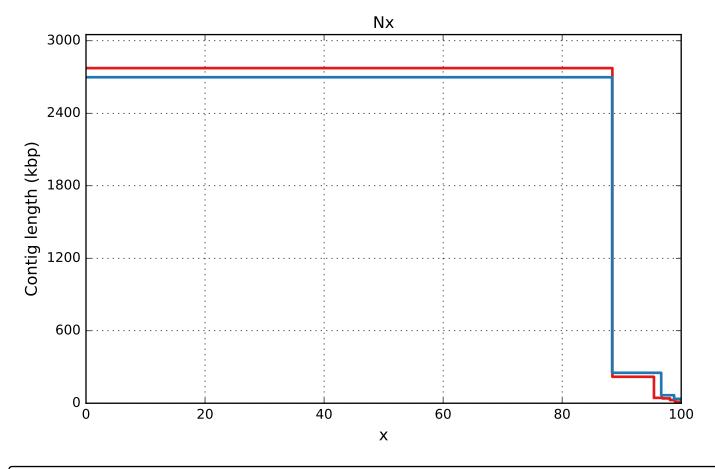
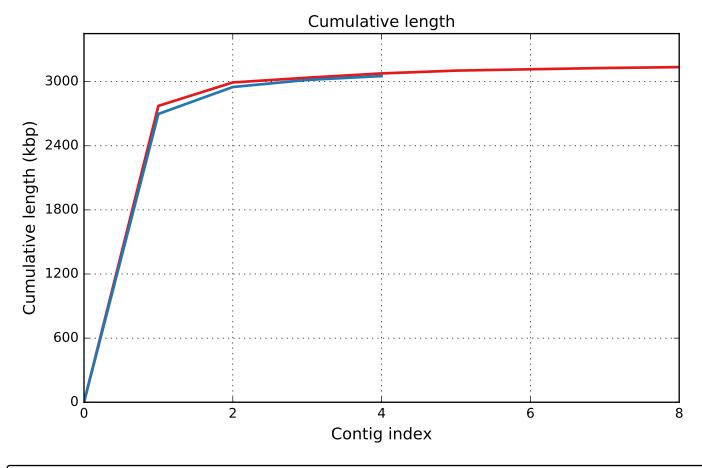
Report

| | 190405_pilon_illumina_to_pacbio | GCF_000174395.2_ASM17439v2_genomic |
|----------------------------|---------------------------------|------------------------------------|
| # contigs (>= 0 bp) | 8 | 4 |
| # contigs (>= 1000 bp) | 8 | 4 |
| # contigs (>= 5000 bp) | 8 | 4 |
| # contigs (>= 10000 bp) | 7 | 4 |
| # contigs (>= 25000 bp) | 5 | 4 |
| # contigs (>= 50000 bp) | 2 | 3 |
| Total length (>= 0 bp) | 3136314 | 3052572 |
| Total length (>= 1000 bp) | 3136314 | 3052572 |
| Total length (>= 5000 bp) | 3136314 | 3052572 |
| Total length (>= 10000 bp) | 3128014 | 3052572 |
| Total length (>= 25000 bp) | 3103592 | 3052572 |
| Total length (>= 50000 bp) | 2992679 | 3016310 |
| # contigs | 8 | 4 |
| Largest contig | 2773702 | 2698137 |
| Total length | 3136314 | 3052572 |
| GC (%) | 37.79 | 37.87 |
| N50 | 2773702 | 2698137 |
| N75 | 2773702 | 2698137 |
| L50 | 1 | 1 |
| L75 | 1 | 1 |
| # N's per 100 kbp | 0.00 | 0.00 |

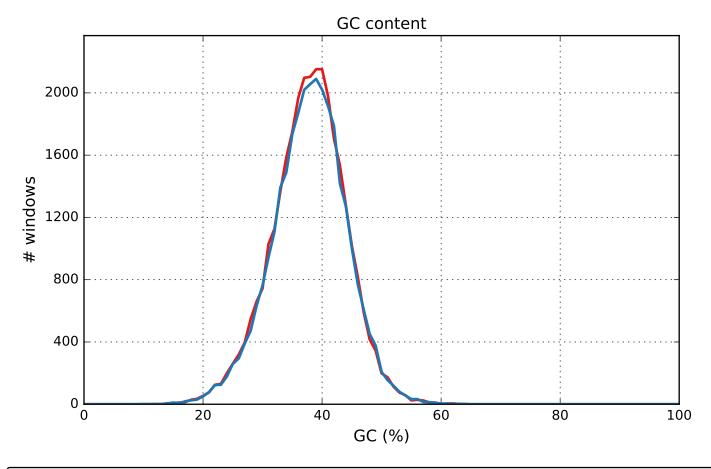
All statistics are based on contigs of size \geq 500 bp, unless otherwise noted (e.g., "# contigs (\geq 0 bp)" and "Total length (\geq 0 bp)" include all contigs).



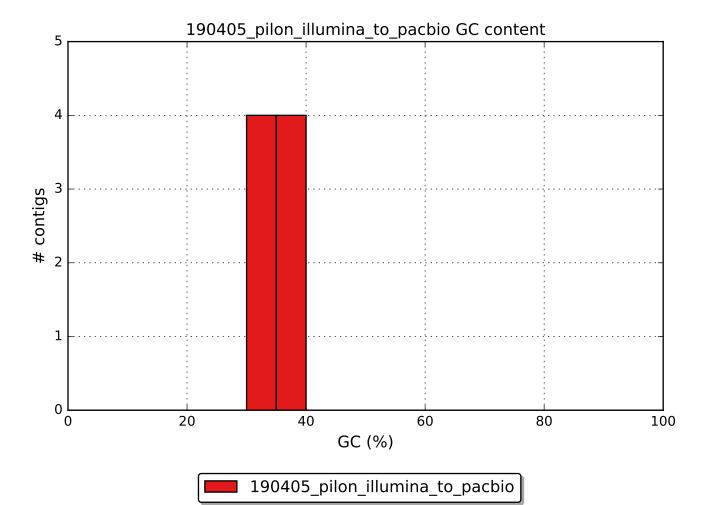
— 190405_pilon_illumina_to_pacbio — GCF_000174395.2_ASM17439v2_genomic

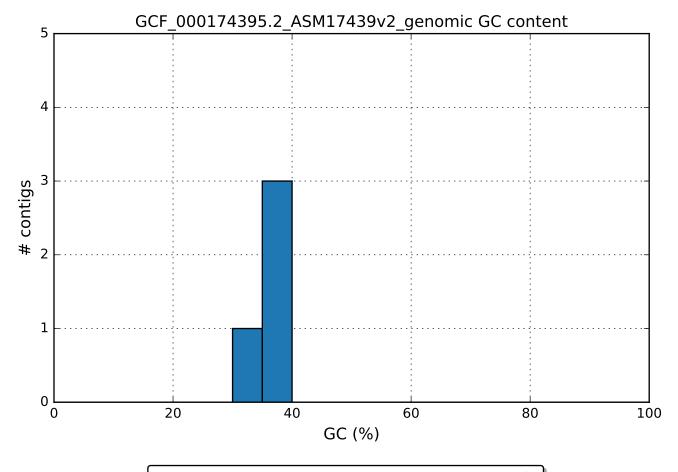


— 190405_pilon_illumina_to_pacbio — GCF_000174395.2_ASM17439v2_genomic



— 190405_pilon_illumina_to_pacbio — GCF_000174395.2_ASM17439v2_genomic





GCF_000174395.2_ASM17439v2_genomic