

| | MP_helen_r10_bins_1 | MP_helen_r10_bins_2 | MP_helen_r10_bins_3 | MP_r10_bins_1 | MP_r10_bins_2 | MP_r10_bins_3 | r1_medaka_r10_bins_1 | r1_medaka_r10_bins_2 | r1_medaka_r10_bins_3 | r2_medaka_r10_bins_1 | r2_medaka_r10_bins_2 | r2_medaka_r10_bins_3 | racon_r1_r10_bins_1 | racon_r1_r10_bins_2 | racon_r1_r10_bins_3 | racon_r2_r10_bins_1 | racon_r2_r10_bins_2 | racon_r2_r10_bins_3 | raw_r10_bins_1 | raw_r10_bins_2 | raw_r10 | |
|-----------------------------|---------------------|---------------------|---------------------|---------------|---------------|---------------|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|----------------|----------------|----------|----------|
| # contigs (>= 5000 bp) | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 |
| # contigs (>= 10000 bp) | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 |
| # contigs (>= 25000 bp) | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 |
| # contigs (>= 50000 bp) | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 |
| Total length (>= 5000 bp) | 24069494 | 24072731 | 24070272 | 24075146 | 24075931 | 24075224 | 24072592 | 24074864 | 24072788 | 24064948 | 24068153 | 24065807 | 24059729 | 24064644 | 24059333 | 24055079 | 24060632 | 24052503 | 24060236 | 24062591 | 24060236 | 24062591 |
| Total length (>= 10000 bp) | 24069494 | 24072731 | 24070272 | 24075146 | 24075931 | 24075224 | 24072592 | 24074864 | 24072788 | 24064948 | 24068153 | 24065807 | 24059729 | 24064644 | 24059333 | 24055079 | 24060632 | 24052503 | 24060236 | 24062591 | 24060236 | 24062591 |
| Total length (>= 25000 bp) | 24069494 | 24072731 | 24070272 | 24075146 | 24075931 | 24075224 | 24072592 | 24074864 | 24072788 | 24064948 | 24068153 | 24065807 | 24059729 | 24064644 | 24059333 | 24055079 | 24060632 | 24052503 | 24060236 | 24062591 | 24060236 | 24062591 |
| Total length (>= 50000 bp) | 24069494 | 24072731 | 24070272 | 24075146 | 24075931 | 24075224 | 24072592 | 24074864 | 24072788 | 24064948 | 24068153 | 24065807 | 24059729 | 24064644 | 24059333 | 24055079 | 24060632 | 24052503 | 24060236 | 24062591 | 24060236 | 24062591 |
| # contigs | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 | 7 |
| Largest contig | 4765299 | 4765370 | 4765301 | 4765922 | 4765391 | 4765920 | 4765362 | 4765395 | 4765360 | 4765364 | 4765350 | 4764643 | 4764770 | 4764652 | 4764716 | 4764723 | 4764656 | 4763471 | 4763493 | 4763493 | 4763493 | 4763493 |
| Total length | 24069494 | 24072731 | 24070272 | 24075146 | 24075931 | 24075224 | 24072592 | 24074864 | 24072788 | 24064948 | 24068153 | 24065807 | 24059729 | 24064644 | 24059333 | 24055079 | 24060632 | 24052503 | 24060236 | 24062591 | 24060236 | 24062591 |
| Reference length | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | 2820294 | |
| GC (%) | 44.80 | 44.81 | 44.80 | 44.80 | 44.80 | 44.80 | 44.80 | 44.80 | 44.80 | 44.80 | 44.80 | 44.80 | 44.80 | 44.80 | 44.80 | 44.80 | 44.80 | 44.77 | 44.77 | 44.77 | 44.77 | |
| Reference GC (%) | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | 37.44 | |
| N50 | 4045615 | 4045634 | 4045636 | 4045594 | 4045608 | 4045601 | 4045589 | 4045607 | 4045601 | 4045589 | 4045605 | 4045604 | 4045233 | 4045287 | 4045256 | 4045364 | 4045333 | 4045318 | 4042946 | 4043032 | 4043032 | 4043032 |
| NG50 | 4765299 | 4765370 | 4765301 | 4765922 | 4765391 | 4765920 | 4765362 | 4765395 | 4765360 | 4765364 | 4765350 | 4764643 | 4764770 | 4764652 | 4764716 | 4764723 | 4764656 | 4763471 | 4763493 | 4763493 | 4763493 | 4763493 |
| N75 | 2845363 | 2845435 | 2845364 | 2845421 | 2845426 | 2845426 | 2845430 | 2845432 | 2845434 | 2845429 | 2845433 | 2845432 | 2845290 | 2845293 | 2845290 | 2845315 | 2845310 | 2845306 | 2843359 | 2843856 | 2843856 | 2843856 |
| NG75 | 4765299 | 4765370 | 4765301 | 4765922 | 4765391 | 4765920 | 4765362 | 4765395 | 4765360 | 4765364 | 4765350 | 4764643 | 4764770 | 4764652 | 4764716 | 4764723 | 4764656 | 4763471 | 4763493 | 4763493 | 4763493 | 4763493 |
| L50 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | |
| LG50 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| L75 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 |
| LG75 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| # misassemblies | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| # misassembled contigs | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| Misassembled contigs length | 2845363 | 2845435 | 2845364 | 2845421 | 2845426 | 2845426 | 2845430 | 2845432 | 2845434 | 2845429 | 2845433 | 2845432 | 2845290 | 2845293 | 2845290 | 2845315 | 2845310 | 2845306 | 2843359 | 2843856 | 2843856 | 2843856 |
| # local misassemblies | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 21 | 20 | 20 | 20 |
| # scaffold gap ext. mis. | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| # scaffold gap loc. mis. | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| # unaligned mis. contigs | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 |
| # unaligned contigs | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 6 part | 0 + 7 part | 0 + 7 part | 0 | 0 | |
| Unaligned length | 21091102 | 21095401 | 21091844 | 21098041 | 21098126 | 21097410 | 21094241 | 21096364 | 21095648 | 21086969 | 21089492 | 21090040 | 21086435 | 21093151 | 21087019 | 21079298 | 21083038 | 21078060 | 21150146 | 21154353 | 2 | 2 |
| Genome fraction (%) | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.954 | 99.931 | 99.948 | 99.948 | 99.948 | |
| Duplication ratio | 1.057 | 1.056 | 1.057 | 1.056 | 1.057 | 1.057 | 1.057 | 1.057 | 1.057 | 1.056 | 1.057 | 1.057 | 1.056 | 1.055 | 1.054 | 1.055 | 1.056 | 1.053 | 1.032 | 1.032 | 1.032 | |
| | | | | | | | | | | | | | | | | | | | | | | |

All statistics are based on contigs of size \geq 5000 bp, unless otherwise noted
(e.g., "# contigs (\geq 0 bp)" and "Total length (\geq 0 bp)" include all contigs).

Misassemblies report

| | MP_helen_r10_bins_1 | MP_helen_r10_bins_2 | MP_helen_r10_bins_3 | MP_r10_bins_1 | MP_r10_bins_2 | MP_r10_bins_3 | r1_medaka_r10_bins_1 | r1_medaka_r10_bins_2 | r1_medaka_r10_bins_3 | r2_medaka_r10_bins_1 | r2_medaka_r10_bins_2 | r2_medaka_r10_bins_3 | racon_r1_r10_bins_1 | racon_r1_r10_bins_2 | racon_r1_r10_bins_3 | racon_r2_r10_bins_1 | racon_r2_r10_bins_2 | racon_r2_r10_bins_3 | raw_r10_bins_1 | raw_r10_bins_2 | raw_r10_bins_3 |
|--|---------------------|---------------------|---------------------|---------------|---------------|---------------|----------------------|----------------------|----------------------|----------------------|----------------------|----------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|----------------|----------------|----------------|
| # misassemblies | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| # contig misassemblies | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| # c. relocations | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| # c. translocations | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| # c. inversions | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| # scaffold misassemblies | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| # s. relocations | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| # s. translocations | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| # s. inversions | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| # misassembled contigs | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| Misassembled contigs length | 2845363 | 2845435 | 2845364 | 2845421 | 2845426 | 2845426 | 2845430 | 2845432 | 2845434 | 2845429 | 2845433 | 2845432 | 2845290 | 2845293 | 2845290 | 2845315 | 2845310 | 2845306 | 2843359 | 2843856 | 2843855 |
| # possibly misassembled contigs | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 | 1 | 1 |
| # possible misassemblies | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 6 | 6 | 6 |
| # local misassemblies | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 16 | 21 | 20 | 20 |
| # scaffold gap ext. mis. | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| # scaffold gap loc. mis. | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| # misassemblies caused by fragmented reference | 19 | 19 | 19 | 19 | 19 | 19 | 19 | 19 | 19 | 19 | 19 | 19 | 19 | 19 | 19 | 19 | 19 | 11 | 10 | 12 | |
| # unaligned mis. contigs | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 |
| # mismatches | 7116 | 7013 | 7130 | 7004 | 7058 | 7061 | 7082 | 7120 | 6974 | 7059 | 7132 | 6861 | 6608 | 6435 | 6479 | 6874 | 7045 | 6761 | 4268 | 4140 | 3934 |
| # indels | 438 | 380 | 444 | 442 | 430 | 429 | 397 | 410 | 406 | 389 | 416 | 400 | 605 | 582 | 557 | 600 | 635 | 604 | 3556 | 3565 | 3513 |
| # indels (<= 5 bp) | 394 | 340 | 403 | 401 | 389 | 389 | 356 | 371 | 366 | 348 | 375 | 361 | 564 | 545 | 522 | 563 | 596 | 567 | 3520 | 3527 | 3479 |
| # indels (> 5 bp) | 44 | 40 | 41 | 41 | 40 | 41 | 39 | 40 | 41 | 37 | 35 | 39 | 40 | 41 | 37 | 35 | 37 | 36 | 38 | 34 | |
| Indels length | 2474 | 2379 | 2417 | 2460 | 2439 | 2384 | 2363 | 2390 | 2335 | 2365 | 2425 | 2319 | 2645 | 2550 | 2492 | 2617 | 2681 | 2536 | 6136 | 6267 | 5933 |

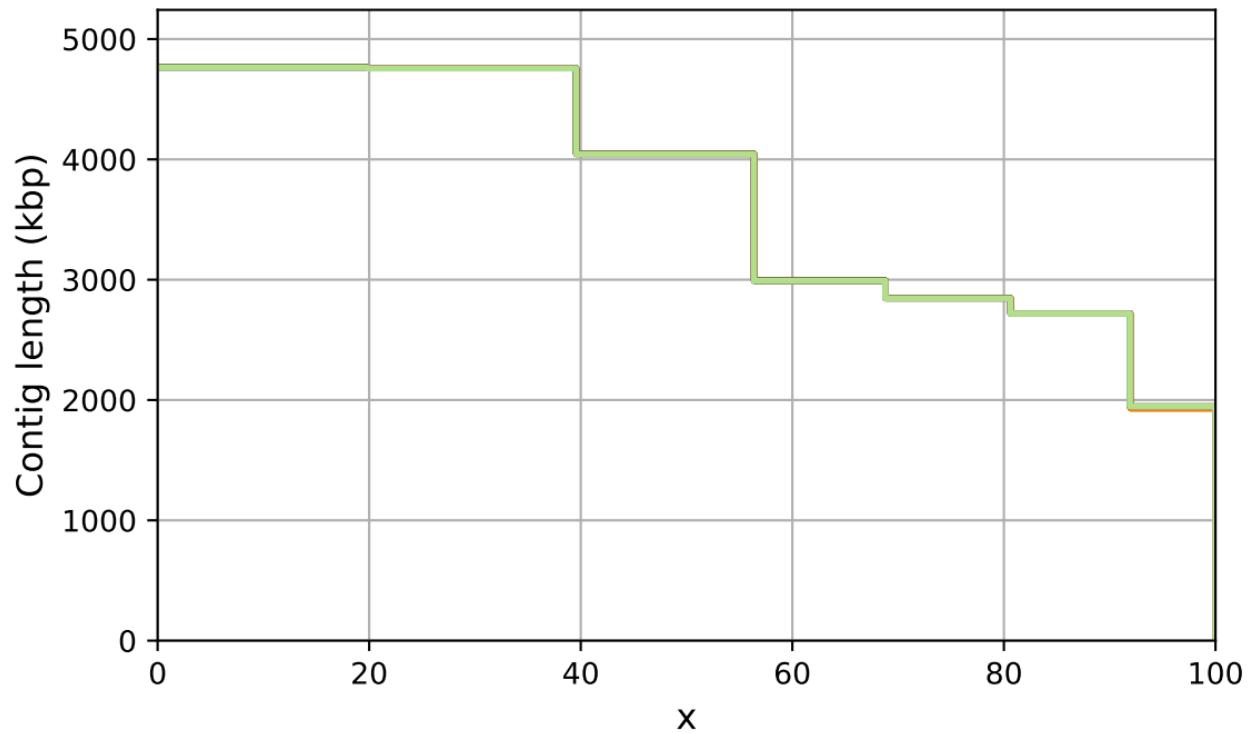
All statistics are based on contigs of size ≥ 5000 bp, unless otherwise noted
(e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Unaligned report

| | MP_helen_r10_bins_1 | MP_helen_r10_bins_2 | MP_helen_r10_bins_3 | MP_r10_bins_1 | MP_r10_bins_2 | MP_r10_bins_3 | r1_medaka_r10_bins_1 | r1_medaka_r10_bins_2 | r1_medaka_r10_bins_3 | racon_r1_r10_bins_1 | racon_r1_r10_bins_2 | racon_r1_r10_bins_3 | racon_r2_r10_bins_1 | racon_r2_r10_bins_2 | racon_r2_r10_bins_3 | raw_r10_bins_1 | raw_r10_bins_2 | raw_r10_bins_3 | |
|-------------------------------|---------------------|---------------------|---------------------|---------------|---------------|---------------|----------------------|----------------------|----------------------|---------------------|---------------------|---------------------|---------------------|---------------------|---------------------|----------------|----------------|----------------|----------|
| # fully unaligned contigs | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | |
| Fully unaligned length | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | |
| # partially unaligned contigs | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 7 | 7 | 7 | 7 | 7 | |
| Partially unaligned length | 21091102 | 21095401 | 21091844 | 21098041 | 21098126 | 21097410 | 21094241 | 21096364 | 21095648 | 21086969 | 21089492 | 21090040 | 21086435 | 21093151 | 21087019 | 21079298 | 21083038 | 21078060 | 21150146 |
| # N's | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | |

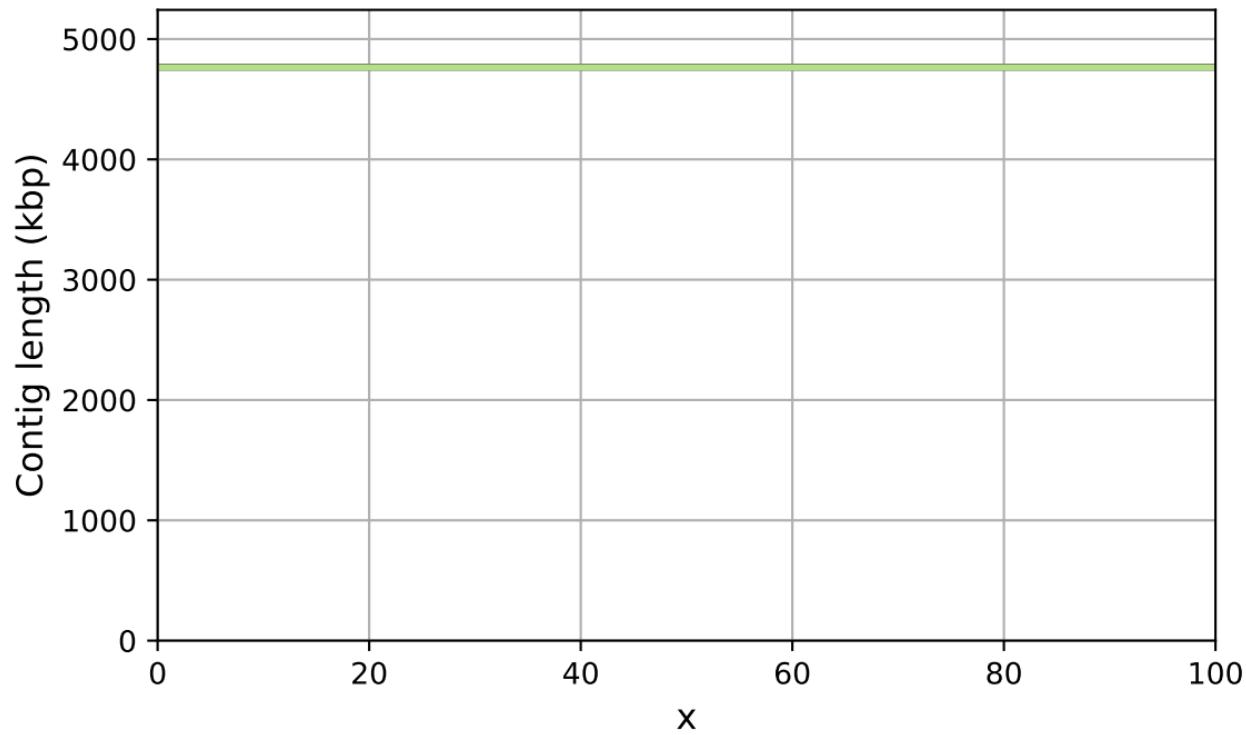
All statistics are based on contigs of size ≥ 5000 bp, unless otherwise noted
(e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Nx



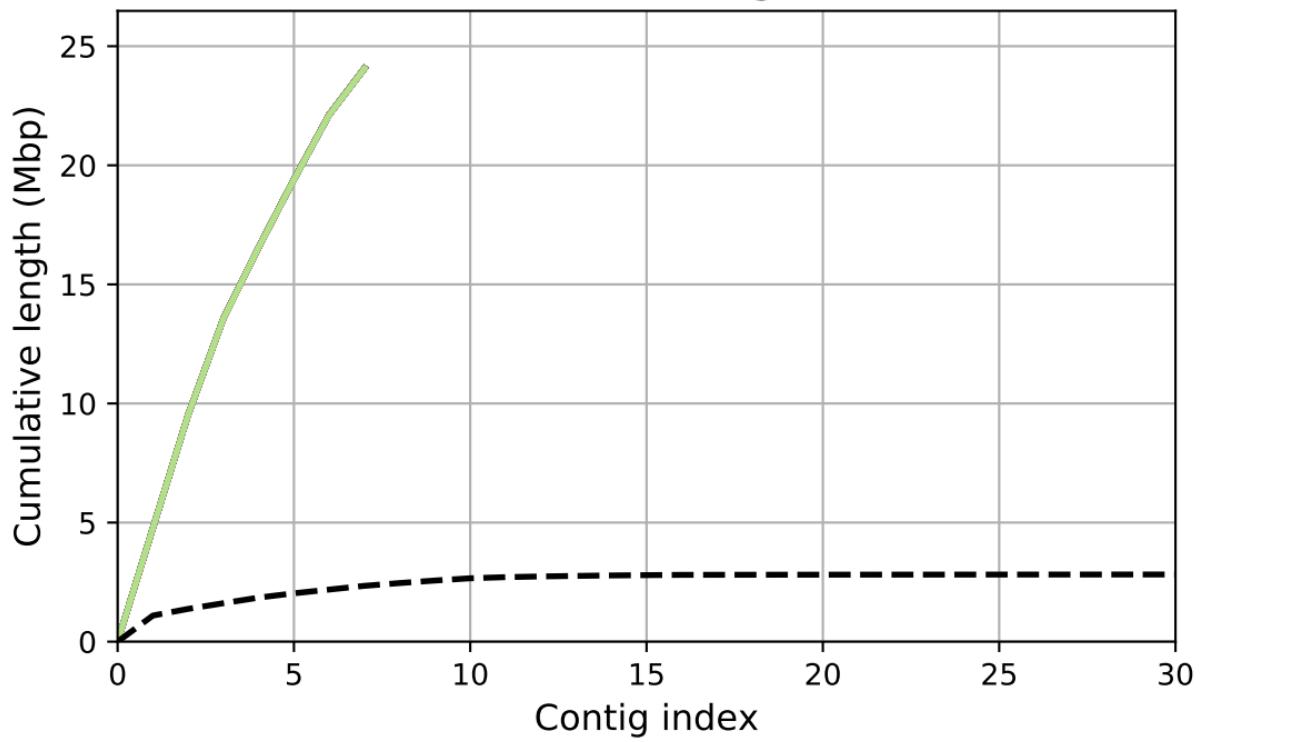
- MP_helen_r10_bins_1
- MP_helen_r10_bins_2
- MP_helen_r10_bins_3
- MP_r10_bins_1
- r1_medaka_r10_bins_2
- r1_medaka_r10_bins_3
- r2_medaka_r10_bins_1
- r2_medaka_r10_bins_2
- racon_r1_r10_bins_3
- racon_r2_r10_bins_1
- racon_r2_r10_bins_2
- racon_r2_r10_bins_3

NGx



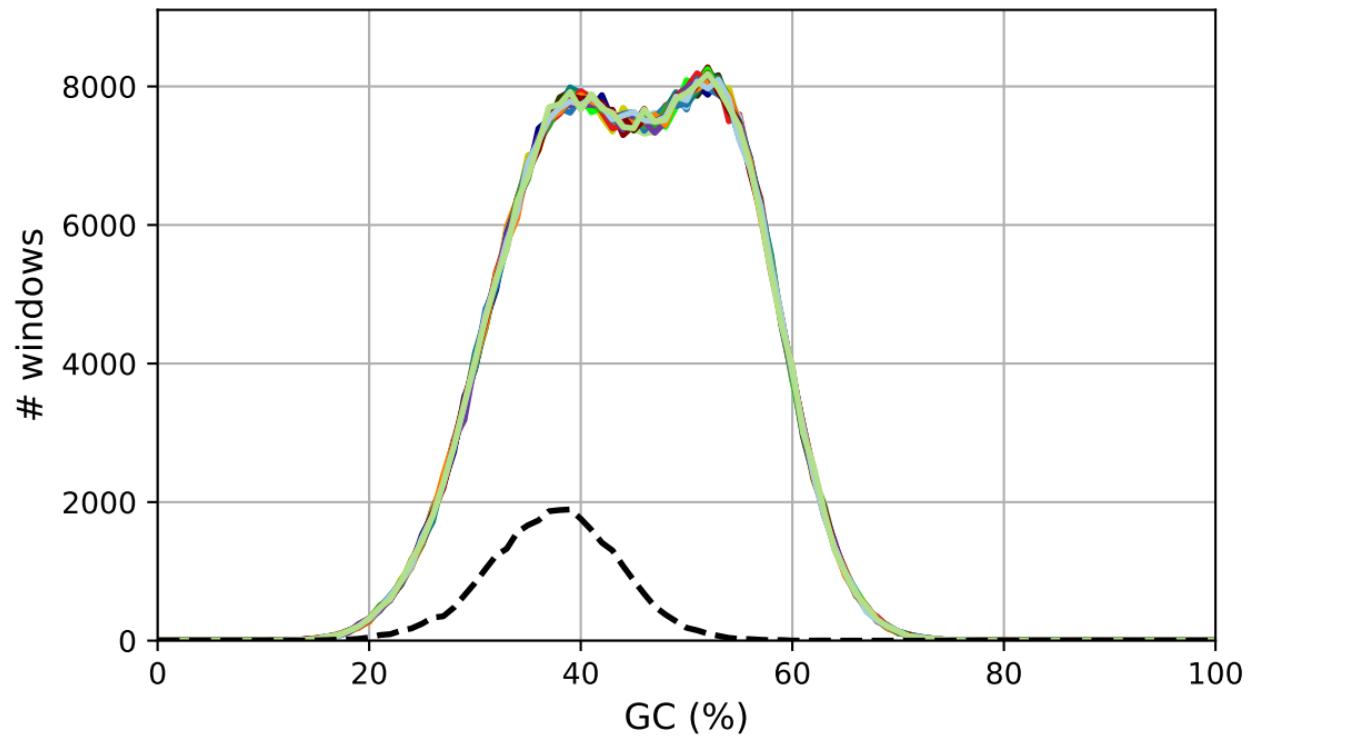
- MP_helen_r10_bins_1
- MP_helen_r10_bins_2
- MP_helen_r10_bins_3
- MP r10 bins 1
- r1_medaka_r10_bins_2
- r1_medaka_r10_bins_3
- r2_medaka_r10_bins_1
- r2_medaka_r10_bins_2
- racon_r1_r10_bins_3
- racon_r2_r10_bins_1
- racon_r2_r10_bins_2
- racon_r2_r10_bins_3

Cumulative length



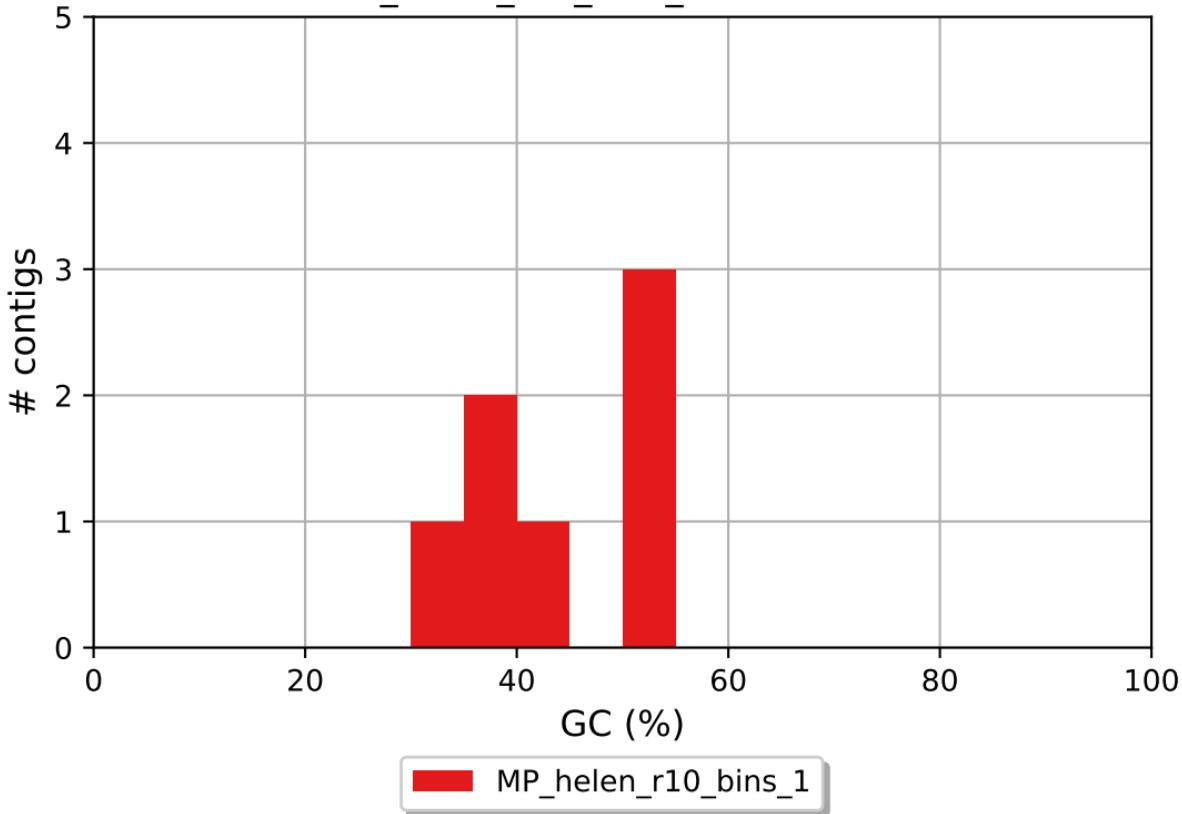
- MP_helen_r10_bins_1
- MP_helen_r10_bins_2
- MP_helen_r10_bins_3
- MP_r10_bins_1
- r1_medaka_r10_bins_3
- r2_medaka_r10_bins_1
- r2_medaka_r10_bins_2
- r2_medaka_r10_bins_3
- racon_r2_r10_bins_1
- racon_r2_r10_bins_2
- racon_r2_r10_bins_3
- raw_r10_bins_1

GC content

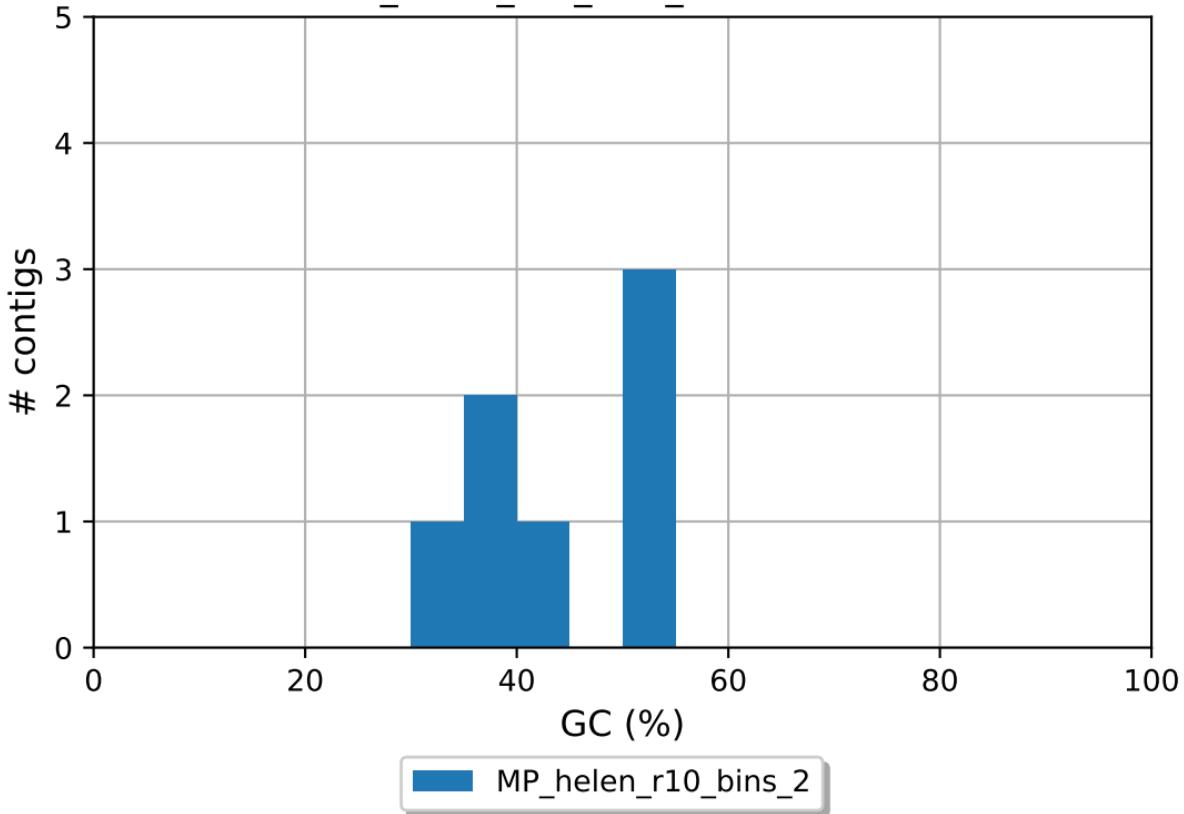


- MP_helen_r10_bins_1
- MP_helen_r10_bins_2
- MP_helen_r10_bins_3
- MP r10 bins 1
- r1_medaka_r10_bins_3
- r2_medaka_r10_bins_1
- r2_medaka_r10_bins_2
- r2_medaka_r10_bins_3
- racon_r2_r10_bins_1
- racon_r2_r10_bins_2
- racon_r2_r10_bins_3
- raw r10 bins 1

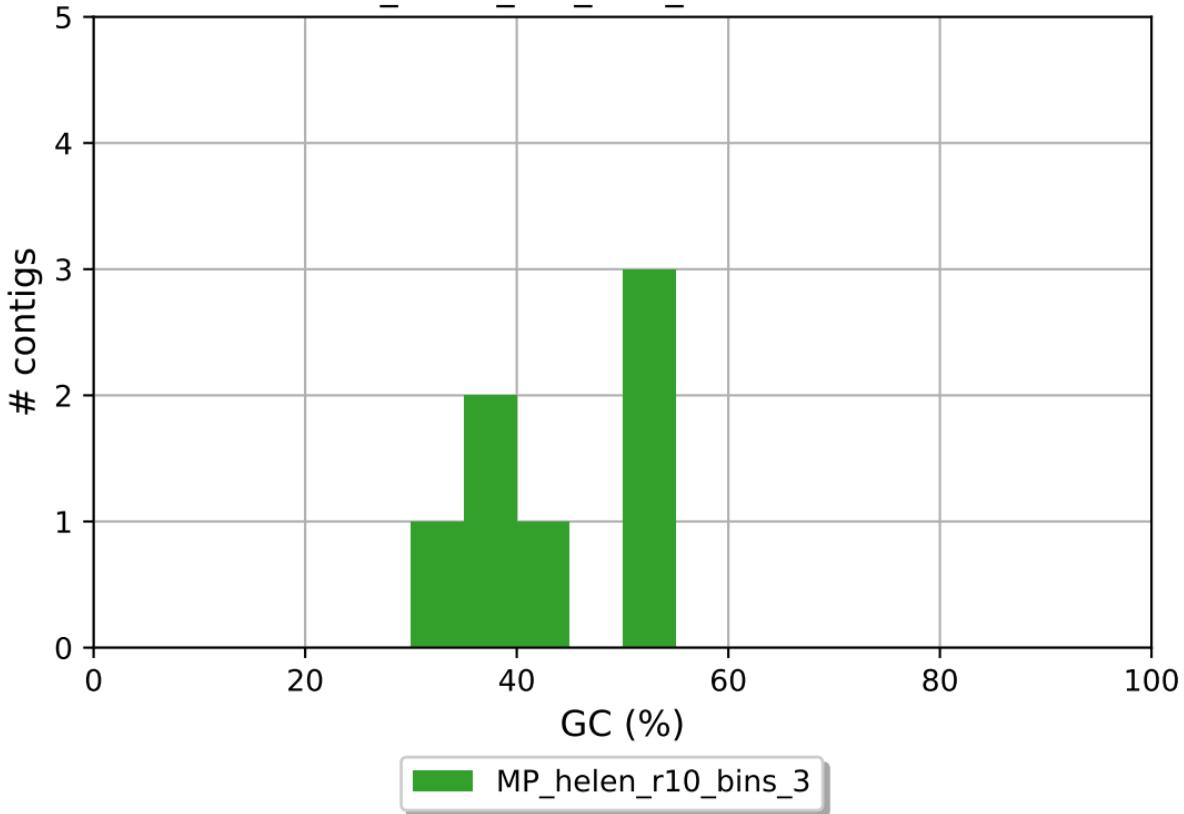
MP_helen_r10_bins_1 GC content



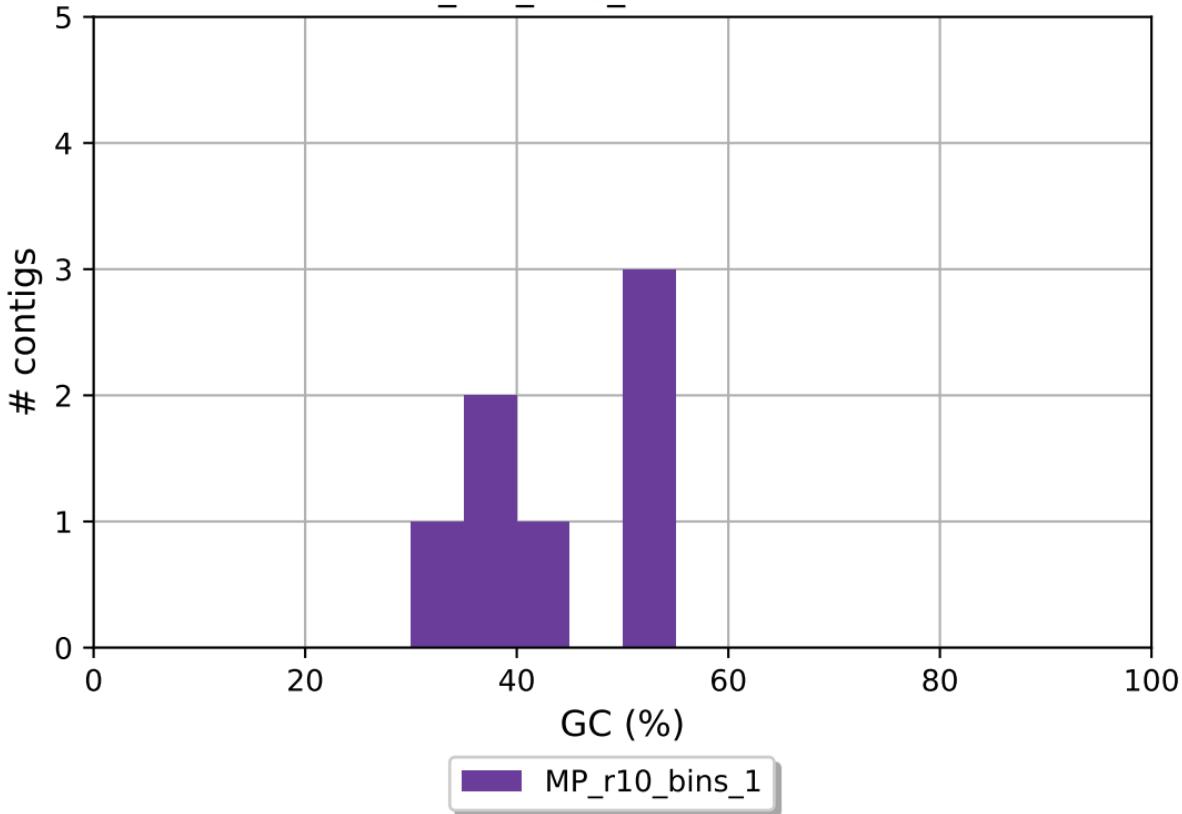
MP_helen_r10_bins_2 GC content



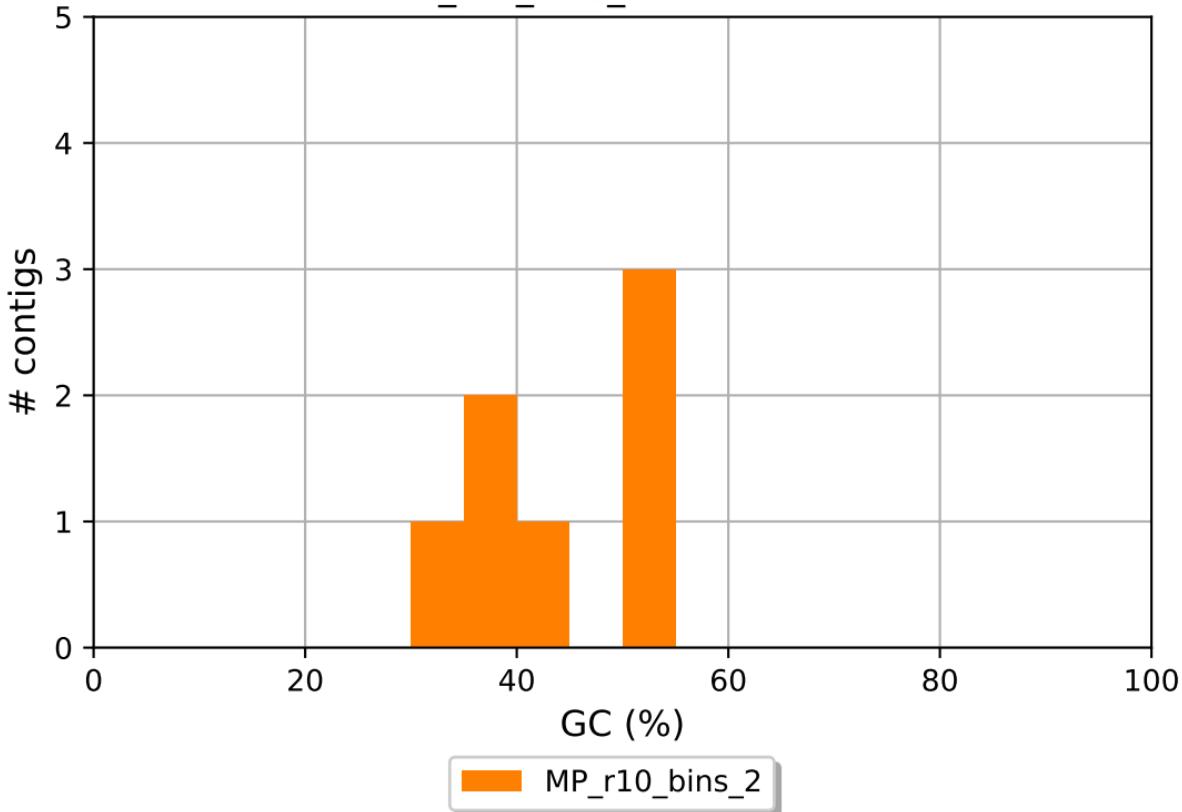
MP_helen_r10_bins_3 GC content



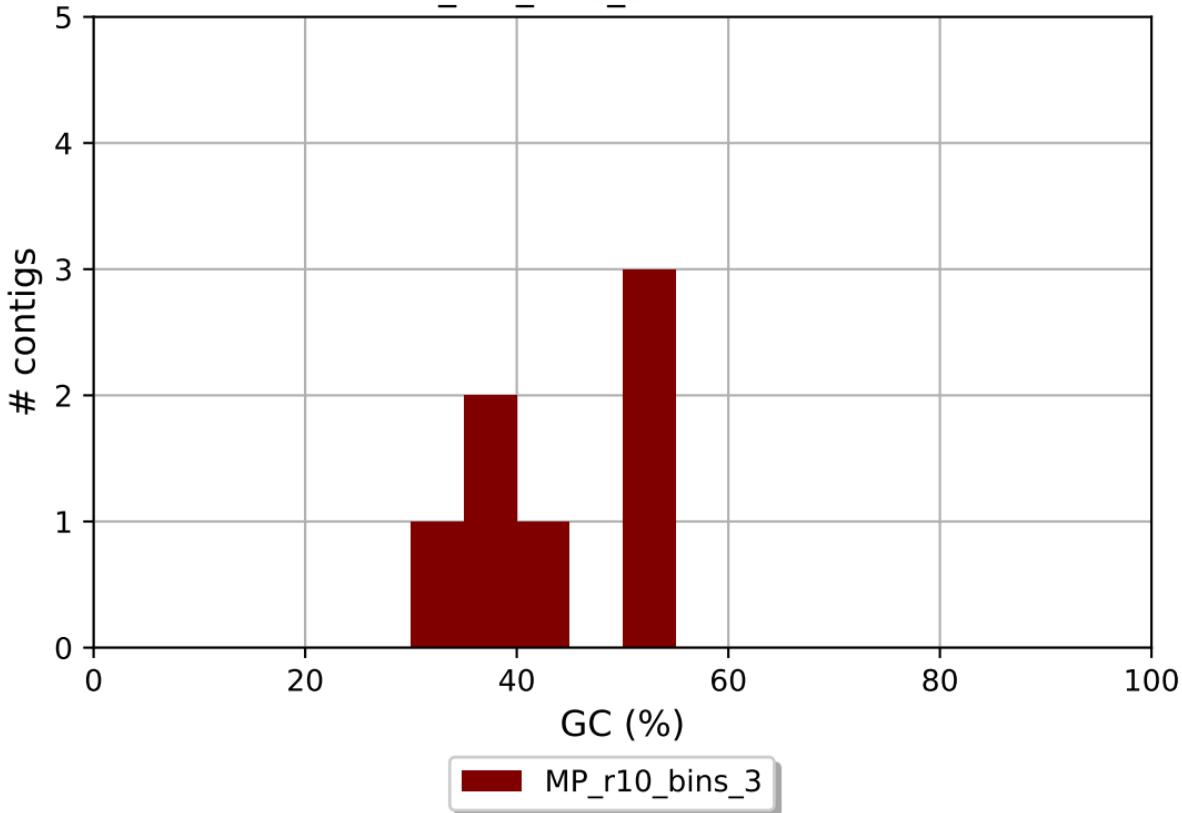
MP_r10_bins_1 GC content



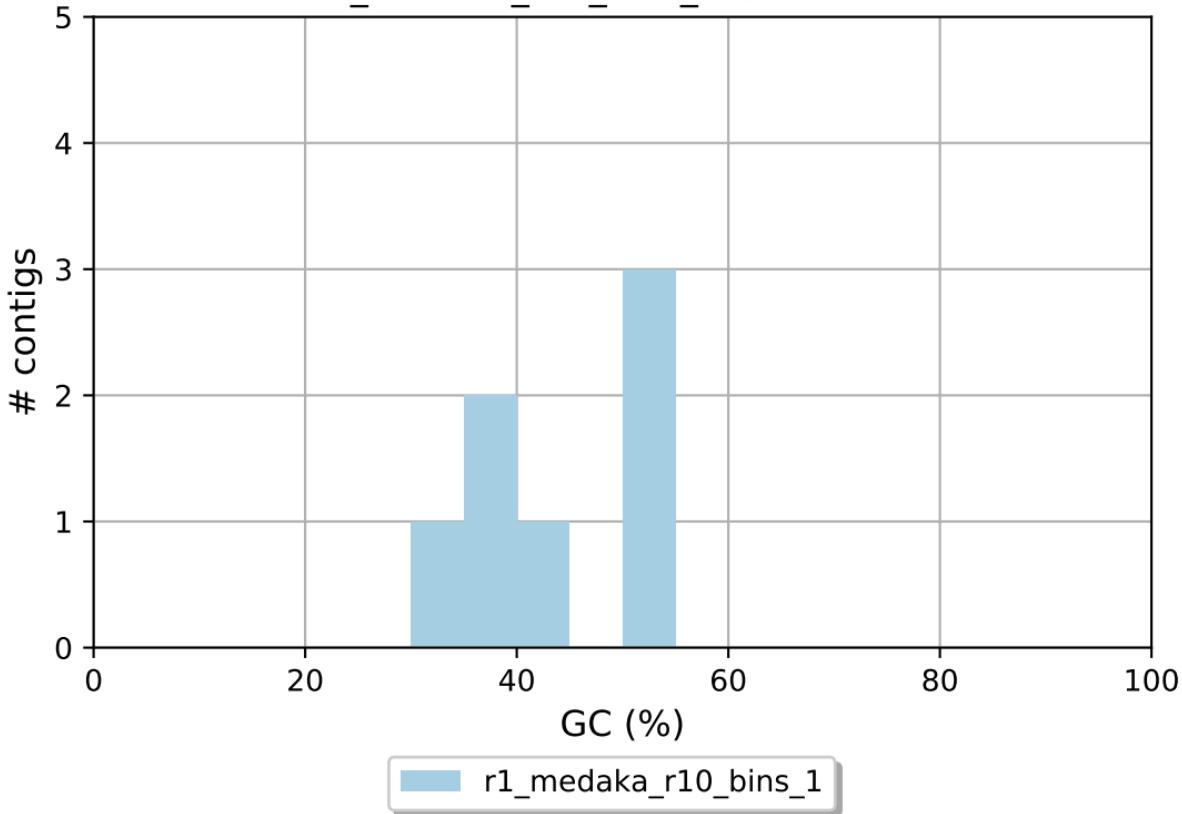
MP_r10_bins_2 GC content



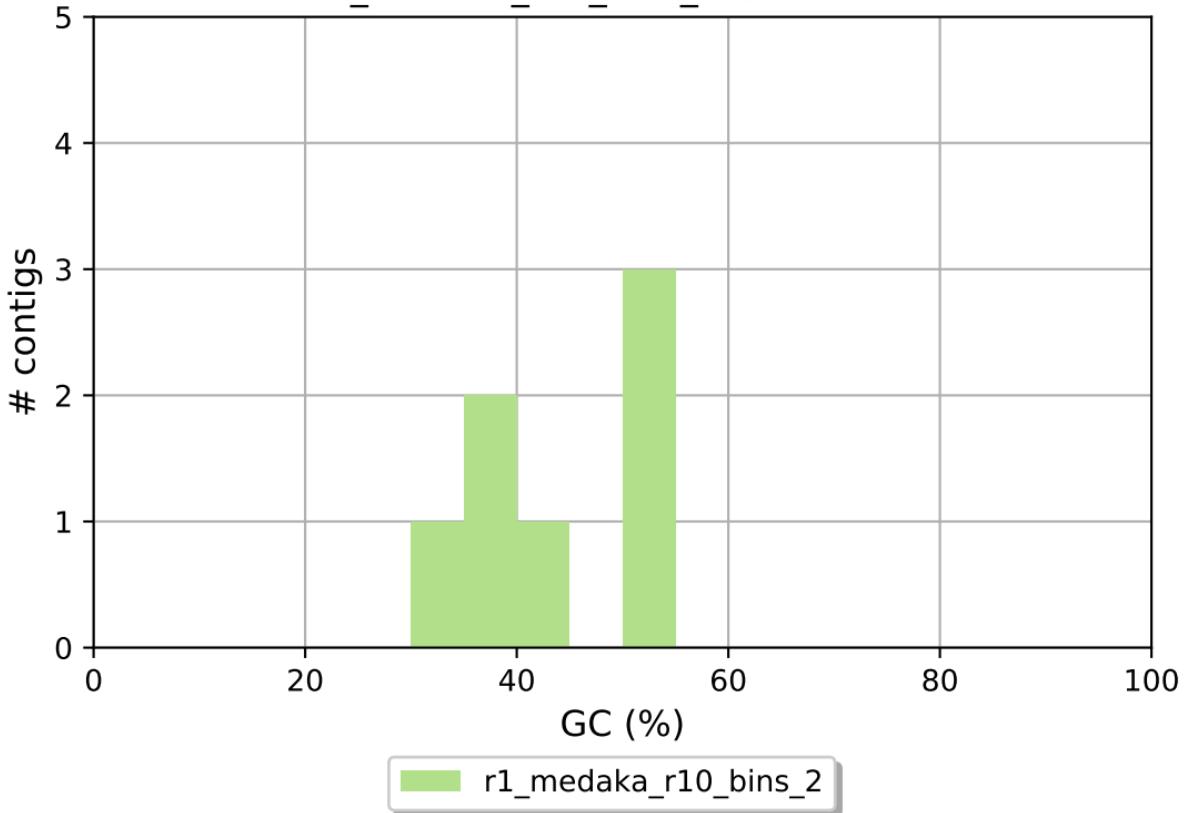
MP_r10_bins_3 GC content



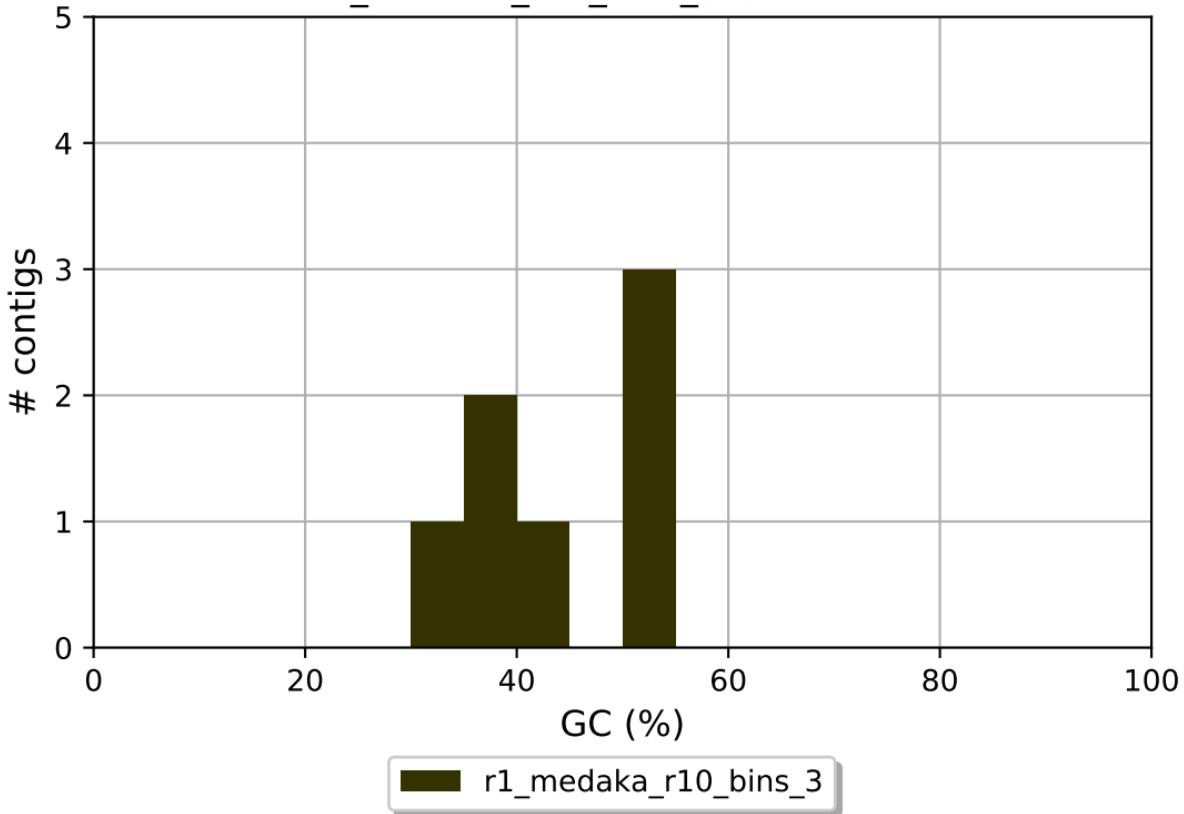
r1_medaka_r10_bins_1 GC content



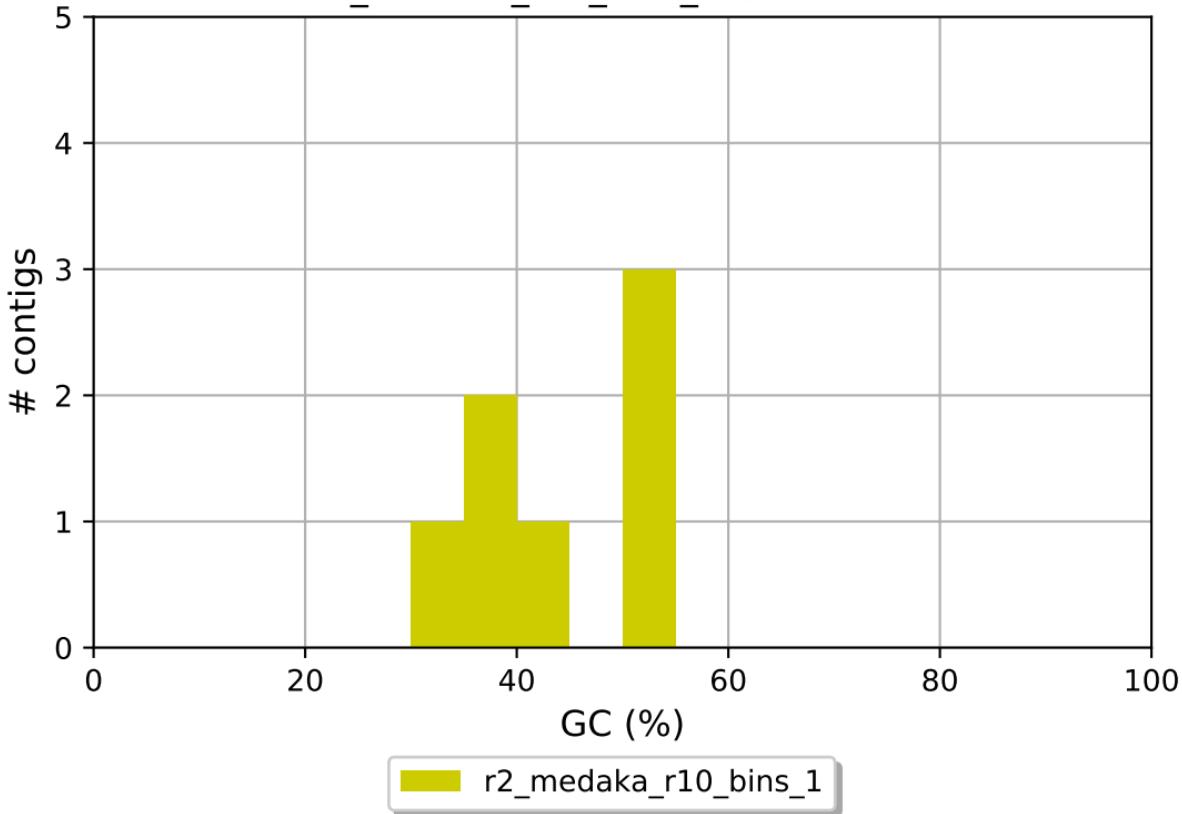
r1_medaka_r10_bins_2 GC content



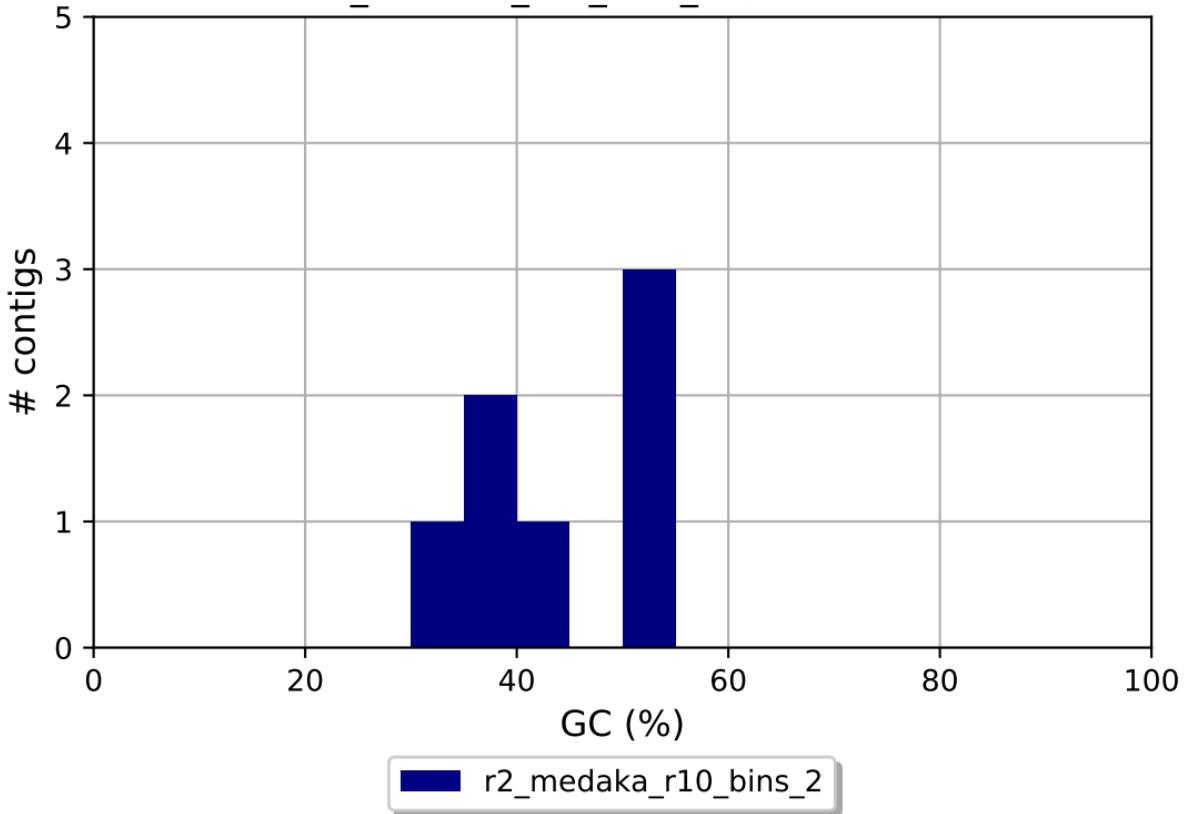
r1_medaka_r10_bins_3 GC content



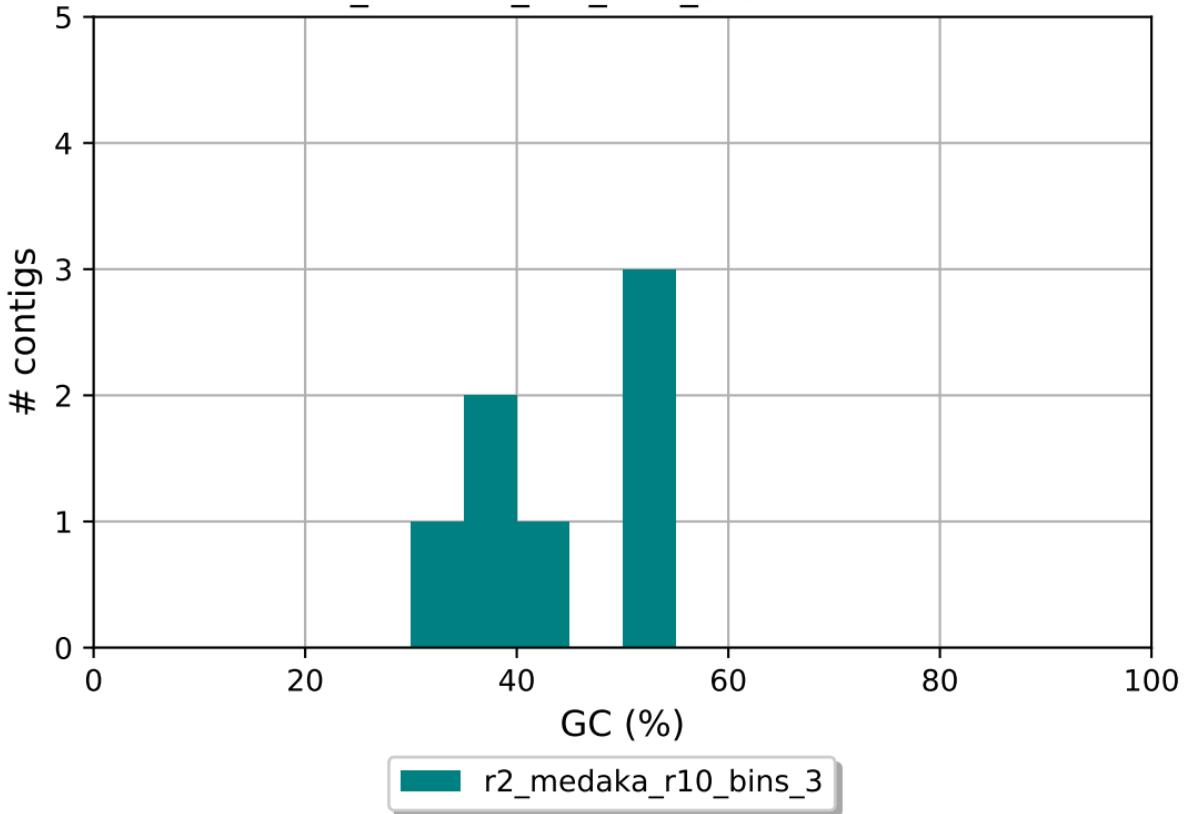
r2_medaka_r10_bins_1 GC content



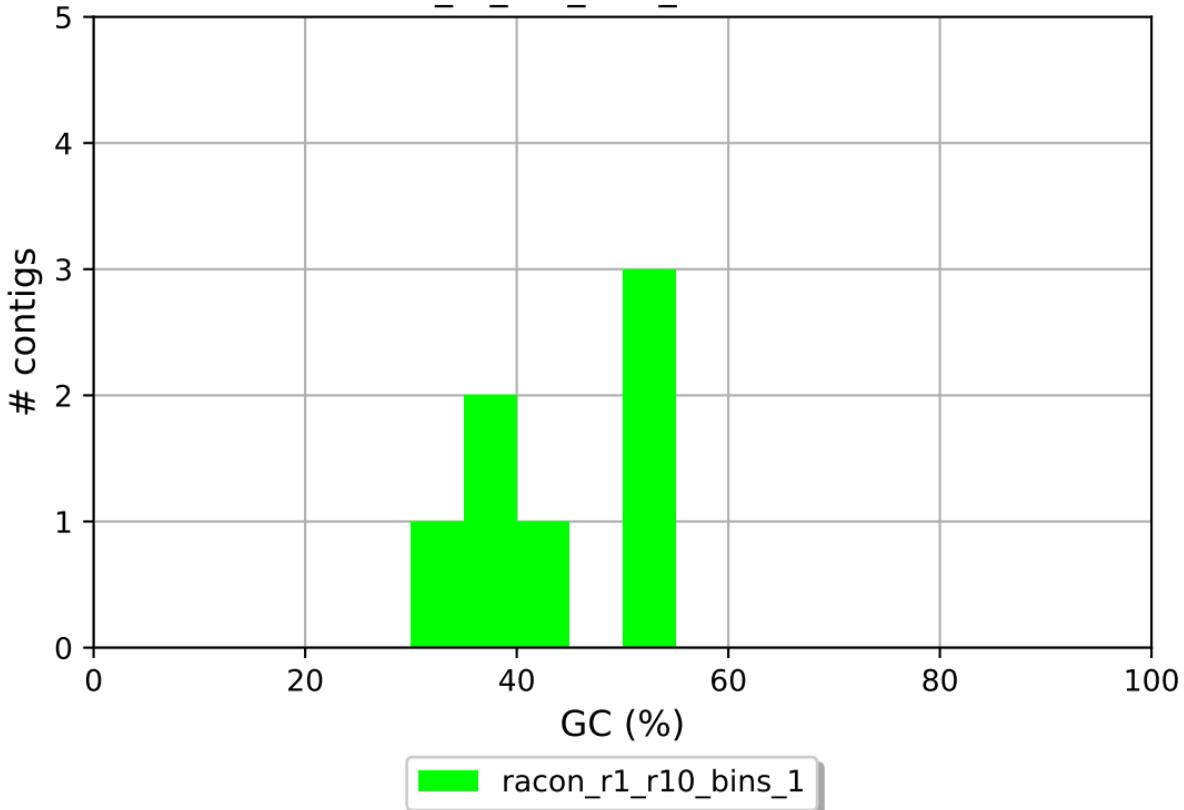
r2_medaka_r10_bins_2 GC content



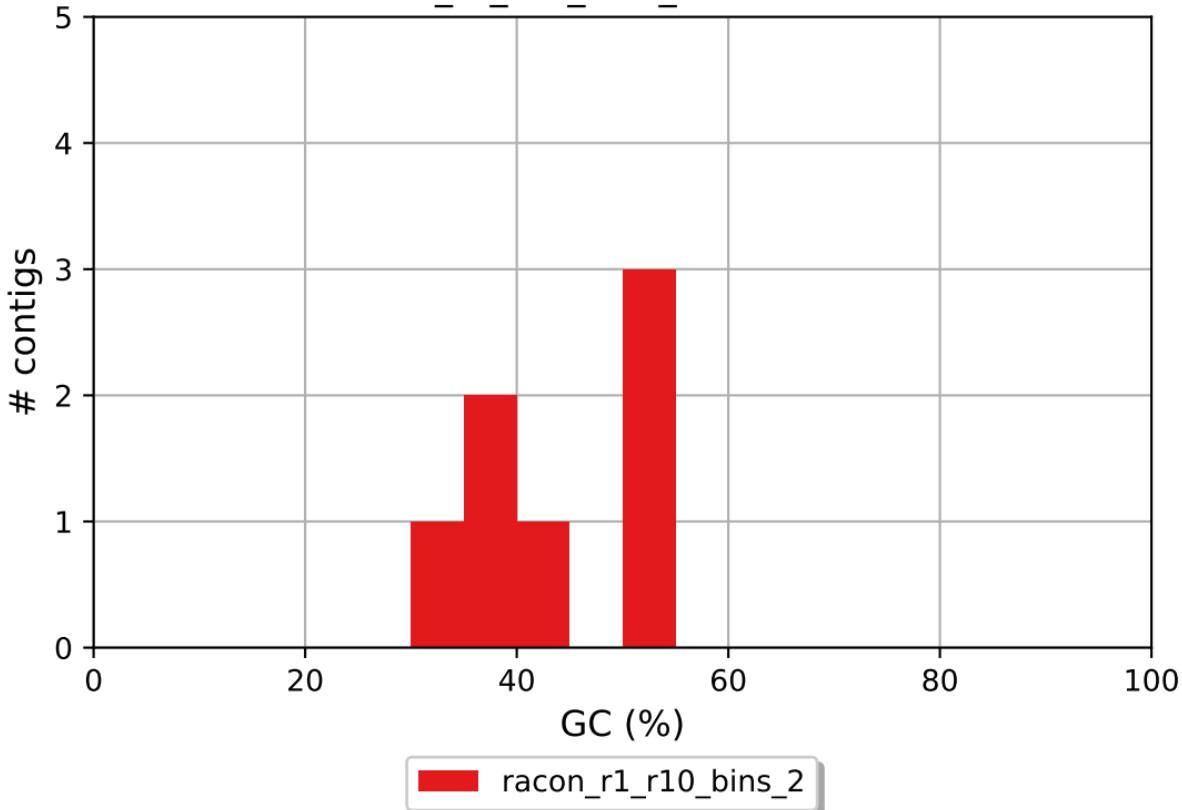
r2_medaka_r10_bins_3 GC content



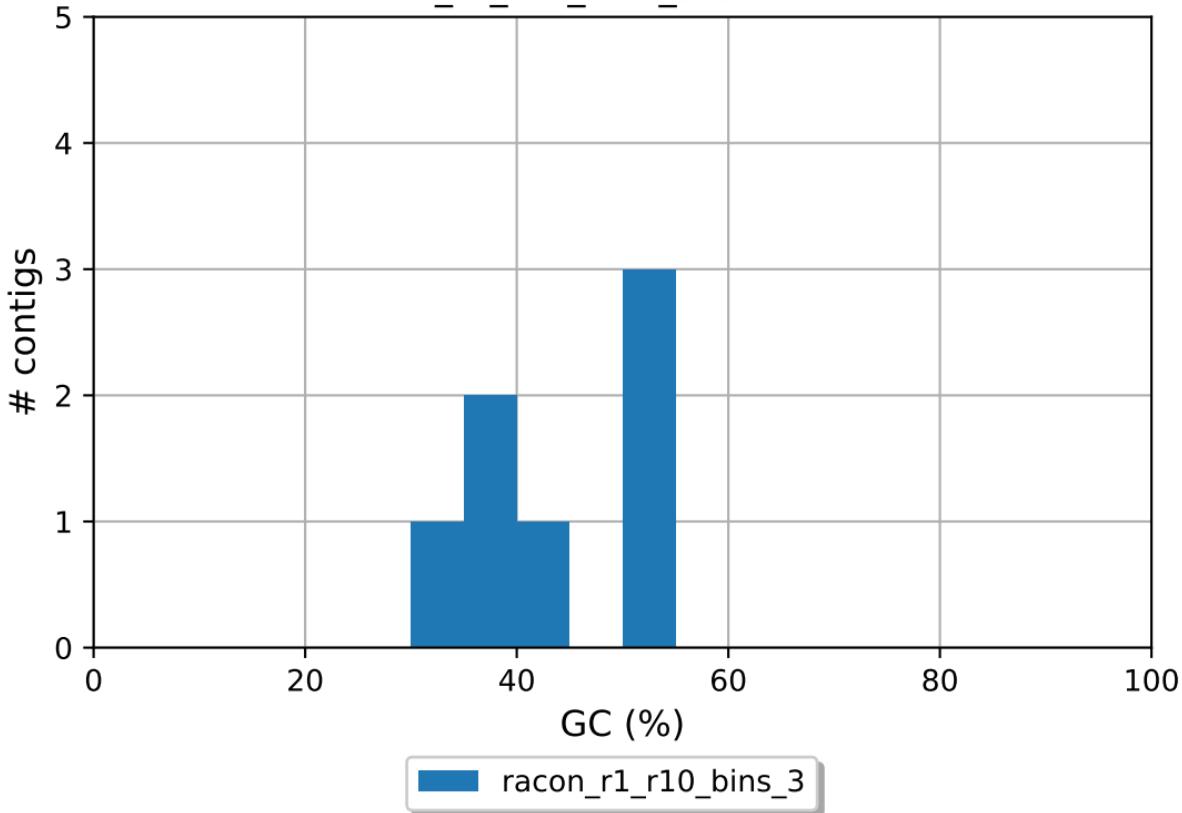
racon_r1_r10_bins_1 GC content



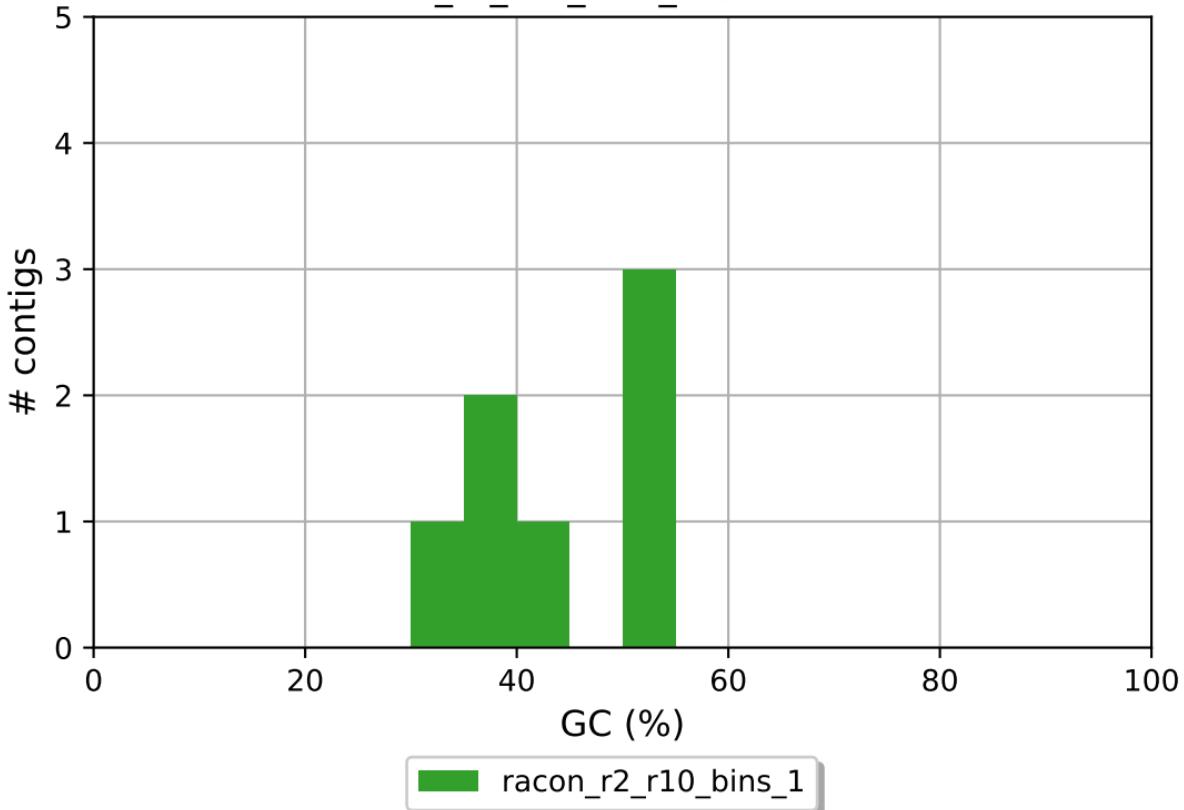
racon_r1_r10_bins_2 GC content



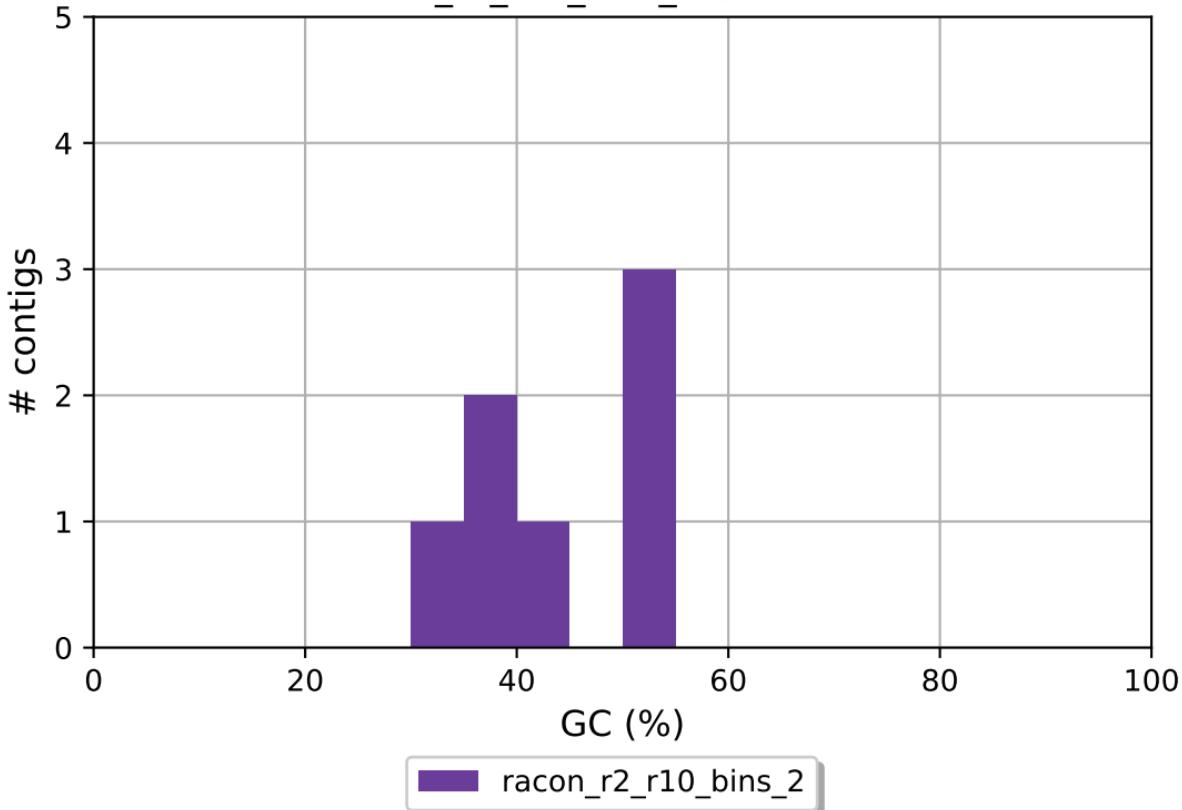
racon_r1_r10_bins_3 GC content



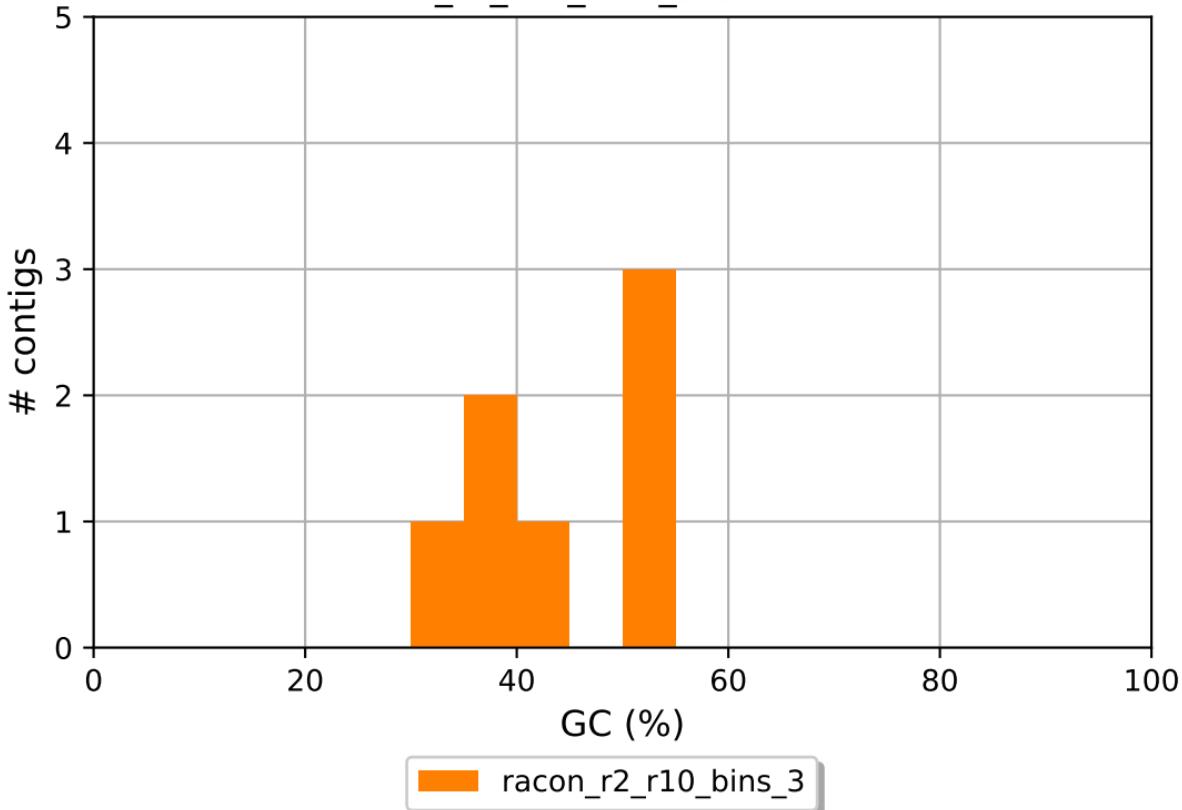
racon_r2_r10_bins_1 GC content



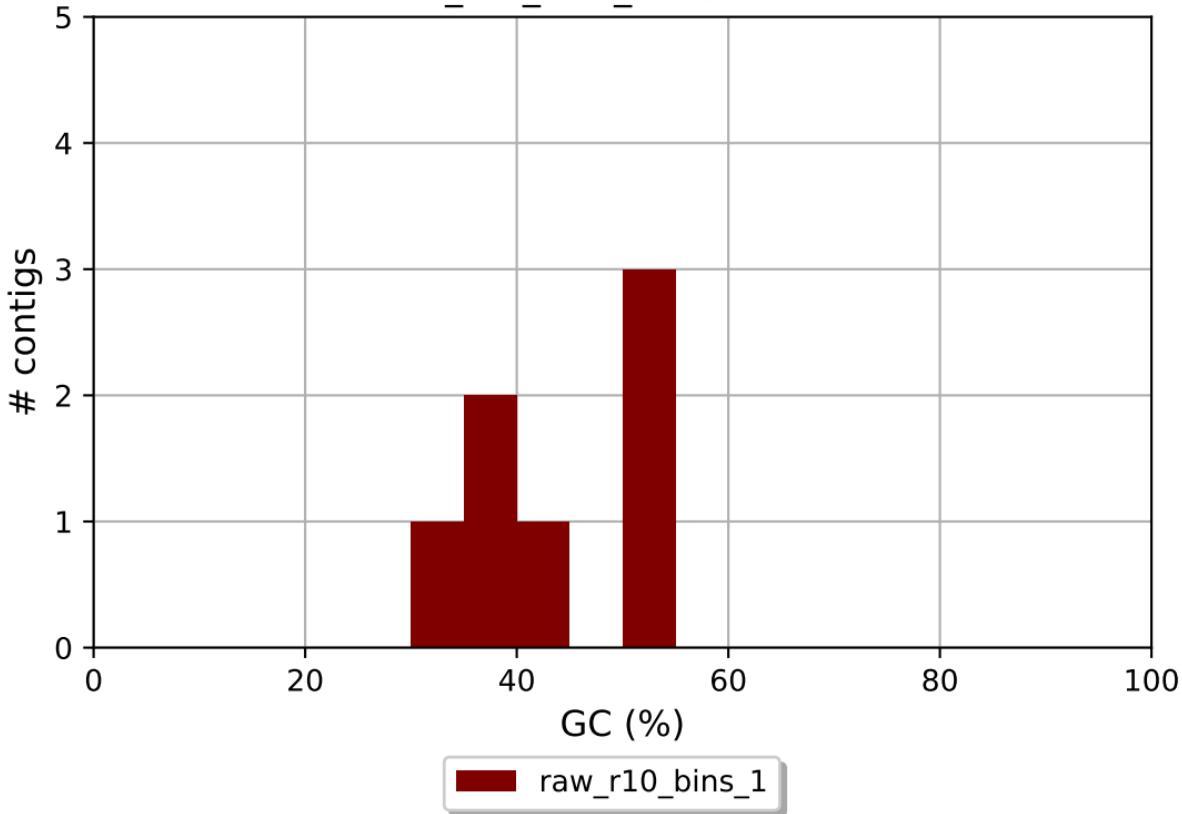
racon_r2_r10_bins_2 GC content



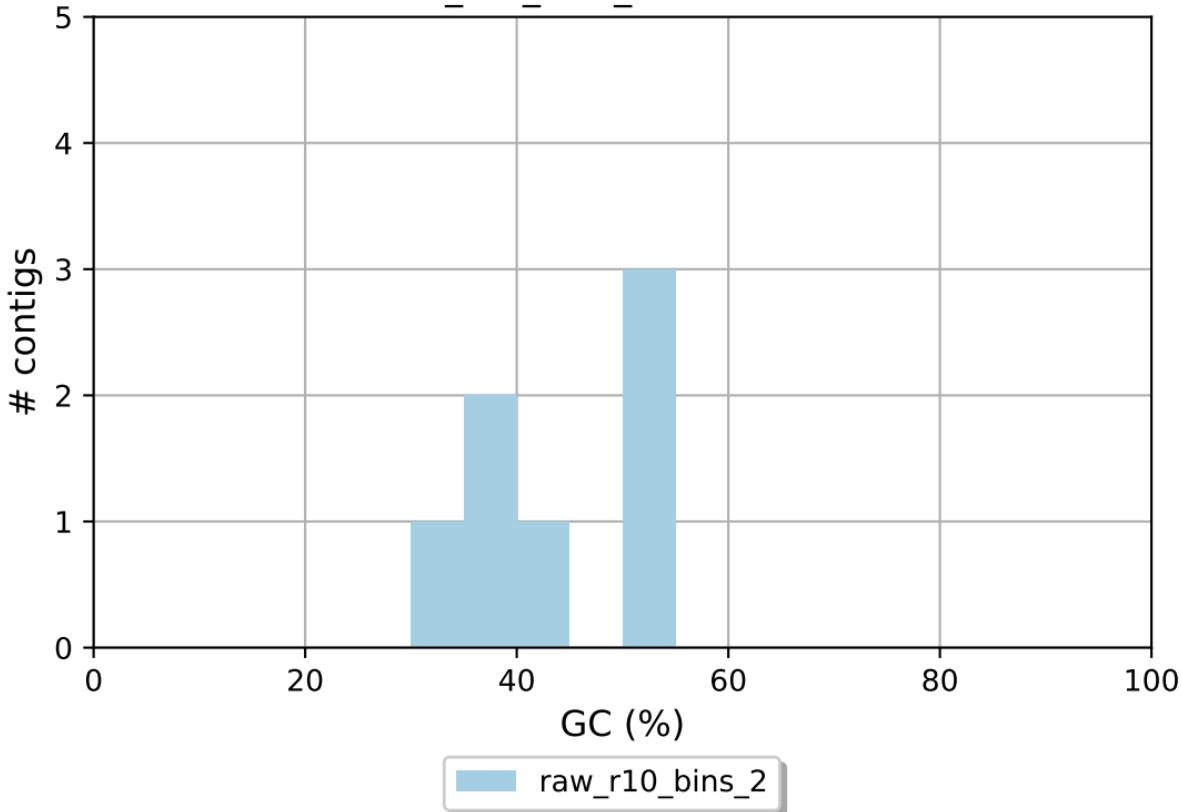
racon_r2_r10_bins_3 GC content



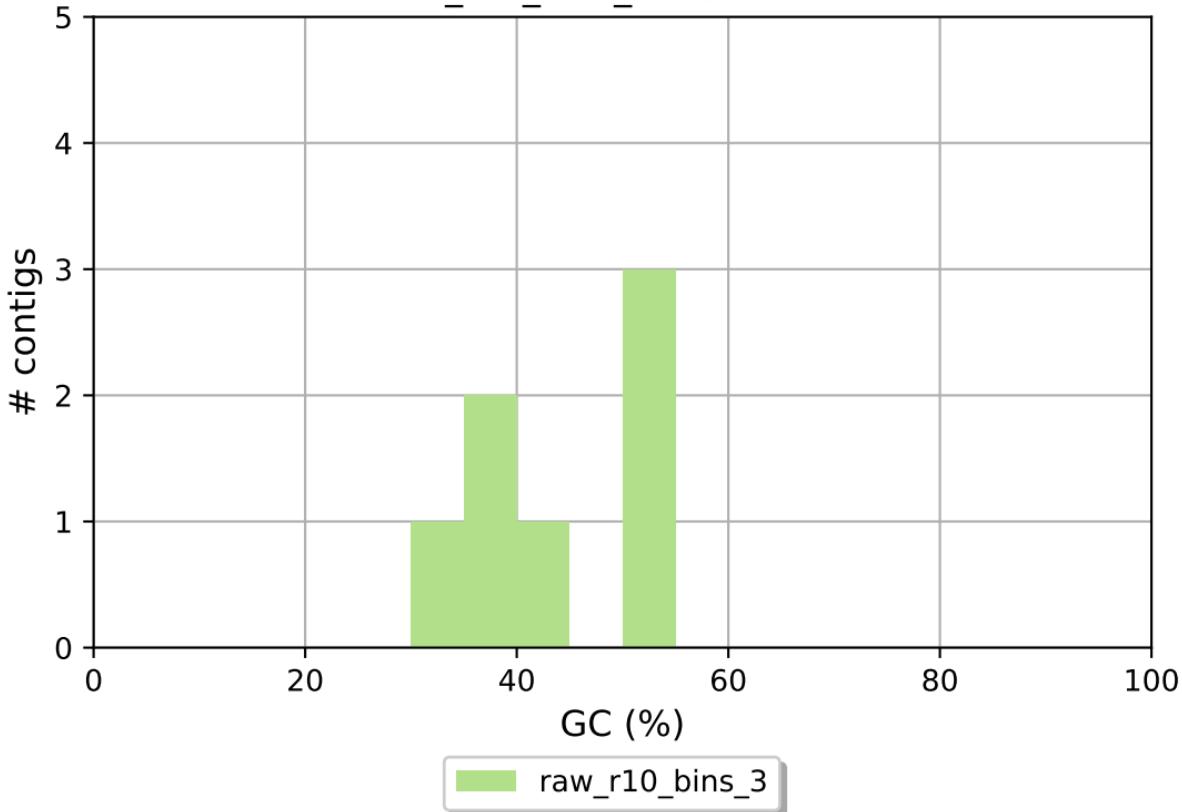
raw_r10_bins_1 GC content



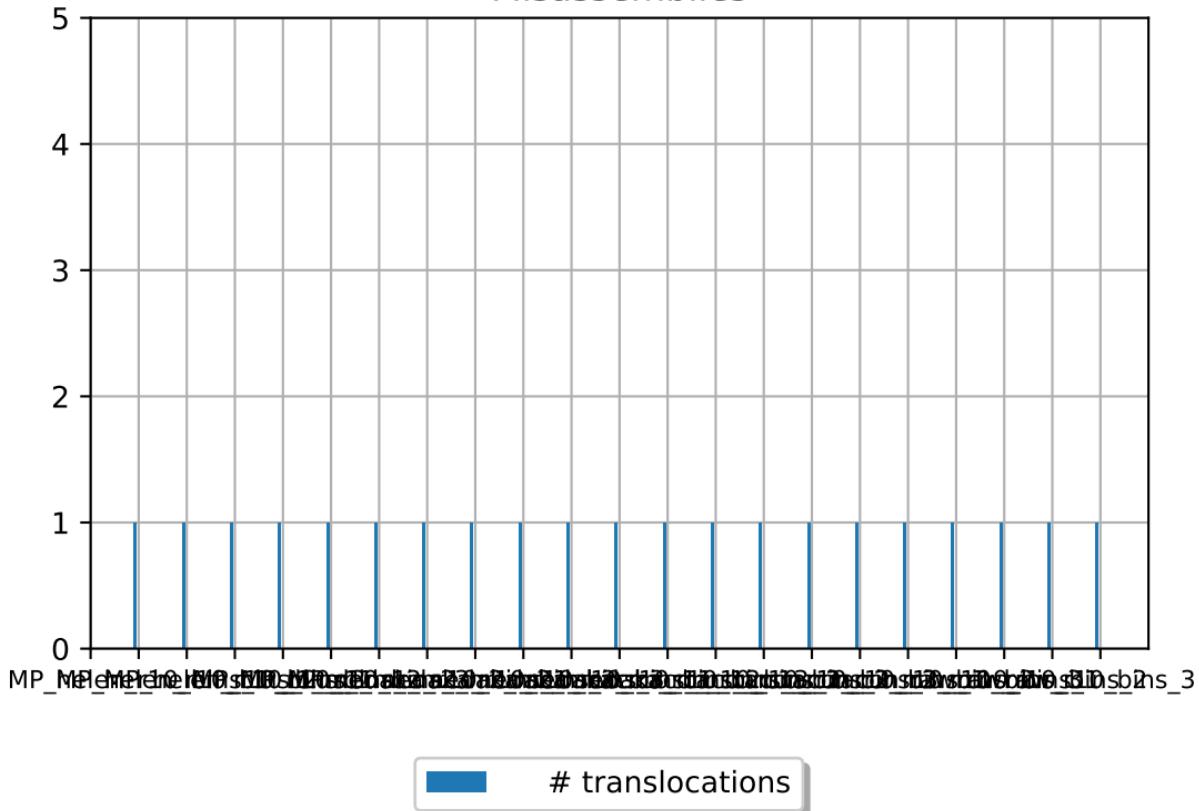
raw_r10_bins_2 GC content



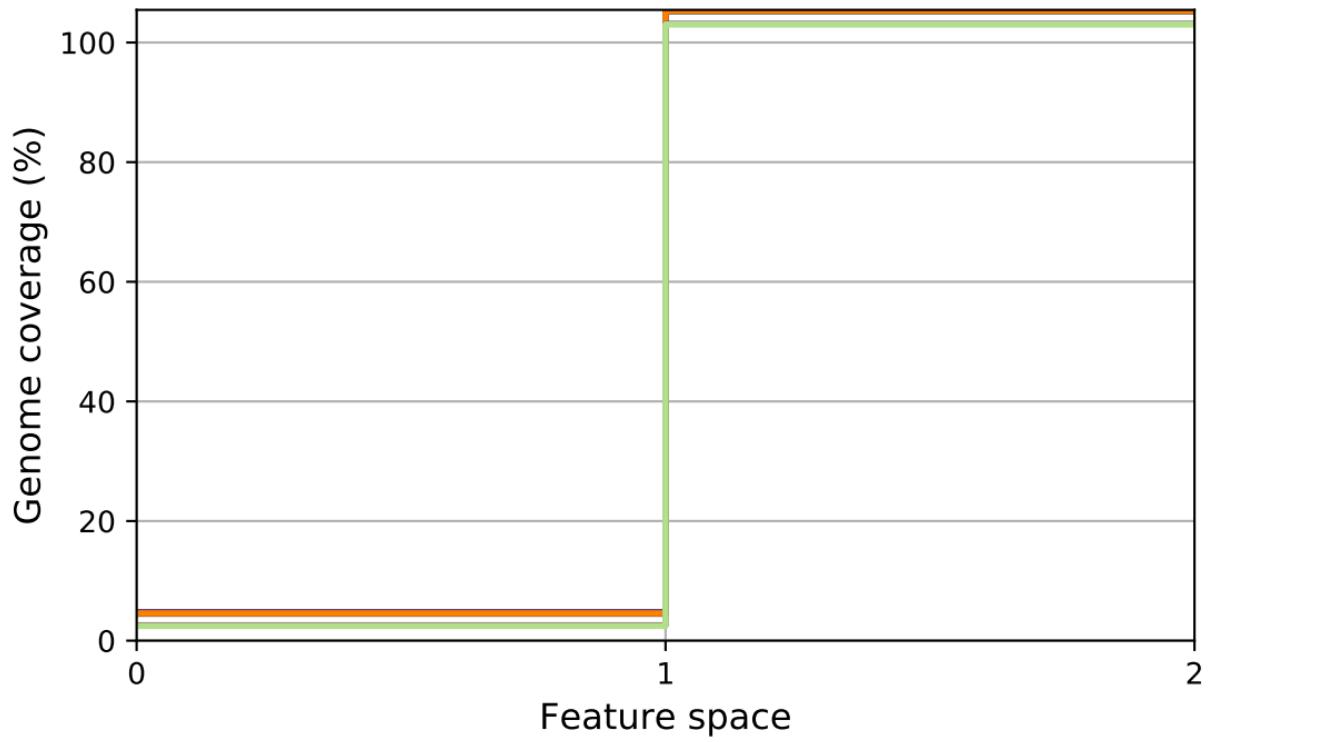
raw_r10_bins_3 GC content



Misassemblies



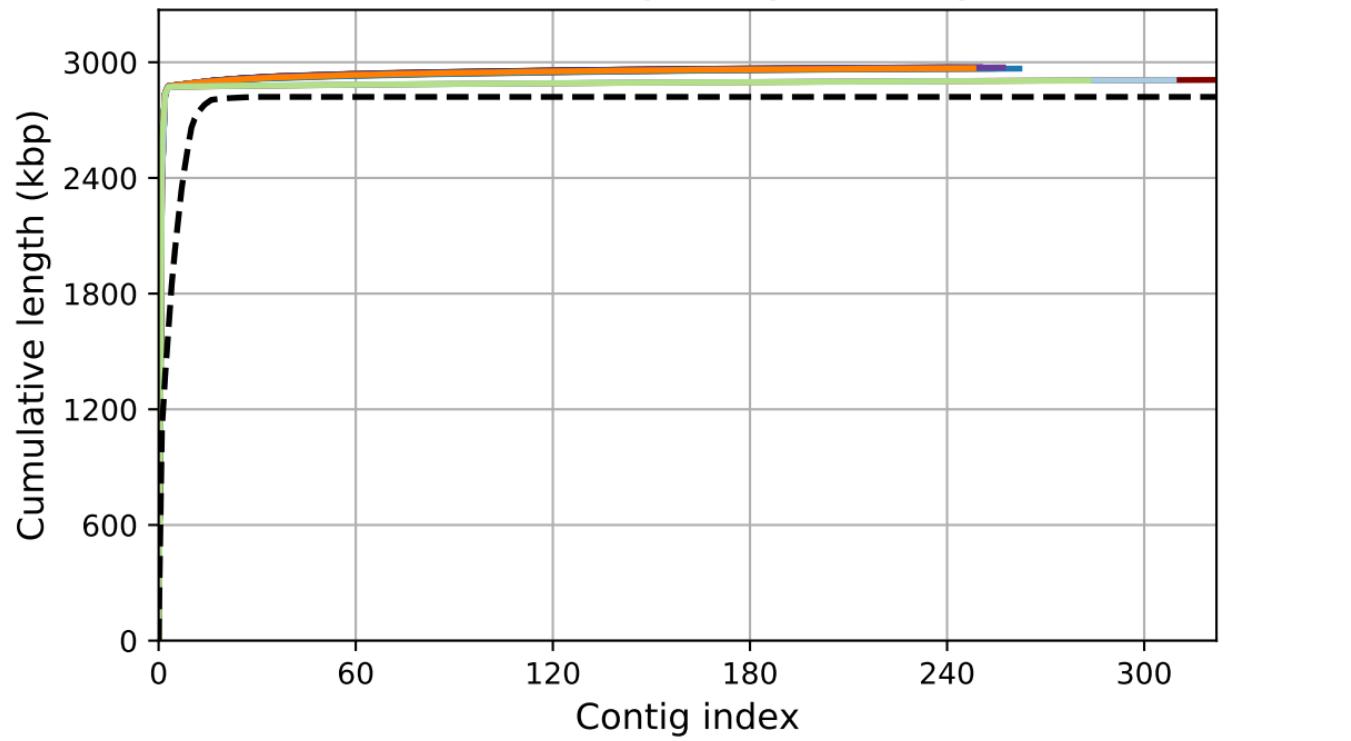
FRCurve (misassemblies)



Legend:

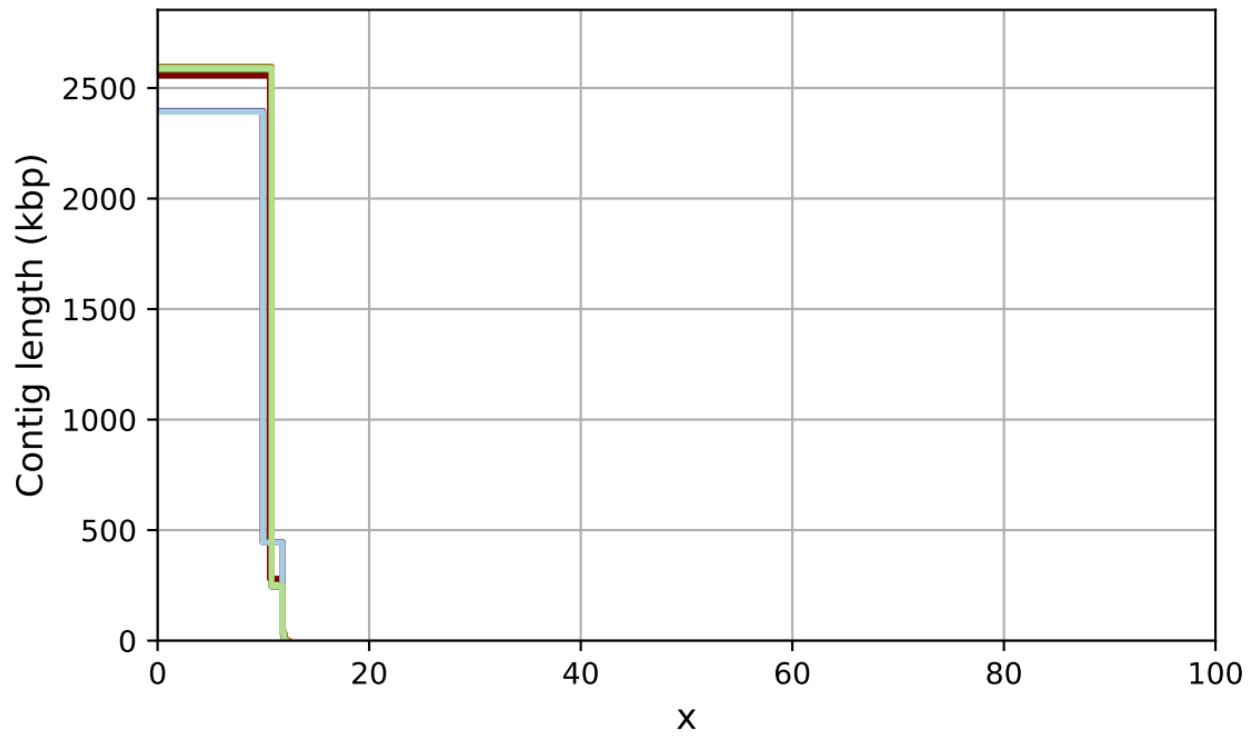
- MP_helen_r10_bins_1
- MP_helen_r10_bins_2
- MP_helen_r10_bins_3
- MP r10 bins 1
- r1_medaka_r10_bins_2
- r1_medaka_r10_bins_3
- r2_medaka_r10_bins_1
- racon_r1_r10_bins_3
- racon_r2_r10_bins_1
- racon_r2_r10_bins_2
- racon_r2_r10_bins_3

Cumulative length (aligned contigs)



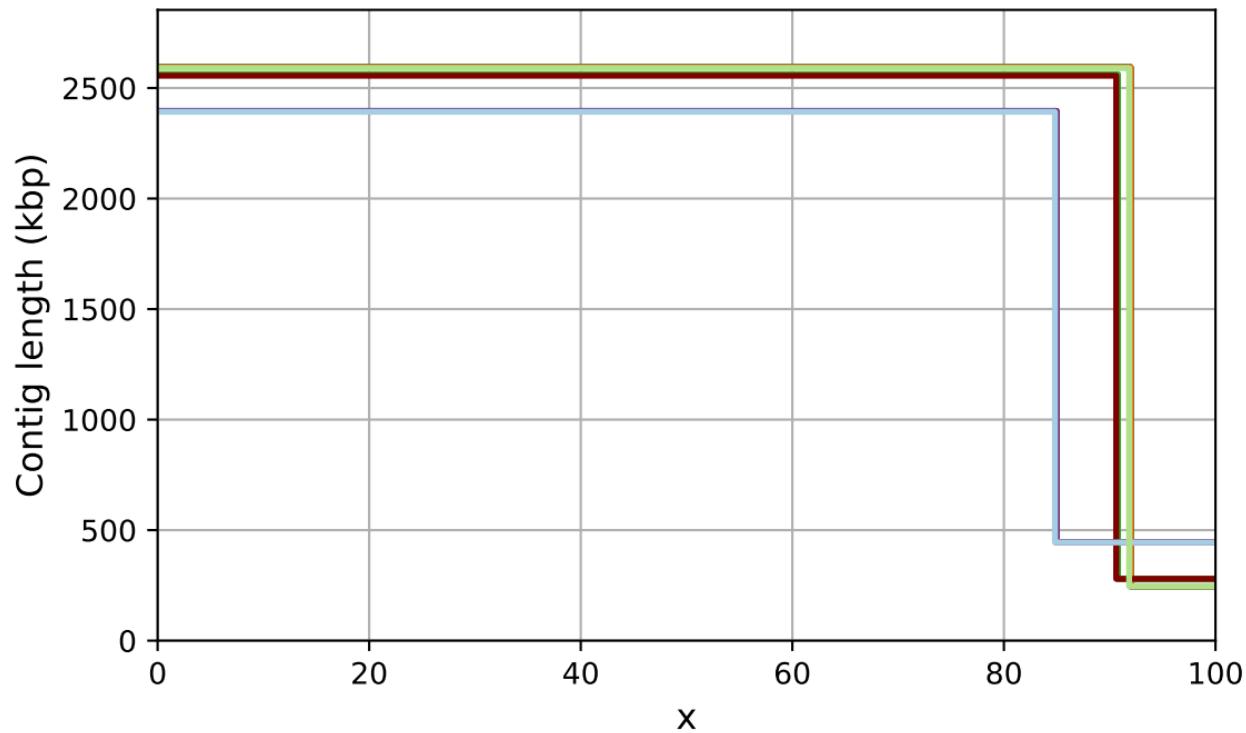
- MP_helen_r10_bins_1
- MP_helen_r10_bins_2
- MP_helen_r10_bins_3
- MP r10 bins 1
- r1_medaka_r10_bins_3
- r2_medaka_r10_bins_1
- r2_medaka_r10_bins_2
- r2_medaka_r10_bins_3
- racon_r2_r10_bins_1
- racon_r2_r10_bins_2
- racon_r2_r10_bins_3
- raw_r10_bins_1

NAx



- MP_helen_r10_bins_1
- MP_helen_r10_bins_2
- MP_helen_r10_bins_3
- MP r10 bins 1
- r1_medaka_r10_bins_2
- r1_medaka_r10_bins_3
- r2_medaka_r10_bins_1
- r2_medaka_r10_bins_2
- racon_r1_r10_bins_3
- racon_r2_r10_bins_1
- racon_r2_r10_bins_2
- racon_r2_r10_bins_3

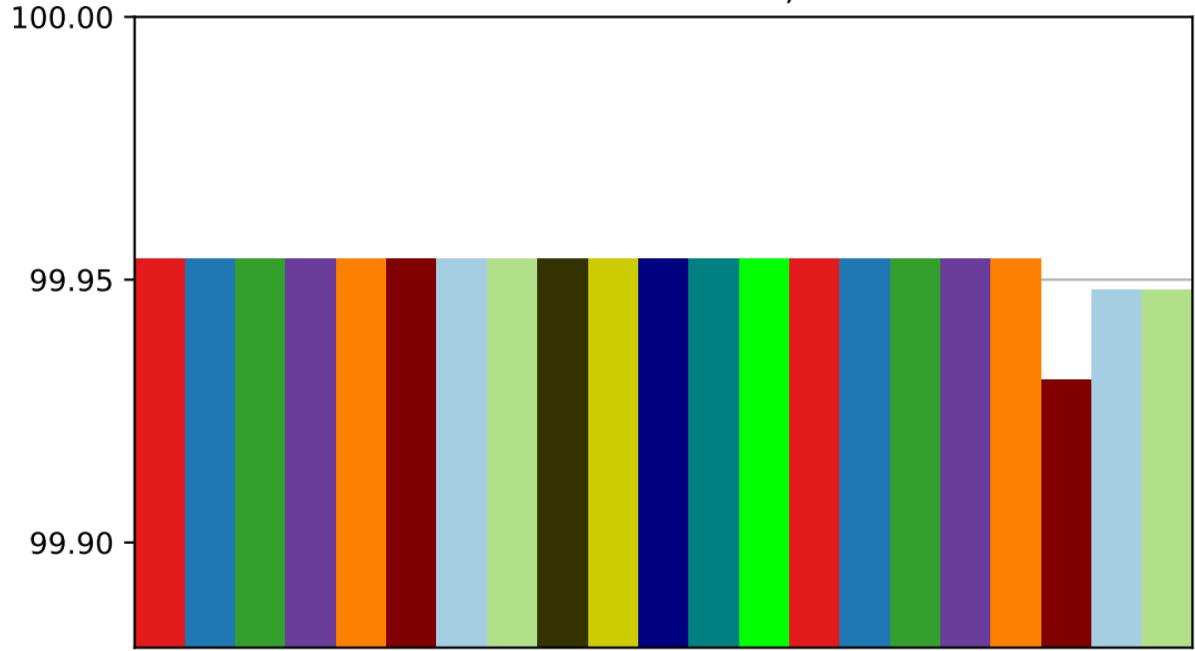
NGAx



Legend for the assembly comparison plot:

- MP_helen_r10_bins_1
- MP_helen_r10_bins_2
- MP_helen_r10_bins_3
- MP_r10_bins_1
- r1_medaka_r10_bins_2
- r1_medaka_r10_bins_3
- r2_medaka_r10_bins_1
- r2_medaka_r10_bins_2
- racon_r1_r10_bins_3
- racon_r2_r10_bins_1
- racon_r2_r10_bins_2
- racon_r2_r10_bins_3

Genome fraction, %



- █ MP_helen_r10_bins_1
- █ MP_helen_r10_bins_2
- █ MP_helen_r10_bins_3
- █ MP_r10_bins_1
- █ r1_medaka_r10_bins_2
- █ r1_medaka_r10_bins_3
- █ r2_medaka_r10_bins_1
- █ r2_medaka_r10_bins_2
- █ racon_r1_r10_bins_3
- █ racon_r2_r10_bins_1
- █ racon_r2_r10_bins_2
- █ racon_r2_r10_bins_3