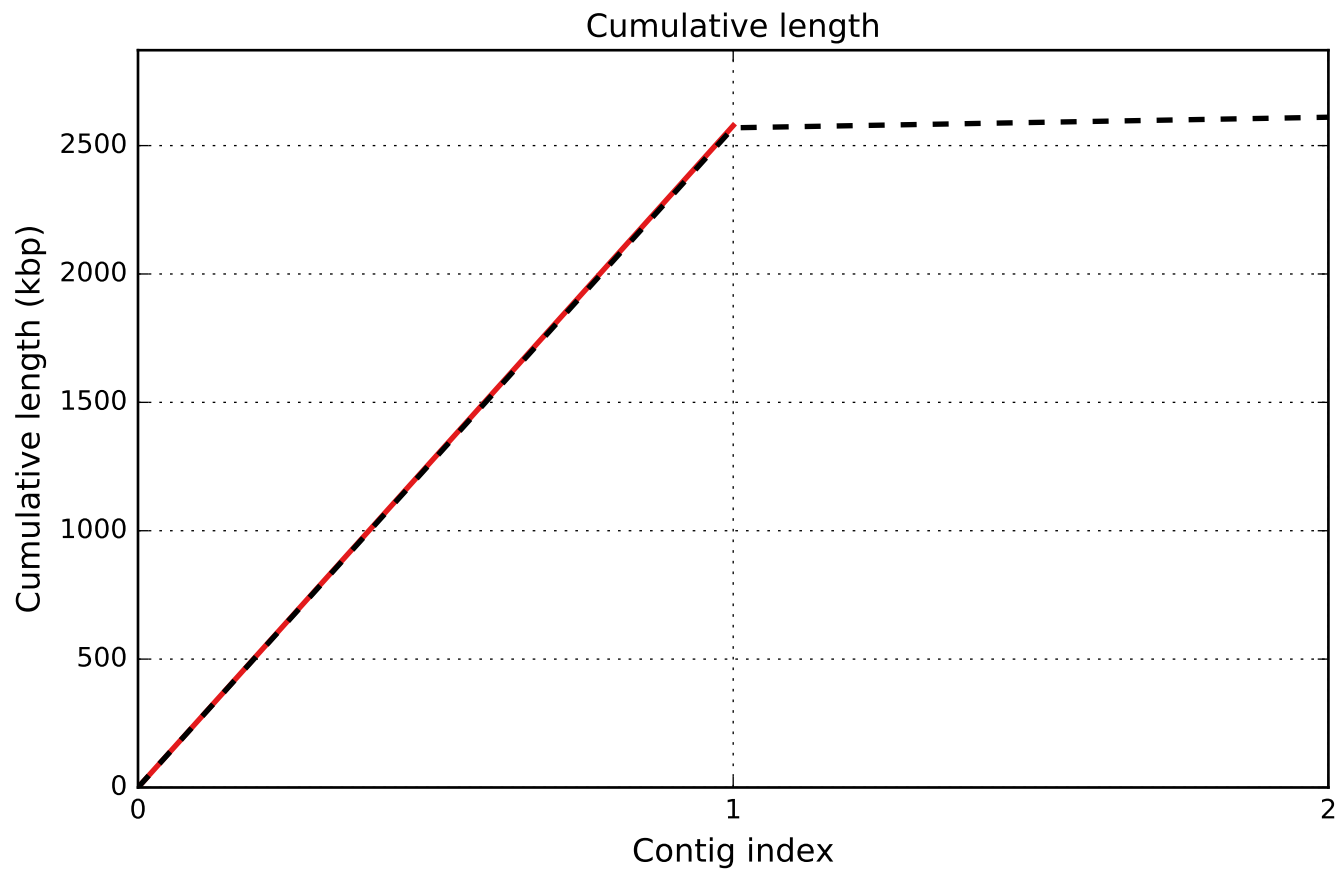
All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).



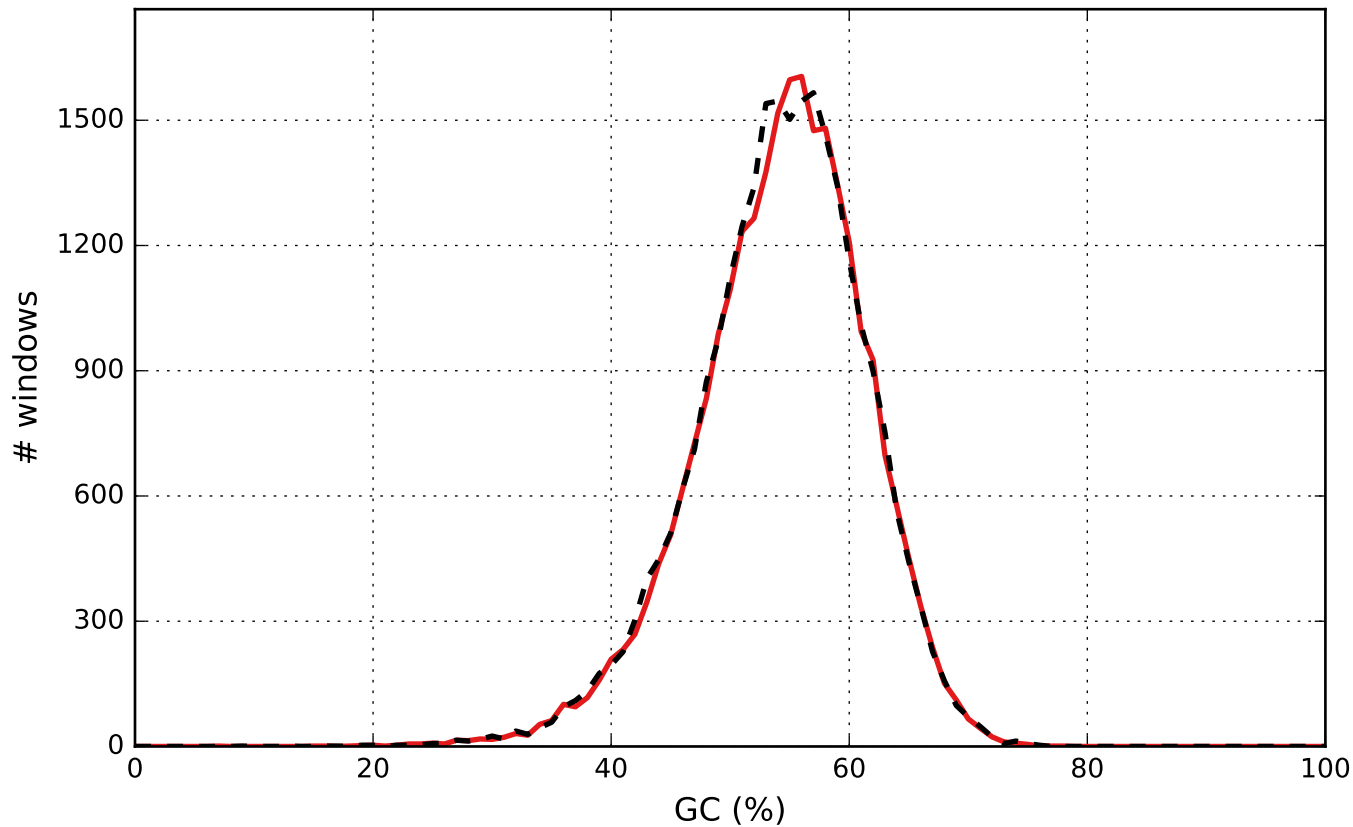
— Iferriphilum_1contig



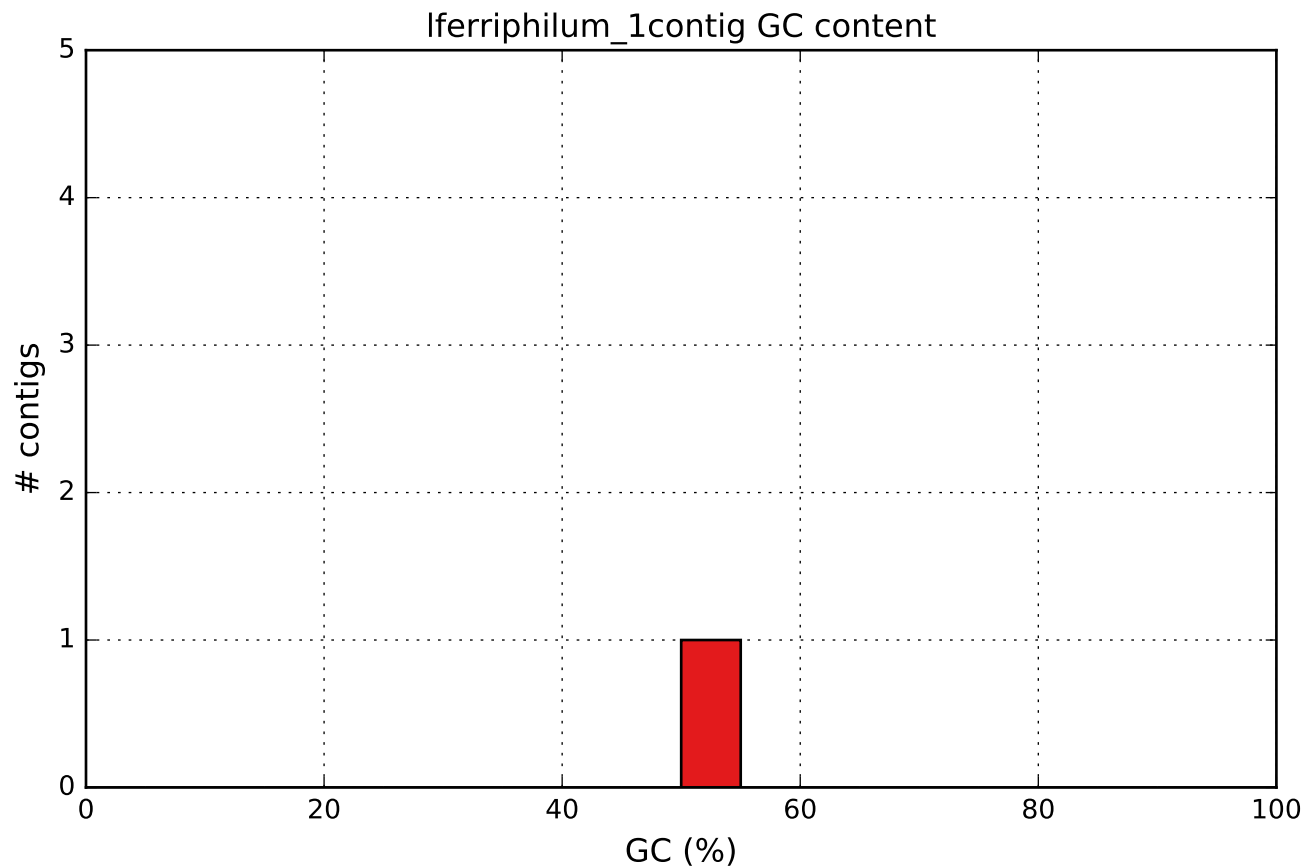
— Iferripphilum_1contig



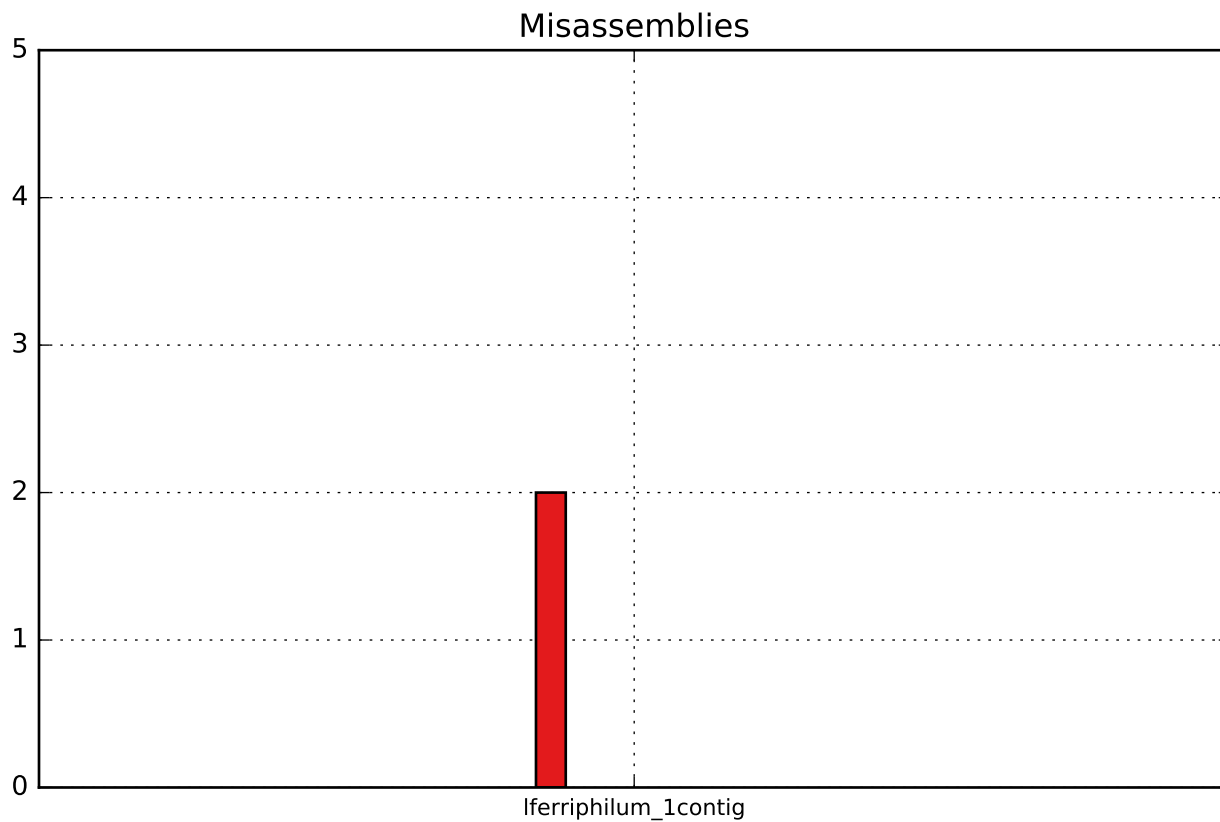
GC content



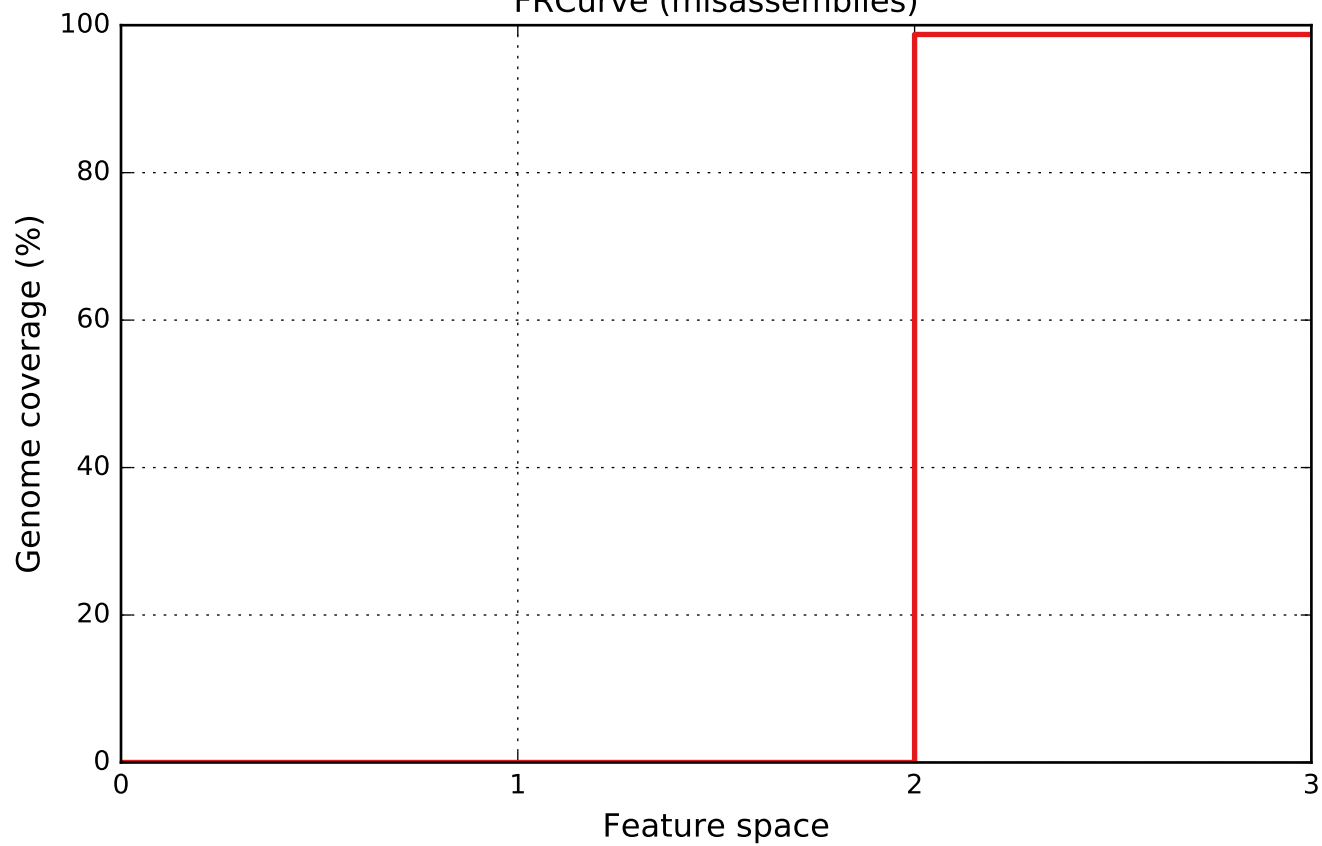
— *Iferriphilum_1contig* - - Reference



lferriphilum_1contig

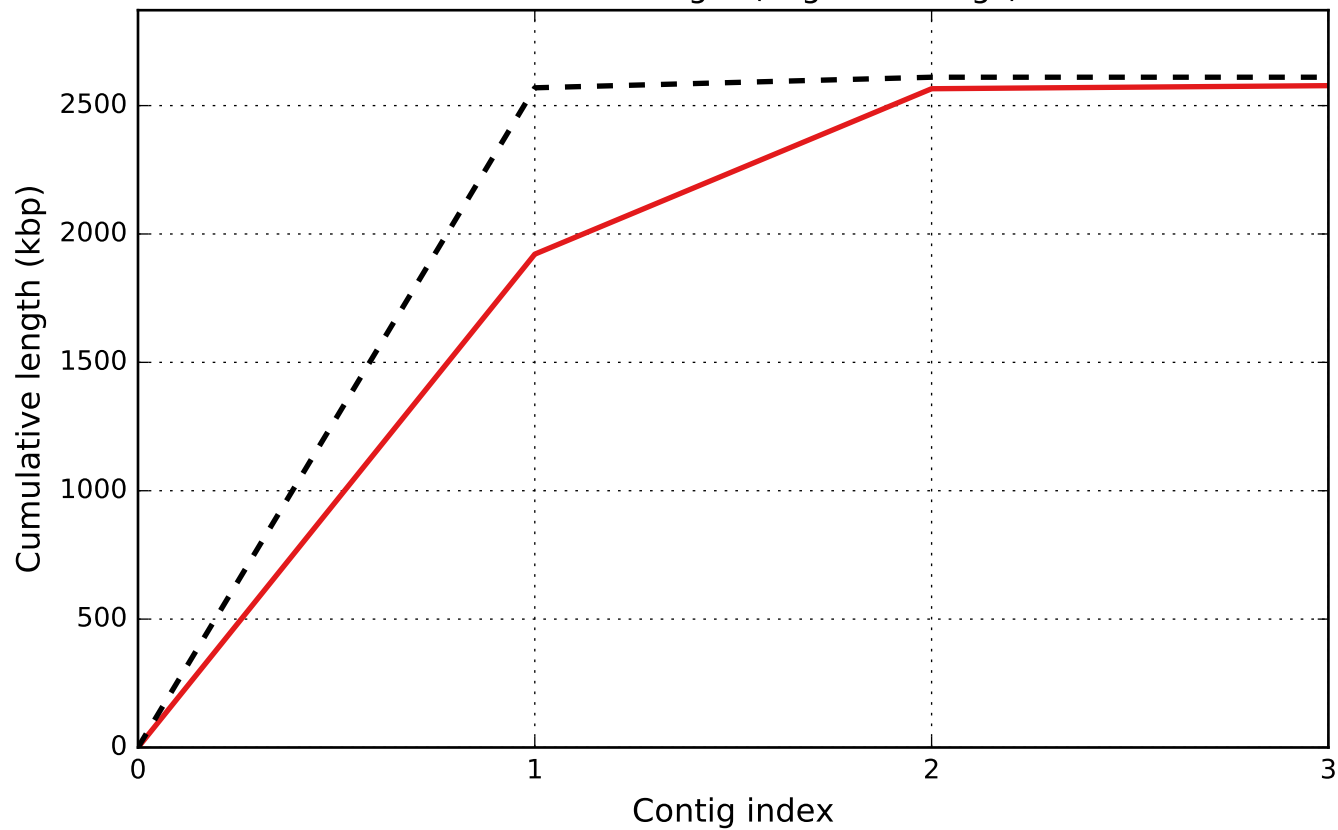


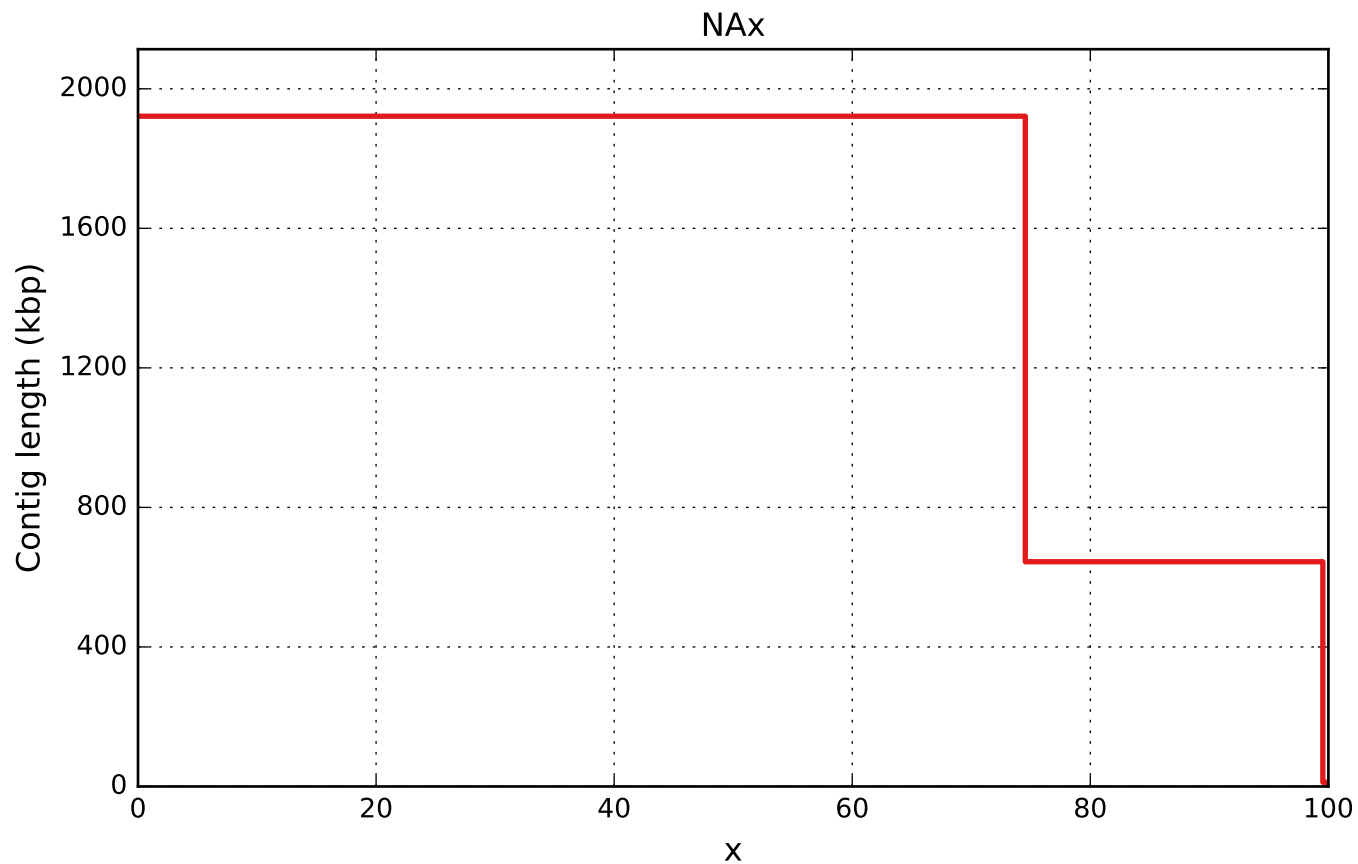
FRCurve (misassemblies)



— Iferriphilum_1contig

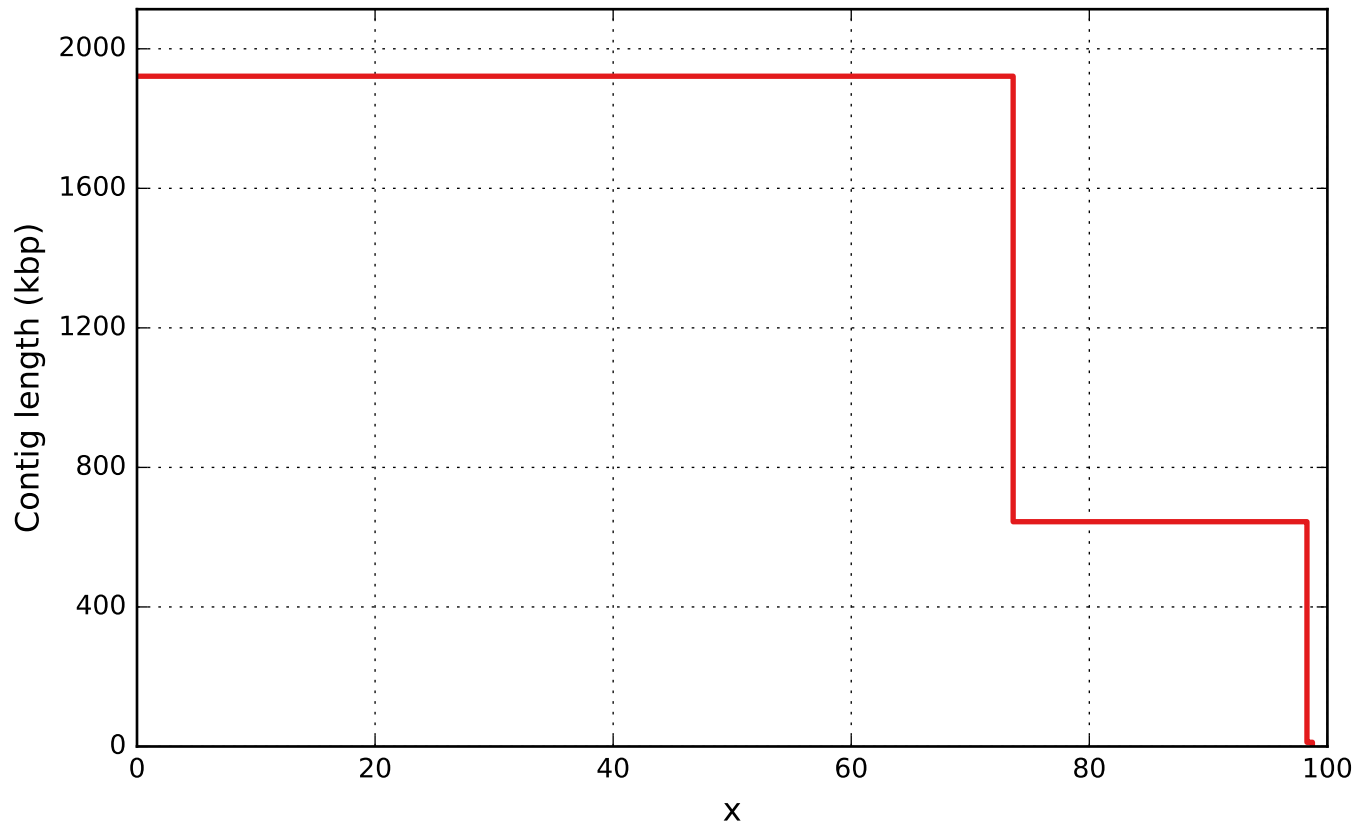
Cumulative length (aligned contigs)





— Iferriphilum_1contig

NGAx



— Iferriphilum_1contig