

MP10k

Summary

General

fastp version:	0.19.4 (https://github.com/OpenGene/fastp)
sequencing:	paired end (151 cycles + 151 cycles)
mean length before filtering:	151bp, 151bp
mean length after filtering:	149bp, 149bp
duplication rate:	70.067730%
Insert size peak:	0

Before filtering

total reads:	446.717352 M
total bases:	67.454320 G
Q20 bases:	62.581933 G (92.776761%)
Q30 bases:	57.303928 G (84.952198%)
GC content:	44.969723%

After filtering

total reads:	334.744784 M
total bases:	50.109674 G
Q20 bases:	48.660270 G (97.107537%)
Q30 bases:	45.704291 G (91.208519%)
GC content:	42.618955%

Filtering result

reads passed filters:	334.744784 M (74.934359%)
reads with low quality:	103.226866 M (23.107870%)
reads with too many N:	2.802000 K (0.000627%)
reads too short:	8.725694 M (1.953292%)
reads with low complexity:	17.206000 K (0.003852%)

Adapters

Adapter or bad ligation of read1

The input has little adapter percentage (~0.372463%), probably it's trimmed before.

Sequence	Occurrences
A	68599
AG	60334
AGA	68812
AGAT	59534
AGATC	

	65833
AGATCG	65081
AGATCGG	56868
AGATCGGA	57555
AGATCGGAA	57759
AGATCGGAAG	54905
AGATCGGAAGA	56033
AGATCGGAAGAG	54606
AGATCGGAAGAGC	51860
AGATCGGAAGAGCA	57361
AGATCGGAAGAGCAC	61168
AGATCGGAAGAGCACA	52506
AGATCGGAAGAGCACAC	51798
AGATCGGAAGAGCACACG	49885
AGATCGGAAGAGCACACGT	44528
AGATCGGAAGAGCACACGTC	46722
AGATCGGAAGAGCACACGTCT	44768
AGATCGGAAGAGCACACGTCTG	43424
AGATCGGAAGAGCACACGTCTGA	43889
AGATCGGAAGAGCACACGTCTGAA	43657
AGATCGGAAGAGCACACGTCTGAAC	43251
AGATCGGAAGAGCACACGTCTGAACT	41374
AGATCGGAAGAGCACACGTCTGAACTC	45168
AGATCGGAAGAGCACACGTCTGAACTCC	39336
AGATCGGAAGAGCACACGTCTGAACTCCA	40941
AGATCGGAAGAGCACACGTCTGAACTCCAG	38276
AGATCGGAAGAGCACACGTCTGAACTCCAGTC	42963
AGATCGGAAGAGCACACGTCTGAACTCCAGTCA	42092
AGATCGGAAGAGCACACGTCTGAACTCCAGTCAC	37147
AGATCGGAAGAGCACACGTCTGAACTCCAGTCACCTTGTA	49551
AGATCGGAAGAGCACACGTCTGAACTCCAGTCACCTTGTAATC	43605
AGATCGGAAGAGCACACGTCTGAACTCCAGTCACCTTGTAATCTCGTATGCCGTCTTCTGCTT	48912
AGATCGGAAGAGCACACGTCTGAACTCCAGTCACCTTGTAATCTCGTATGCCGTCTTCTGCTTGAAAA	46105
AGATCGGAAGAGCACACGTCTGAACTCCAGTCACCTTGTAATCTCGTATGCCGTCTTCTGCTTGAAAA	46532
other adapter sequences	1759669

Adapter or bad ligation of read2

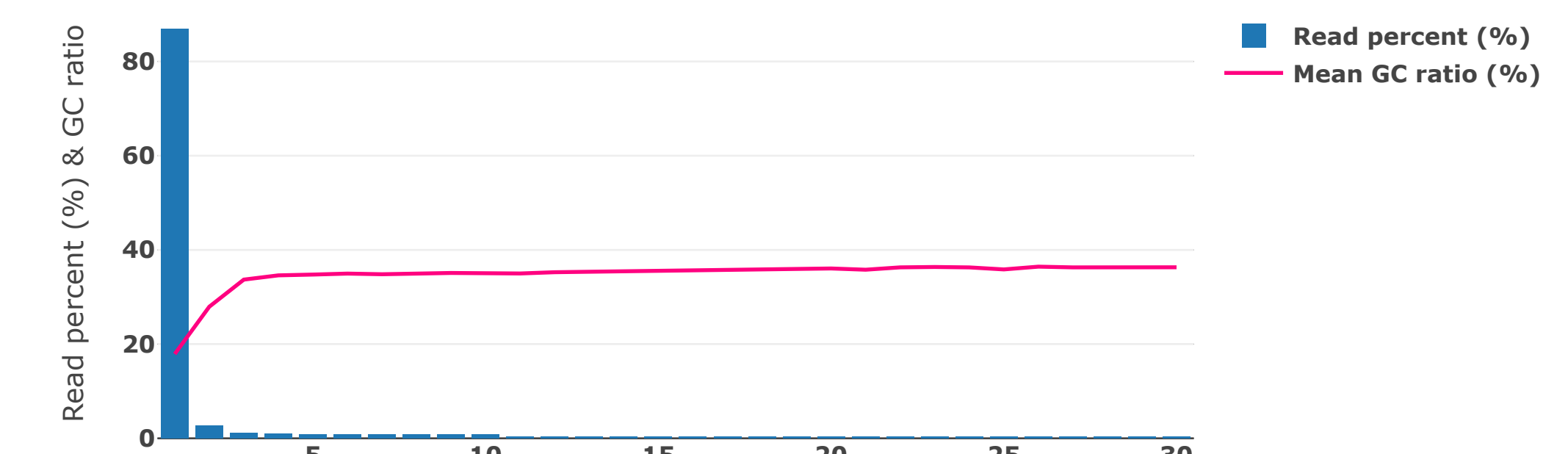
The input has little adapter percentage (~0.327220%), probably it's trimmed before.

Sequence	Occurrences
A	68470
AG	59629
AGA	66817

AGAT	62554
AGATC	70824
AGATCG	66301
AGATCGG	61251
AGATCGGA	58401
AGATCGGAA	56142
AGATCGGAAG	56859
AGATCGGAAGA	61624
AGATCGGAAGAG	57525
AGATCGGAAGAGC	56216
AGATCGGAAGAGCG	63747
AGATCGGAAGAGCGT	48893
AGATCGGAAGAGCGTC	65289
AGATCGGAAGAGCGTCG	57510
AGATCGGAAGAGCGTCGT	50858
AGATCGGAAGAGCGTCGTG	51462
AGATCGGAAGAGCGTCGTGT	45625
AGATCGGAAGAGCGTCGTGTA	57181
AGATCGGAAGAGCGTCGTGTAG	141021
AGATCGGAAGAGCGTCGTGTAGG	118913
AGATCGGAAGAGCGTCGTGTAGGG	50658
AGATCGGAAGAGCGTCGTGTAGGGA	72040
AGATCGGAAGAGCGTCGTGTAGGGAAA	81911
AGATCGGAAGAGCGTCGTGTAGGGAAAGA	45206
AGATCGGAAGAGCGTCGTGTAGGGAAAGAG	79110
AGATCGGAAGAGCGTCGTGTAGGGAAAGAGT	47063
other adapter sequences	1894156

Duplication

duplication rate (70.067730%)

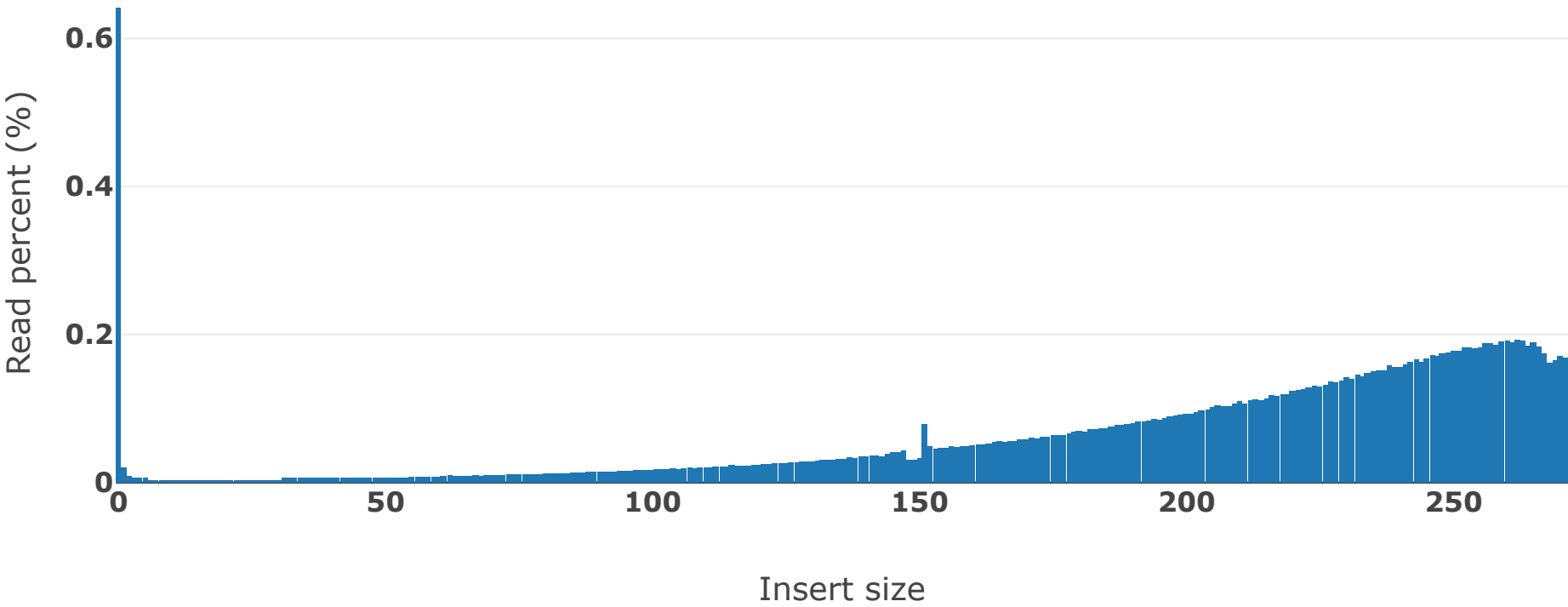


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duplication level

Insert size estimation

Insert size distribution (83.467602% reads are with unknown length)

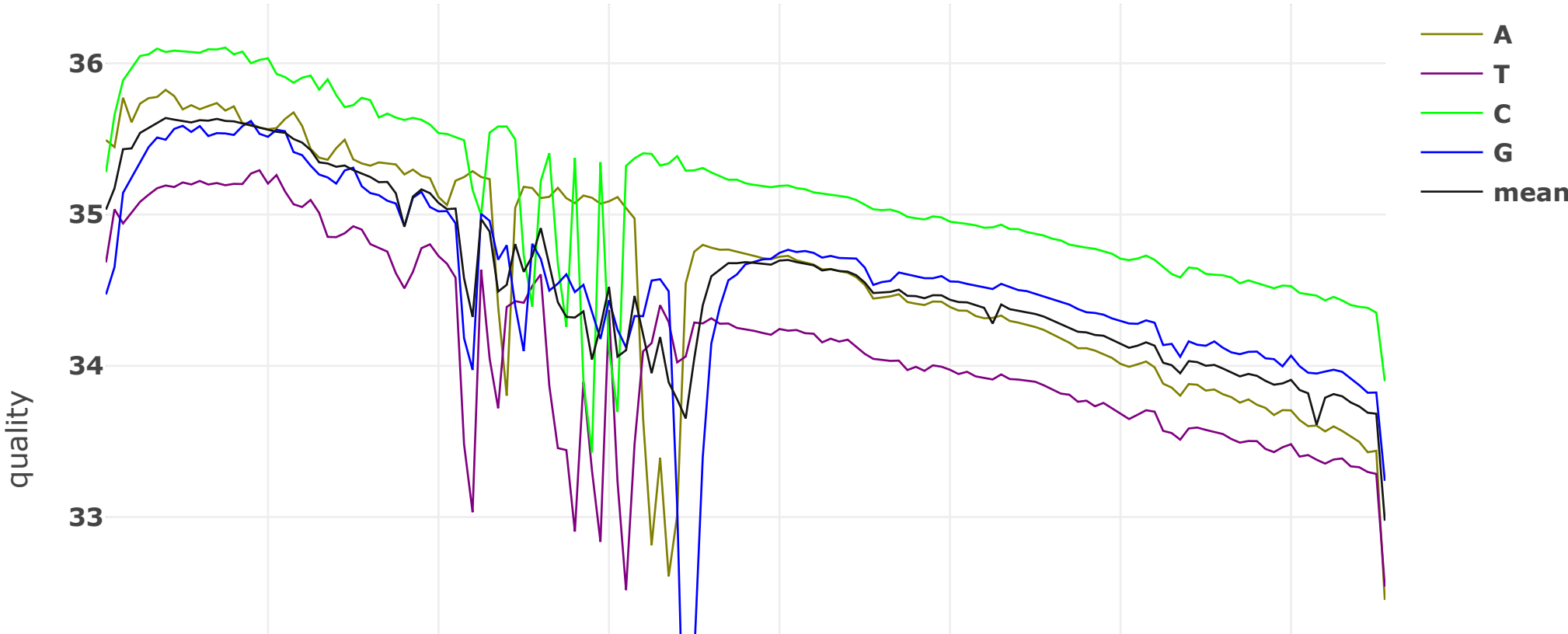


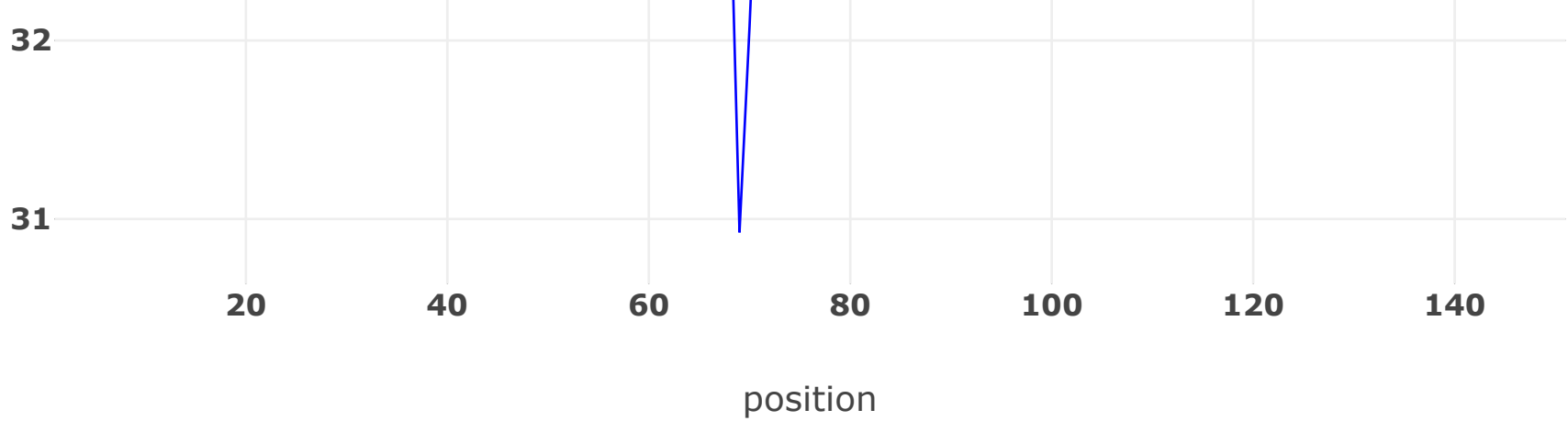
This estimation is based on paired-end overlap analysis, and there are 83.467602% reads found not overlapped. The nonoverlapped read pairs may have insert size <30 or >272, or contain too much sequencing errors to be detected as overlapped.

Before filtering

Before filtering: read1: quality

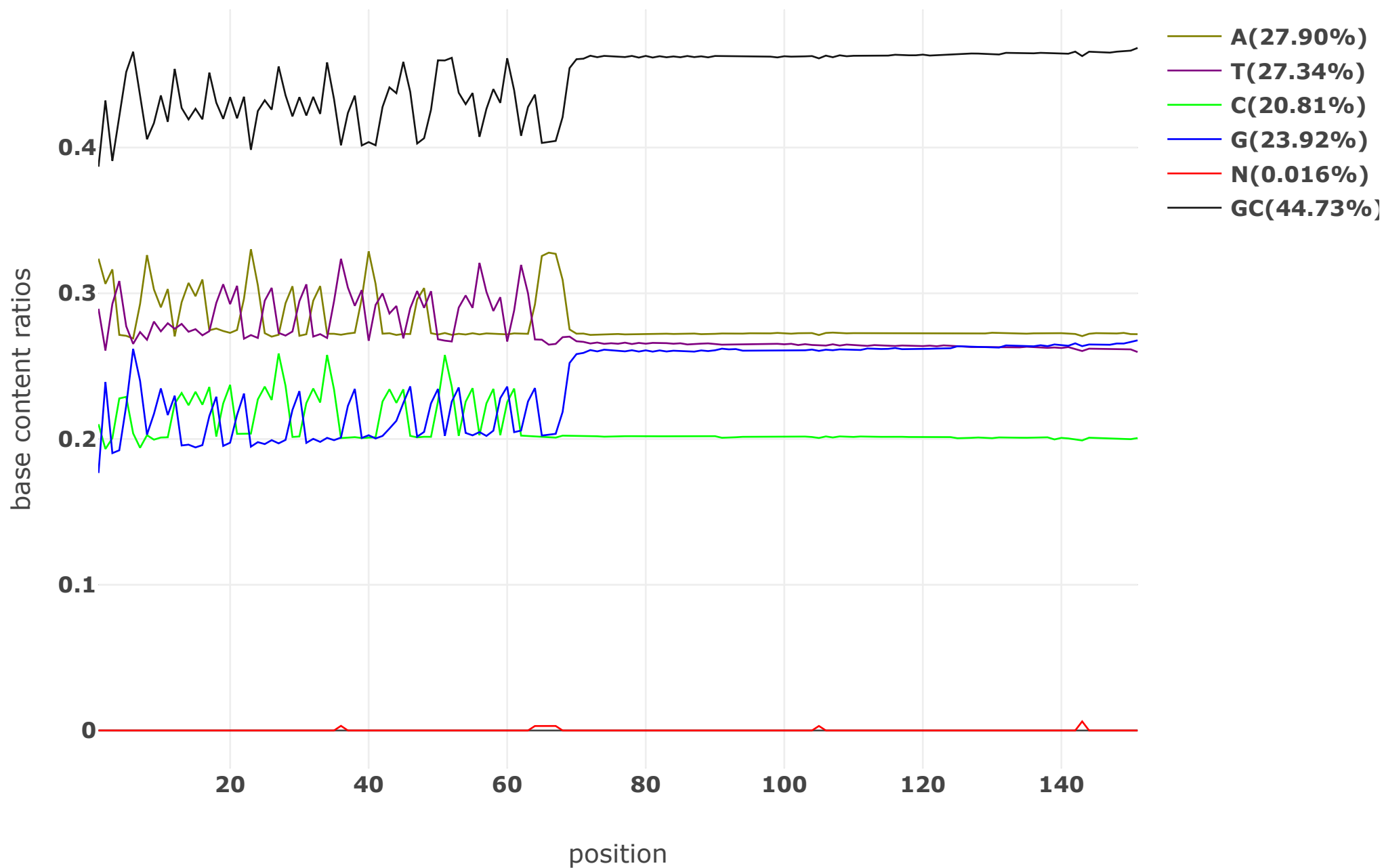
Value of each position will be shown on mouse over.





Before filtering: read1: base contents

Value of each position will be shown on mouse over.



Before filtering: read1: KMER counting















Darker background means larger counts. The count will be shown on mouse over.

[illegible]

AGA	AGAAA	AGAAT	AGAAC	AGAAG	AGATA	AGATT	AGATC	AGATG	AGACA	AGACT	AGACC	AGACG	AGAGA	AGAGT	AGAGC	AGAGG
AGT	AGTAA	AGTAT	AGTAC	AGTAG	AGTTA	AGTTT	AGTTC	AGTTG	AGTCA	AGTCT	AGTCC	AGTCG	AGTGA	AGTGT	AGTGC	AGTGG
AGC	AGCAA	AGCAT	AGCAC	AGCAG	AGCTA	AGCTT	AGCTC	AGCTG	AGCCA	AGCCT	AGCCC	AGCCG	AGCGA	AGCGT	AGCGC	AGCGG
AGG	AGGAA	AGGAT	AGGAC	AGGAG	AGGTA	AGGTT	AGGTC	AGGTG	AGGCA	AGGCT	AGGCC	AGGCG	AGGGA	AGGGT	AGGGC	AGGGG
TAA	TAAAA	TAAAT	TAAAC	TAAAG	TAATA	TAATT	TAATC	TAATG	TAACA	TAACT	TAACC	TAACG	TAAGA	TAAGT	TAAGC	TAAGG
TAT	TATAA	TATAT	TATAC	TATAG	TATTA	TATTT	TATTC	TATTG	TATCA	TATCT	TATCC	TATCG	TATGA	TATGT	TATGC	TATGG
TAC	TACAA	TACAT	TACAC	TACAG	TACTA	TACTT	TACTC	TACTG	TACCA	TACCT	TACCC	TACCG	TACGA	TACGT	TACGC	TACGG
TAG	TAGAA	TAGAT	TAGAC	TAGAG	TAGTA	TAGTT	TAGTC	TAGTG	TAGCA	TAGCT	TAGCC	TAGCG	TAGGA	TAGGT	TAGGC	TAGGG
TTA	TTAAA	TTAAT	TTAAC	TTAAG	TTATA	TTATT	TTATC	TTATG	TTACA	TTACT	TTACC	TTACG	TTAGA	TTAGT	TTAGC	TTAGG
TTT	TTTAA	TTTAT	TTTAC	TTTAG	TTTTA	TTTTT	TTTTC	TTTTG	TTTCA	TTTCT	TTTCC	TTTCG	TTTGA	TTTGT	TTTGC	TTTGG
TTC	TTCAA	TTCAT	TTCAC	TTCAG	TTCTA	TTCTT	TTCTC	TTCTG	TTCCA	TTCCT	TTCCC	TTCCG	TTCGA	TTCGT	TTCGC	TTCGG
TTG	TTGAA	TTGAT	TTGAC	TTGAG	TTGTA	TTGTT	TTGTC	TTGTG	TTGCA	TTGCT	TTGCC	TTGCG	TTGGA	TTGGT	TTGGC	TTGGG
TCA	TCAAA	TCAAT	TCAAC	TCAAG	TCATA	TCATT	TCATC	TCATG	TCACA	TCACT	TCACC	TCACG	TCAGA	TCAGT	TCAGC	TCAGG
TCT	TCTAA	TCTAT	TCTAC	TCTAG	TCTTA	TCTTT	TCTTC	TCTTG	TCTCA	TCTCT	TCTCC	TCTCG	TCTGA	TCTGT	TCTGC	TCTGG
TCC	TCCAA	TCCAT	TCCAC	TCCAG	TCCTA	TCCTT	TCCTC	TCCTG	TCCCA	TCCCT	TCCCC	TCCCG	TCCGA	TCCGT	TCCGC	TCCGG
TCG	TCGAA	TCGAT	TCGAC	TCGAG	TCGTA	TCGTT	TCGTC	TCGTG	TCGCA	TCGCT	TCGCC	TCGCG	TCGGA	TCGGT	TCGGC	TCGGG
TGA	TGAAA	TGAAT	TGAAC	TGAAG	TGATA	TGATT	TGATC	TGATG	TGACA	TGACT	TGACC	TGACG	TGAGA	TGAGT	TGAGC	TGAGG
TGT	TGTAA	TGTAT	TGTAC	TGTAG	TGTTA	TGTTT	TGTTC	TGTTG	TGTCA	TGTCT	TGTCC	TGTCTG	TGTGA	TGTGT	TGTGC	TGTGG
TGC	TGCAA	TGCAT	TGCAC	TGCAG	TGCTA	TGCTT	TGCTC	TGCTG	TGCCA	TGCCT	TGCCC	TGCCG	TGCGA	TGCGT	TGCGC	TGCGG
TGG	TGGAA	TGGAT	TGGAC	TGGAG	TGGTA	TGGTT	TGGTC	TGGTG	TGGCA	TGGCT	TGGCC	TGGCG	TGGGA	TGGGT	TGGGC	TGGGG
CAA	CAAAA	CAAAAT	CAAAC	CAAAG	CAATA	CAATT	CAATC	CAATG	CAACA	CAACT	CAACC	CAACG	CAAGA	CAAGT	CAAGC	CAAGG
CAT	CATAA	CATAT	CATAC	CATAG	CATTA	CATTT	CATTC	CATTG	CATCA	CATCT	CATCC	CATCG	CATGA	CATGT	CATGC	CATGG
CAC	CACAA	CACAT	CACAC	CACAG	CACTA	CACCT	CACCTC	CACCTG	CACCA	CACCT	CACCC	CACCG	CACGA	CACGT	CACGC	CACGG
CAG	CAGAA	CAGAT	CAGAC	CAGAG	CAGTA	CAGTT	CAGTC	CAGTG	CAGCA	CAGCT	CAGCC	CAGCG	CAGGA	CAGGT	CAGGC	CAGGG
CTA	CTAAA	CTAAT	CTAAC	CTAAG	CTATA	CTATT	CTATC	CTATG	CTACA	CTACT	CTACC	CTACG	CTAGA	CTAGT	CTAGC	CTAGG
CTT	CTTAA	CTTAT	CTTAC	CTTAG	CTTTA	CTTTT	CTTTC	CTTTG	CTTCA	CTTCT	CTTCC	CTTCG	CTTGA	CTTGT	CTTGC	CTTGG
CTC	CTCAA	CTCAT	CTCAC	CTCAG	CTCTA	CTCTT	CTCTC	CTCTG	CTCCA	CTCCT	CTCCC	CTCCG	CTCGA	CTCGT	CTCGC	CTCGG
CTG	CTGAA	CTGAT	CTGAC	CTGAG	CTGTA	CTGTT	CTGTG	CTGTG	CTGCA	CTGCT	CTGCC	CTGCG	CTGGA	CTGGT	CTGGC	CTGGG
CCA	CCAAA	CCAAT	CCAAC	CCAAG	CCATA	CCATT	CCATC	CCATG	CCACA	CCACT	CCACC	CCACG	CCAGA	CCAGT	CCAGC	CCAGG
CCT	CCTAA	CCTAT	CCTAC	CCTAG	CCTTA	CCTTT	CCTTC	CCTTG	CCTCA	CCTCT	CCTCC	CCTCG	CCTGA	CCTGT	CCTGC	CCTGG
CCC	CCCAA	CCCAT	CCCAC	CCCAG	CCCTA	CCCTT	CCCTC	CCCTG	CCCCA	CCCCT	CCCCC	CCCCG	CCCGA	CCCGT	CCCGC	CCCGG
CCG	CCGAA	CCGAT	CCGAC	CCGAG	CCGTA	CCGTT	CCGTC	CCGTG	CCGCA	CCGCT	CCGCC	CCGCG	CCGGA	CCGGT	CCGGC	CCGGG
CGA	CGAAA	CGAAT	CGAAC	CGAAG	CGATA	CGATT	CGATC	CGATG	CGACA	CGACT	CGACC	CGACG	CGAGA	CGAGT	CGAGC	CGAGG
CGT	CGTAA	CGTAT	CGTAC	CGTAG	CGTTA	CGTTT	CGTTC	CGTTG	CGTCA	CGTCT	CGTCC	CGTCG	CGTGA	CGTGT	CGTGC	CGTGG
CGC	CGCAA	CGCAT	CGCAC	CGCAG	CGCTA	CGCTT	CGCTC	CGCTG	CGCCA	CGCCT	CGCCC	CGCCG	CGCGA	CGCGT	CGCGC	CGCGG
CGG	CGGAA	CGGAT	CGGAC	CGGAG	CGGTA	CGGTT	CGGTC	CGGTG	CGGCA	CGGCT	CGGCC	CGGCG	CGGGA	CGGGT	CGGGC	CGGGG
GAA	GAAAA	GAAAT	GAAAC	GAAAG	GAATA	GAATT	GAATC	GAATG	GAACA	GAACT	GAACC	GAACG	GAAGA	GAAGT	GAAGC	GAAGG
GAT	GATAA	GATAT	GATAC	GATAG	GATTA	GATTT	GATTC	GATTG	GATCA	GATCT	GATCC	GATCG	GATGA	GATGT	GATGC	GATGG
GAC	GACAA	GACAT	GACAC	GACAG	GACTA	GACTT	GACTC	GACTG	GACCA	GACCT	GACCC	GACCG	GACGA	GACGT	GACGC	GACGG
GAG	GAGAA	GAGAT	GAGAC	GAGAG	GAGTA	GAGTT	GAGTC	GAGTG	GAGCA	GAGCT	GAGCC	GAGCG	GAGGA	GAGGT	GAGGC	GAGGG
GTA	GTAAA	GTAAT	GTAAC	GTAAG	GTATA	GTATT	GTATC	GTATG	GTACA	GTACT	GTACC	GTACG	GTAGA	GTAGT	GTAGC	GTAGG
GTT	GTTAA	GTTAT	GTTAC	GTTAG	GTTTA	GTTTT	GTTTC	GTTTG	GTTCA	GTTCT	GTTCC	GTTCTG	GTTGA	GTTGT	GTTGC	GTTGG
GTC	GTCAA	GTCAT	GTCAC	GTCAG	GTCTA	GTCTT	GTCTC	GTCTG	GTCCA	GTCTT	GTCCC	GTCCG	GTCGA	GTCGT	GTCGC	GTCGG
GTG	GTGAA	GTGAT	GTGAC	GTGAG	GTGTA	GTGTT	GTGTC	GTGTG	GTGCA	GTGCT	GTGCC	GTGCG	GTGGA	GTGGT	GTGGC	GTGGG
GCA	GCAAA	GCAAT	GCAAC	GCAAG	GCATA	GCATT	GCATC	GCATG	GCACA	GCACT	GCACC	GCACG	GCAGA	GCAGT	GCAGC	GCAGG
GCT	GCTAA	GCTAT	GCTAC	GCTAG	GCTTA	GCTTT	GCTTC	GCTTG	GCTCA	GCTCT	GCTCC	GCTCG	GCTGA	GCTGT	GCTGC	GCTGG
GCC	GCCAA	GCCAT	GCCAC	GCCAG	GCCTA	GCCTT	GCCTC	GCCTG	GCCCA	GCCCT	GCCCC	GCCCG	GCCGA	GCCGT	GCCGC	GCCGG
GCG	GCGAA	GCGAT	GCGAC	GCGAG	GCGTA	GCGTT	GCGTC	GCGTG	GCGCA	GCGCT	GCGCC	GCGCG	GCGGA	GCGGT	GCGGC	GCGGG
GGA	GGAAA	GGAAT	GGAAC	GGAAG	GGATA	GGATT	GGATC	GGATG	GGACA	GGACT	GGACC	GGACG	GGAGA	GGAGT	GGAGC	GGAGG
GGT	GGTAA	GGTAT	GGTAC	GGTAG	GGTTA	GGTTT	GGTTC	GGTTG	GGTCA	GGTCT	GGTCC	GGTCG	GGTGA	GGTGT	GGTGC	GGTGG
GGC	GGCAA	GGCAT	GGCAC	GGCAG	GGCTA	GGCTT	GGCTC	GGCTG	GGCCA	GGCCT	GGCCC	GGCCG	GGCGA	GGCGT	GGCGC	GGCGG
GGG	GGGAA	GGGAT	GGGAC	GGGAG	GGGTA	GGGTT	GGGTC	GGGTG	GGGCA	GGGCT	GGGCC	GGGCG	GGGGA	GGGGT	GGGGC	GGGGG

Before filtering: read1: overrepresented sequences

Sampling rate: 1 / 20

overrepresented sequence	count (% of bases)	distribution: cycle 1 ~ cycle 151
AAAGATGTGTATAAGAGACA	86282 (0.102329%)	
AAGATGTGTATAAGAGACAG	149980 (0.177874%)	
AAGCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	17427 (0.041336%)	
ACACACACACACACACACACACACACACACACACAA	2326 (0.005517%)	
ACACACACACACACACACACACACACACACACACAC	58007 (0.137591%)	
ACACACACACACACACACACACACACACACACACAG	1568 (0.003719%)	
ACACACACACACACACACACACCTGTCTCTTATACACATC	15516 (0.036804%)	
ACACACACACACCTGTCTCTTATACACATCTAGATGTGTA	11379 (0.026991%)	
ACACACACACCTGTCTCTTATACACATCTAGATGTGTATA	1309 (0.003105%)	
ACACACACCTGTCTCTTATACACATCTAGATGTGTATAAG	1667 (0.003954%)	
ACACACCTGTCTCTTATACACATCTAGATGTGTATAAGAG	3488 (0.008273%)	
ACACATCTAGATGTGTATAAGAGACAGCACACACACAC	157 (0.000372%)	
ACACATCTAGATGTGTATAAGAGACAGGTGTGTGTGTGTG	157 (0.000372%)	
ACACCTGTCTCTTATACACATCTAGATGTGTATAAGAGAC	21275 (0.050464%)	

[illegible]

[illegible]

[illegible]

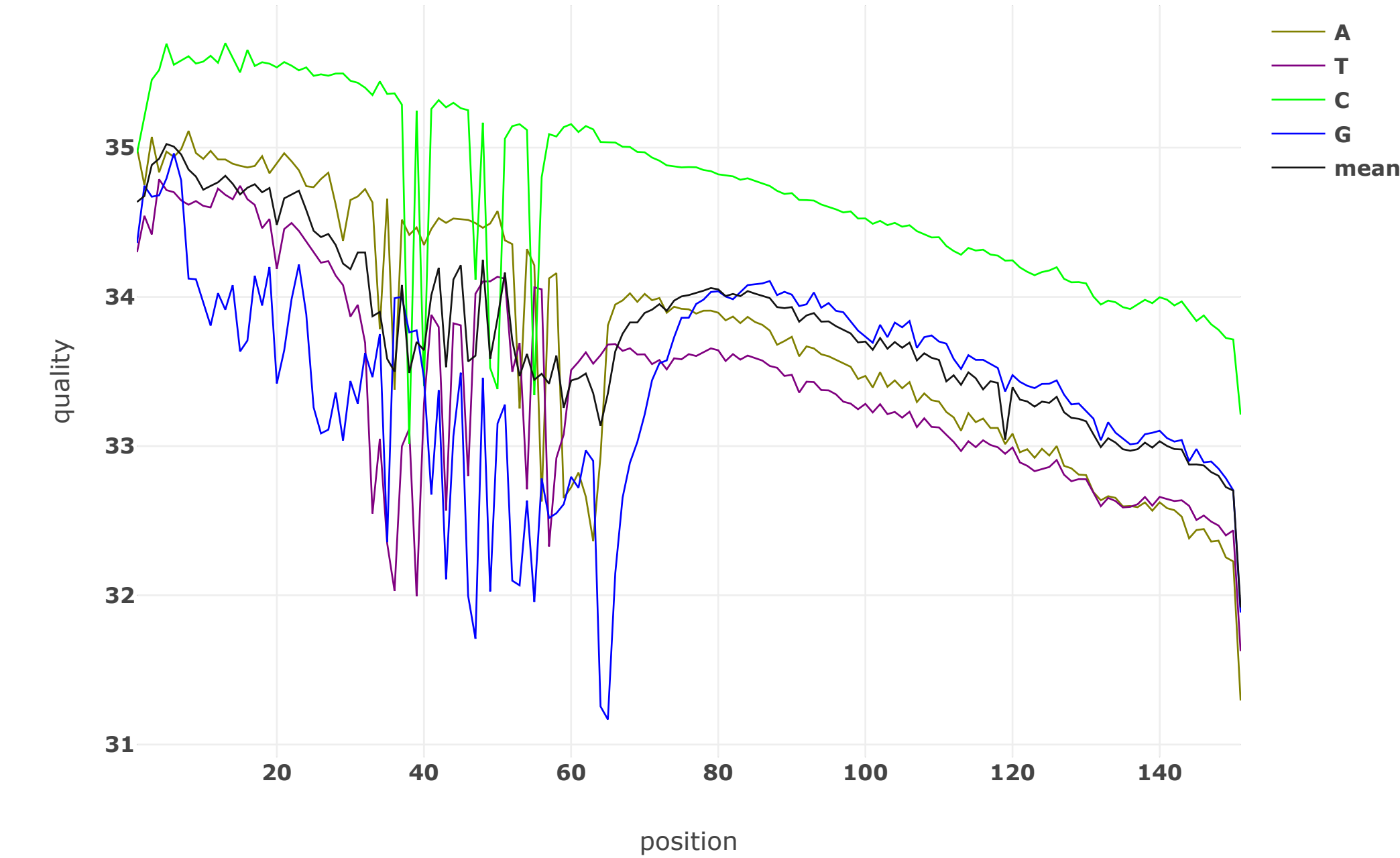
[illegible]

[illegible]

TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCCC	81 (0.000192%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCCT	88 (0.000209%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCTC	95 (0.000225%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCTG	64 (0.000152%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCTT	56 (0.000133%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGAA	68 (0.000161%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGAG	86 (0.000204%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGAT	79 (0.000187%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGCA	62 (0.000147%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGCT	95 (0.000225%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGGG	101 (0.000240%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGGT	88 (0.000209%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGTA	137 (0.000325%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGTC	78 (0.000185%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGTG	261 (0.000619%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGTT	99 (0.000235%)	
TGTCTCTTATACACATCTCA	891 (0.001057%)	
TGTCTCTTATACACATCTCT	623 (0.000739%)	
TGTCTCTTATACACATCTTG	655 (0.000777%)	
TGTCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	17648 (0.041861%)	
TGTGCTGTCTCTTATACACATCTAGATGTGTATAAGAGAC	11772 (0.027923%)	
TGTGTATAAGAGACAGCACACACACACACACACACACA	36 (0.000085%)	
TGTGTATAAGAGACAGGTGTGTGTGTGTGTGTGTGTGT	49 (0.000116%)	
TGTGTGCTGTCTCTTATACACATCTAGATGTGTATAAGAG	4414 (0.010470%)	
TGTGTGCTGTCTCTTATACACATCTAGATGTGTATAAG	2093 (0.004965%)	
TGTGTGCTGTCTCTTATACACATCTAGATGTGTATA	3993 (0.009471%)	
TGTGTGCTGTCTCTTATACACATCTAGATGTGTA	13161 (0.031218%)	
TGTGTGCTGTGTGTGTGTGTGTGTGTGTCTCTTATACATC	16029 (0.038020%)	
TGTGTGCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTG	39916 (0.094680%)	
TGTGTGCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTT	1082 (0.002566%)	
TTATACACATCTAGATGTGTATAAGAGACAGCACACACAC	70 (0.000166%)	
TTATACACATCTAGATGTGTATAAGAGACAGGTGTGTGTG	36 (0.000085%)	
TTCCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	20890 (0.049551%)	

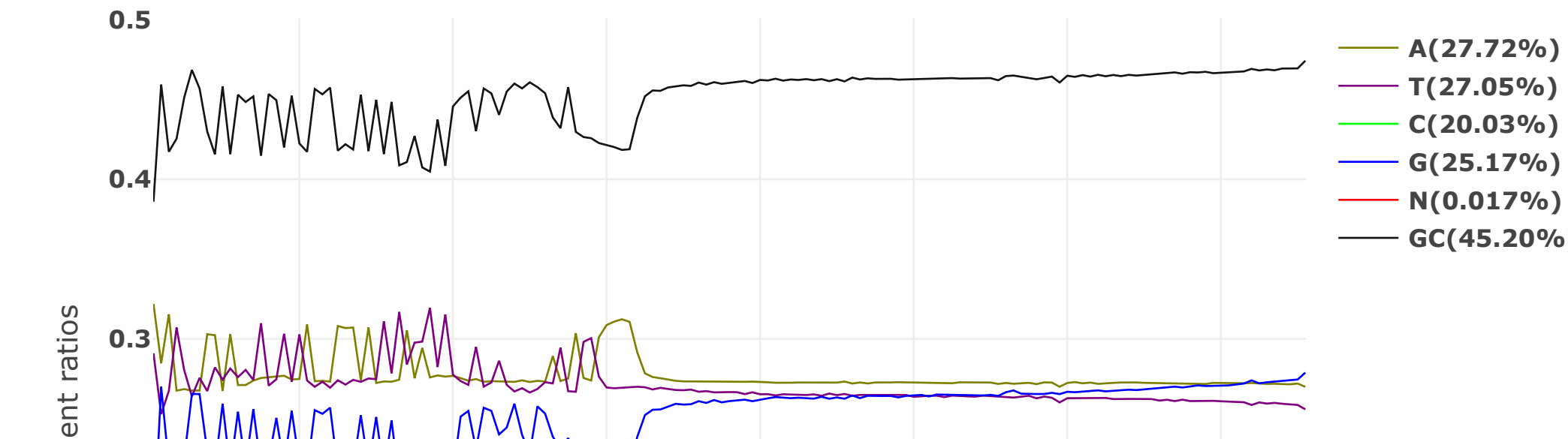
Before filtering: read2: quality

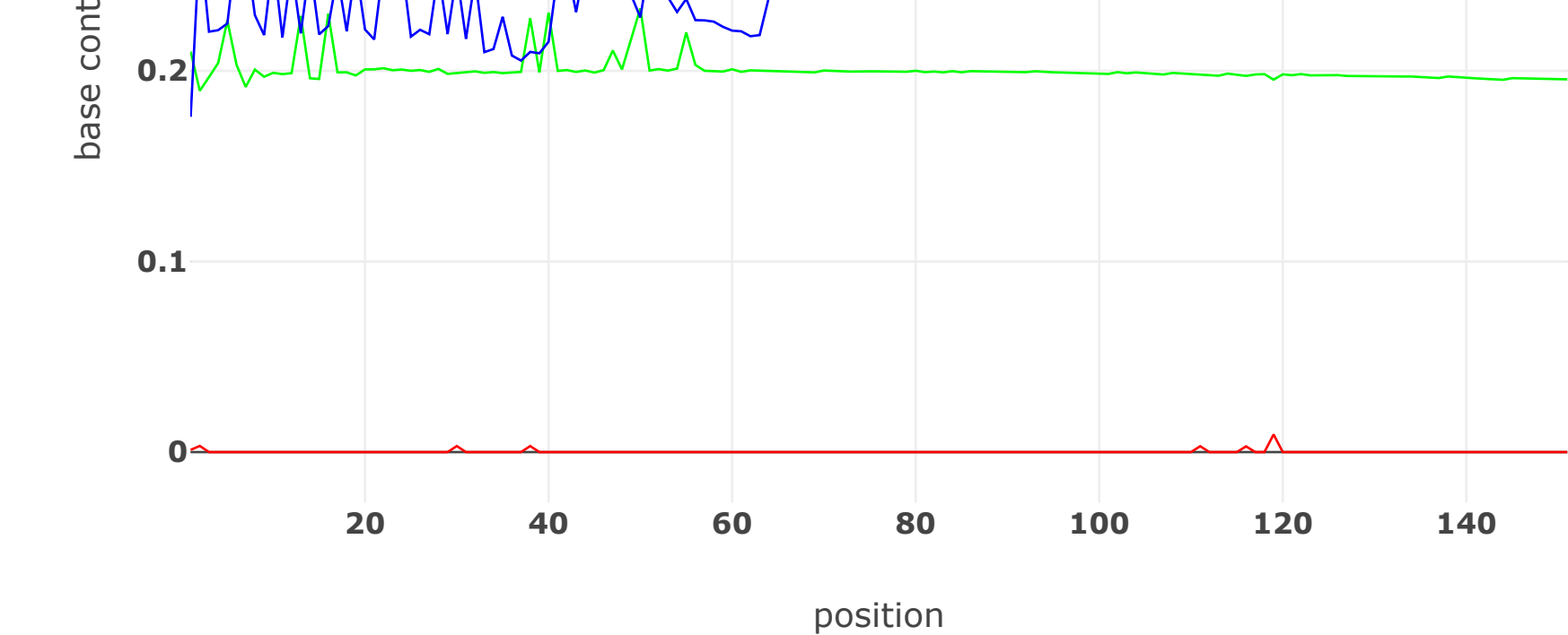
Value of each position will be shown on mouse over.



Before filtering: read2: base contents

Value of each position will be shown on mouse over.





Before filtering: read2: KMER counting

Darker background means larger counts. The count will be shown on mouse over.

	AA	AT	AC	AG	TA	TT	TC	TG	CA	CT	CC	CG	GA	GT	GC	GG
AAA	AAAAA	AAAAT	AAAAC	AAAAG	AAATA	AAATT	AAATC	AAATG	AAACA	AAACT	AAACC	AAACG	AAAGA	AAAGT	AAAGC	AAAGG
AAT	AATAA	AATAT	AATAC	AATAG	AATTA	AATTT	AATTC	AATTG	AATCA	AATCT	AATCC	AATCG	AATGA	AATGT	AATGC	AATGG
AAC	AACAA	AACAT	AACAC	AACAG	AACTA	AACTT	AACTC	AACTG	AACCA	AACCT	AACCC	AACCG	AACGA	AACGT	AACGC	AACGG
AAG	AAGAA	AAGAT	AAGAC	AAGAG	AAGTA	AAGTT	AAGTC	AAGTG	AAGCA	AAGCT	AAGCC	AAGCG	AAGGA	AAGGT	AAGGC	AAGGG
ATA	ATAAA	ATAAT	ATAAC	ATAAG	ATATA	ATATT	ATATC	ATATG	ATACA	ATACT	ATACC	ATACG	ATAGA	ATAGT	ATAGC	ATAGG
ATT	ATTAA	ATTAT	ATTAC	ATTAG	ATTTA	ATTTT	ATTTT	ATTTG	ATTCA	ATTCT	ATTCC	ATTCT	ATTGA	ATTGT	ATTGC	ATTGG
ATC	ATCAA	ATCAT	ATCAC	ATCAG	ATCTA	ATCTT	ATCTC	ATCTG	ATCCA	ATCCT	ATCCC	ATCCG	ATCGA	ATCGT	ATCGC	ATCGG
ATG	ATGAA	ATGAT	ATGAC	ATGAG	ATGTA	ATGTT	ATGTC	ATGTG	ATGCA	ATGCT	ATGCC	ATGCG	ATGGA	ATGGT	ATGGC	ATGGG
ACA	ACAAA	ACAAT	ACAAC	ACAAG	ACATA	ACATT	ACATC	ACATG	ACACA	ACACT	ACACC	ACACG	ACAGA	ACAGT	ACAGC	ACAGG
ACT	ACTAA	ACTAT	ACTAC	ACTAG	ACTTA	ACTTT	ACTTC	ACTTG	ACTCA	ACTCT	ACTCC	ACTCG	ACTGA	ACTGT	ACTGC	ACTGG
ACC	ACCAA	ACCAT	ACCAC	ACCAG	ACCTA	ACCTT	ACCTC	ACCTG	ACCCA	ACCCT	ACCCC	ACCCG	ACCGA	ACCGT	ACCGC	ACCGG
ACG	ACGAA	ACGAT	ACGAC	ACGAG	ACGTA	ACGTT	ACGTC	ACGTG	ACGCA	ACGCT	ACGCC	ACGCG	ACGGA	ACGGT	ACGGC	ACGGG
AGA	AGAAA	AGAAT	AGAAC	AGAAG	AGATA	AGATT	AGATC	AGATG	AGACA	AGACT	AGACC	AGACG	AGAGA	AGAGT	AGAGC	AGAGG
AGT	AGTAA	AGTAT	AGTAC	AGTAG	AGTTA	AGTTT	AGTTC	AGTTG	AGTCA	AGTCT	AGTCC	AGTCG	AGTGA	AGTGT	AGTGC	AGTGG
AGC	AGCAA	AGCAT	AGCAC	AGCAG	AGCTA	AGCTT	AGCTC	AGCTG	AGCCA	AGCCT	AGCCC	AGCCG	AGCGA	AGCGT	AGCGC	AGCGG
AGG	AGGAA	AGGAT	AGGAC	AGGAG	AGGTA	AGGTT	AGGTC	AGGTG	AGGCA	AGGCT	AGGCC	AGGCG	AGGGA	AGGGT	AGGGC	AGGGG
TAA	TAAAA	TAAAT	TAAAC	TAAAG	TAATA	TAATT	TAATC	TAATG	TAACA	TAACT	TAACC	TAACG	TAAGA	TAAGT	TAAGC	TAAGG
TAT	TATAA	TATAT	TATAC	TATAG	TATTA	TATTT	TATTC	TATTG	TATCA	TATCT	TATCC	TATCG	TATGA	TATGT	TATGC	TATGG
TAC	TACAA	TACAT	TACAC	TACAG	TACTA	TACTT	TACTC	TACTG	TACCA	TACCT	TACCC	TACCG	TACGA	TACGT	TACGC	TACGG
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Before filtering: read2: overrepresented sequences

Sampling rate: 1 / 20

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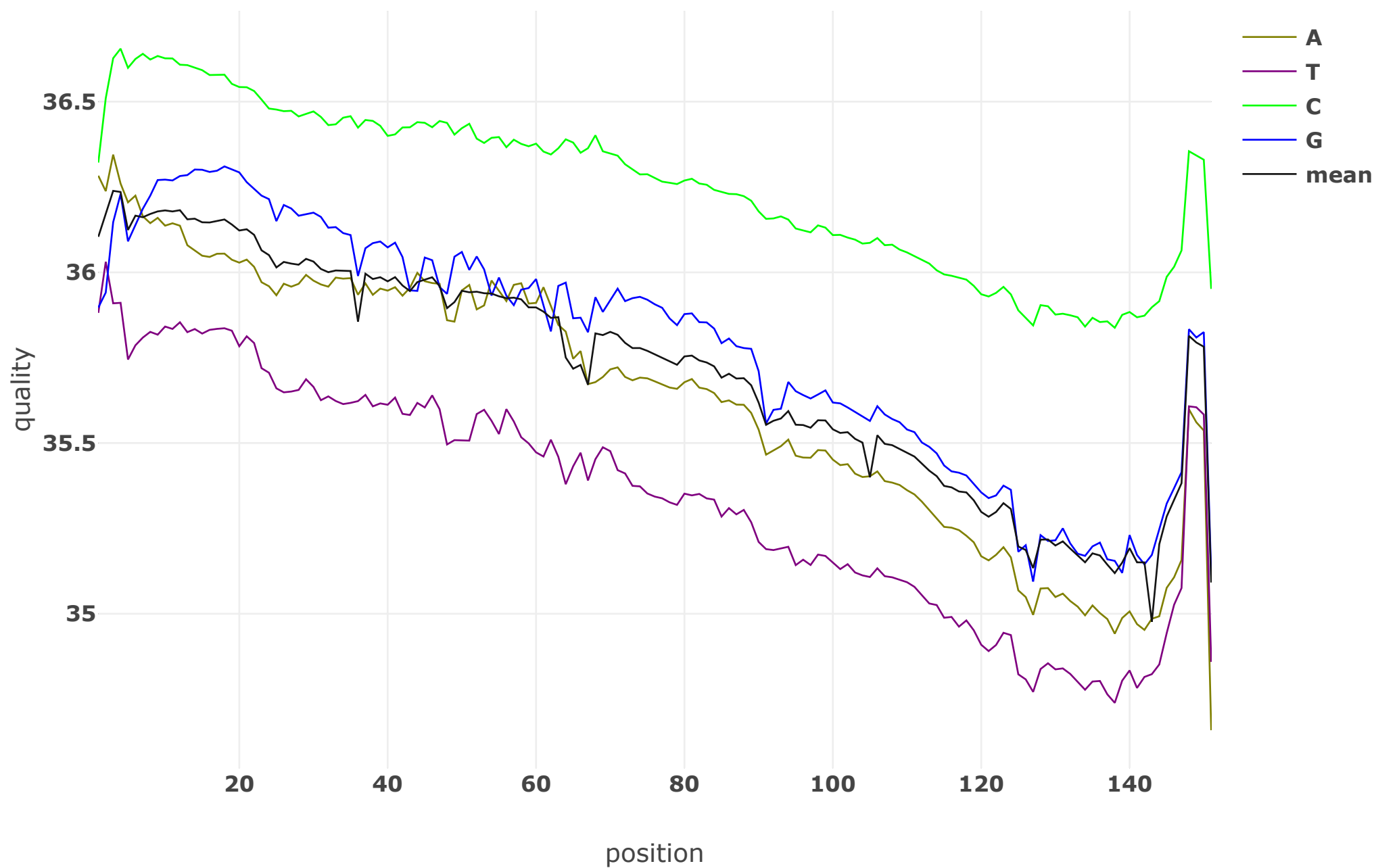
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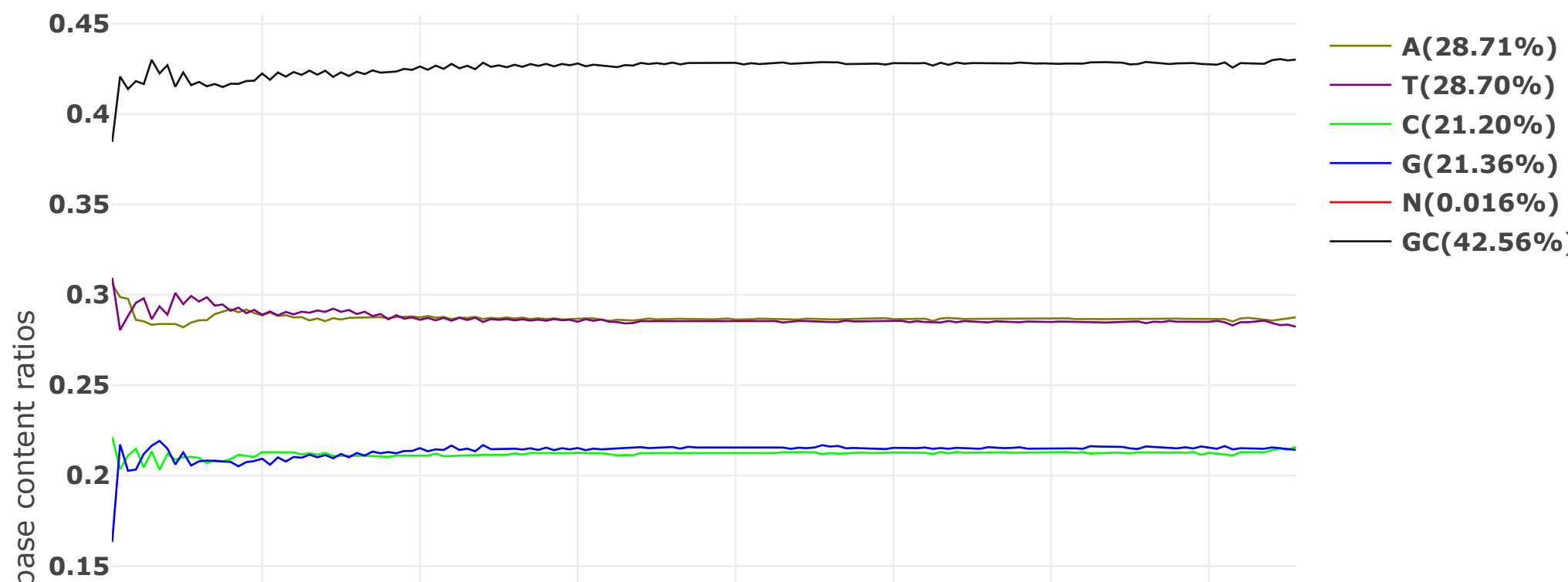
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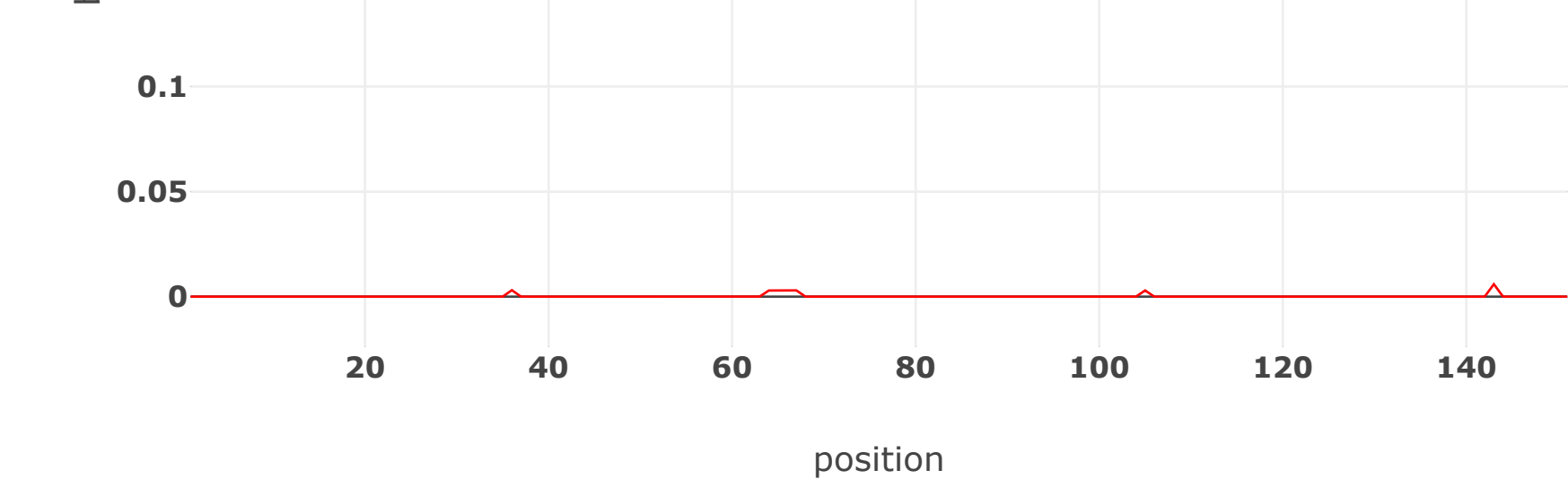
Value of each position will be shown on mouse over.



After filtering: read1: base contents

Value of each position will be shown on mouse over.





After filtering: read1: KMER counting

















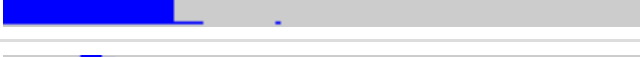
















Darker background means larger counts. The count will be shown on mouse over.


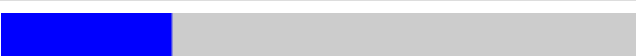


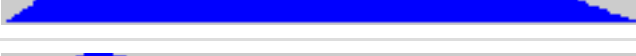
























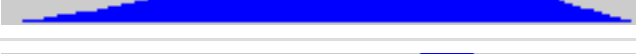




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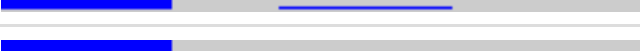

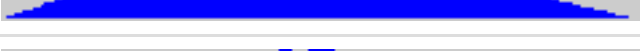

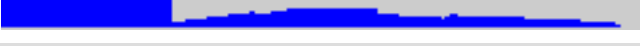














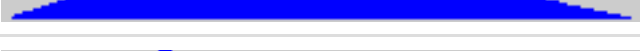






After filtering: read1: overrepresented sequences

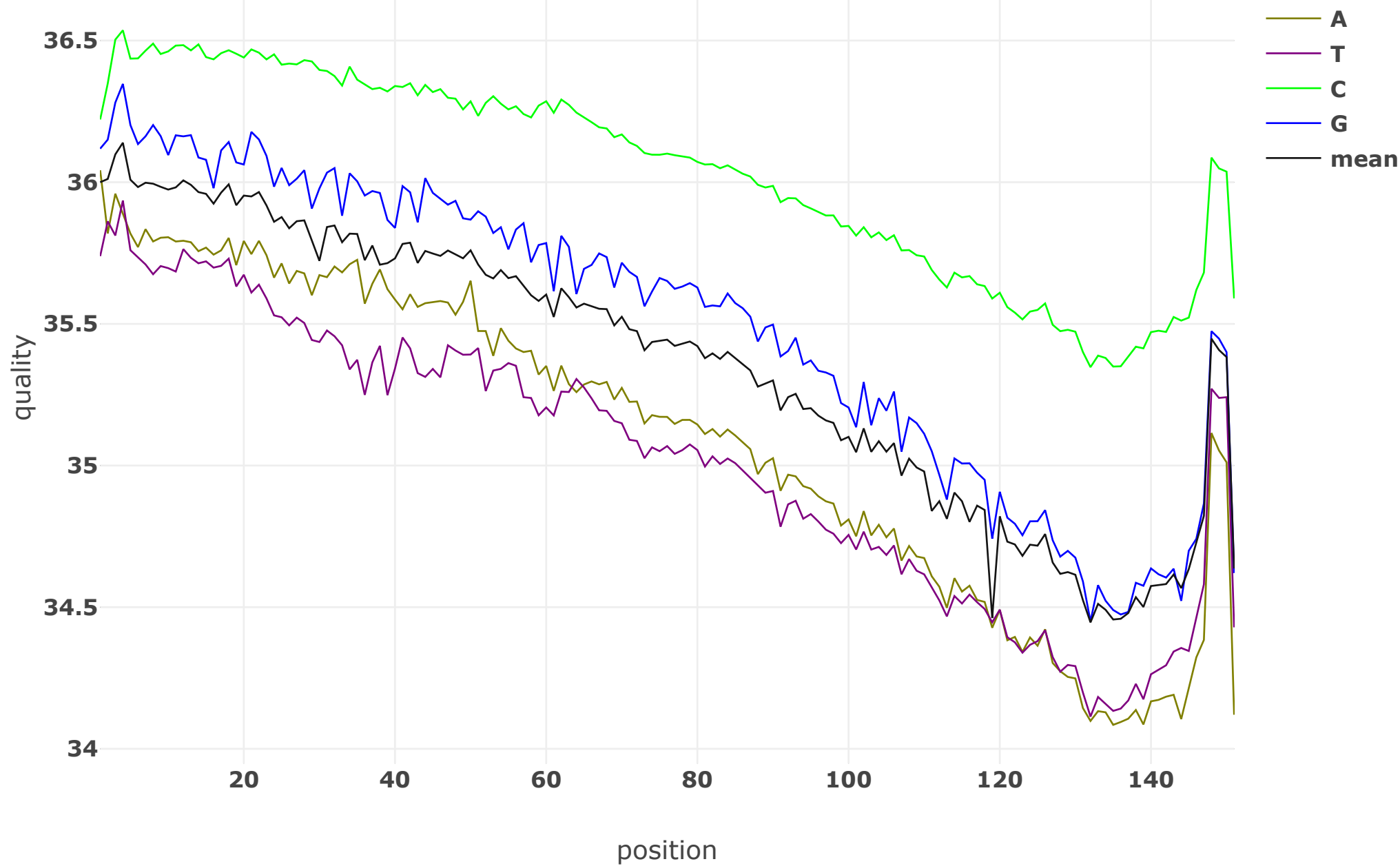
Sampling rate: 1 / 20

overrepresented sequence	count (% of bases)	distribution: cycle 1 ~ cycle 151
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ATGTGTATAAGAGACAGGTGTGTGTGTGTGTGTGTGTGTG	70 (0.000223%)	
ATGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT	12830 (0.040923%)	
CAAGATGTGTATAAGAGACA	93357 (0.148886%)	
CACACACACACACACACACACACACACACACACACAAA	549 (0.001751%)	
CACACACACACACACACACACACACACACACACACACA	10934 (0.034875%)	
CACACACACACACACACACCTGTCTCTTATACACATCT	1550 (0.004944%)	
CACACACACACCTGTCTCTTATACACATCTAGATGTGTAT	1243 (0.003965%)	
CACACACACCTGTCTCTTATACACATCTAGATGTGTATAA	973 (0.003103%)	
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CACACCTGTCTCTTATACACATCTAGATGTGTATAAGAGA	5322 (0.016975%)	
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CAGATGTGTATAAGAGACAG	100940 (0.160979%)	
CATCTAGATGTGTATAAGAGACAGGTGTGTGTGTGTGTGT	136 (0.000434%)	
CCAGATGTGTATAAGAGACA	92686 (0.147816%)	
CCCCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	19727 (0.062921%)	
CCCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	39315 (0.125399%)	
CCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGA	717 (0.002287%)	
CCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGC	1580 (0.005040%)	
CCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGG	2231 (0.007116%)	
CCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGT	340 (0.001084%)	
CTCACACACACACACACACACACACACACACACACACA	8885 (0.028340%)	
CTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT	2731 (0.008711%)	
CTCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	20355 (0.064924%)	
CTCTTATACACATCTAGATGTGTATAAGAGACAGCACACA	50 (0.000159%)	
CTCTTATACACATCTAGATGTGTATAAGAGACAGGTGTGT	40 (0.000128%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGAC	2393 (0.007633%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGAT	2816 (0.008982%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCA	517 (0.001649%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCC	305 (0.000973%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCT	368 (0.001174%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGA	297 (0.000947%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGC	315 (0.001005%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGG		

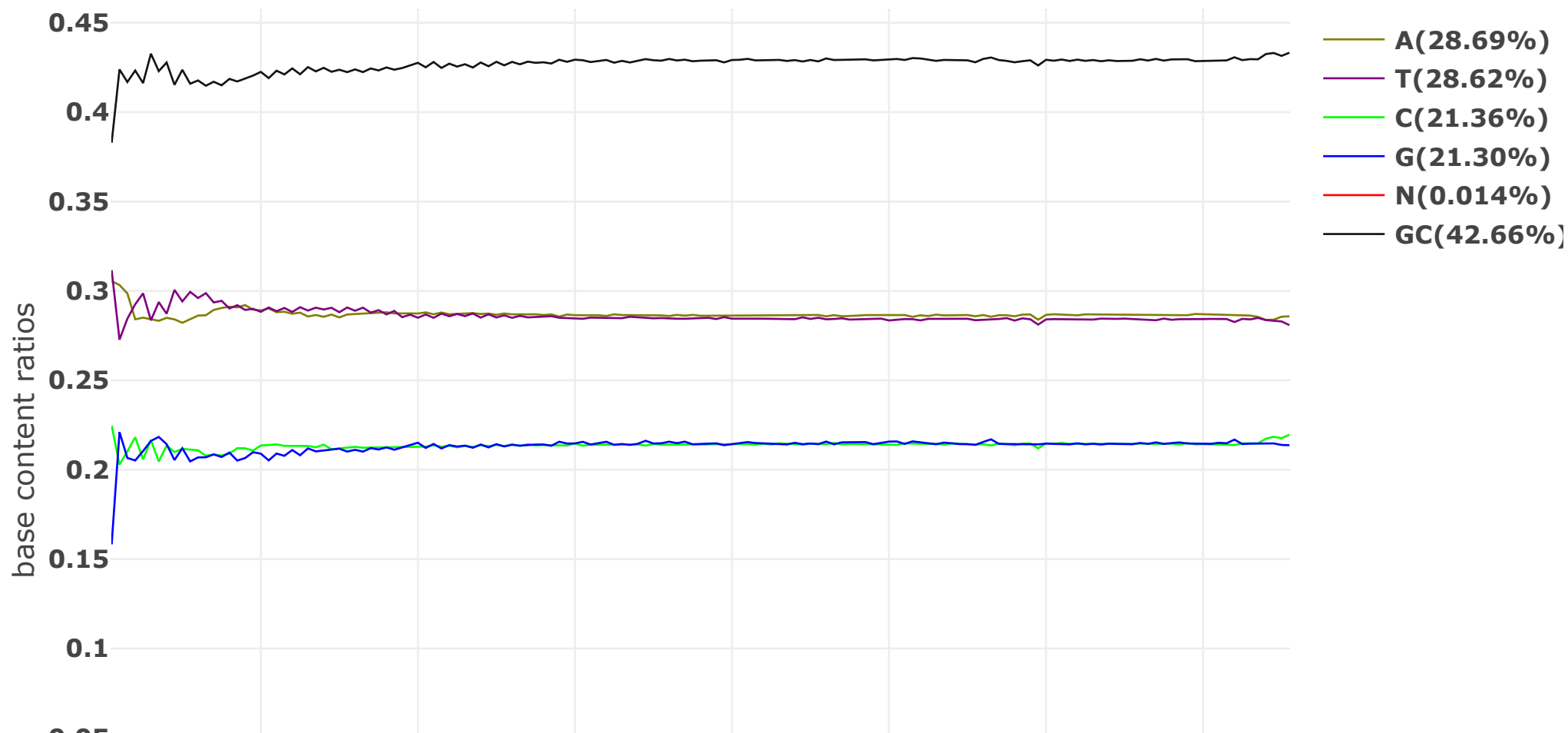
	399 (0.001273%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGT	685 (0.002185%)	
CTGTCTCTTATACACATCTC	328033 (0.523147%)	
CTGTCTCTTATACACATCTG	284417 (0.453588%)	
CTGTCTCTTATACACATCTT	297700 (0.474772%)	
CTTATACACATCTAGATGTGTATAAGAGACAGCACACACA	38 (0.000121%)	
CTTATACACATCTAGATGTGTATAAGAGACAGGTGTGTGT	31 (0.000099%)	
GACCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	25552 (0.081501%)	
GACTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	23644 (0.075415%)	
GA	7522 (0.023992%)	
GA	187 (0.001491%)	
GAGATGTGTATAAGAGACAG	205762 (0.328149%)	
GAGCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	21446 (0.068404%)	
GATGTGTATAAGAGACAGCACACACACACACACACACACA	93 (0.000297%)	
GATGTGTATAAGAGACAGGTGTGTGTGTGTGTGTGTGTGT	120 (0.000383%)	
GCACACACACACACACACACACACACACACACACACACAC	14496 (0.046236%)	
GCCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	6903 (0.022018%)	
GCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGA	2376 (0.007578%)	
GCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGC	4773 (0.015224%)	
GCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGG	7441 (0.023734%)	
GCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGT	1251 (0.003990%)	
GGCCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	22985 (0.073313%)	
GGCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	18209 (0.058079%)	
GGGCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	22132 (0.070592%)	
GGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTG	6891 (0.021980%)	
GTATAAGAGACAGGTGTGTGTGTGTGTGTGTGTGTGTGTG	28 (0.000089%)	
GTCTCTTATACACATCTAGATGTGTATAAGAGACAGCACA	88 (0.000281%)	
GTCTCTTATACACATCTAGATGTGTATAAGAGACAGGTAT	32 (0.000102%)	
GTCTCTTATACACATCTAGATGTGTATAAGAGACAGGTGT	109 (0.000348%)	
GTCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	25434 (0.081124%)	
GTGCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	14445 (0.046074%)	
GTGTATAAGAGACAGGTGTGTGTGTGTGTGTGTGTGTGTG	58 (0.000185%)	
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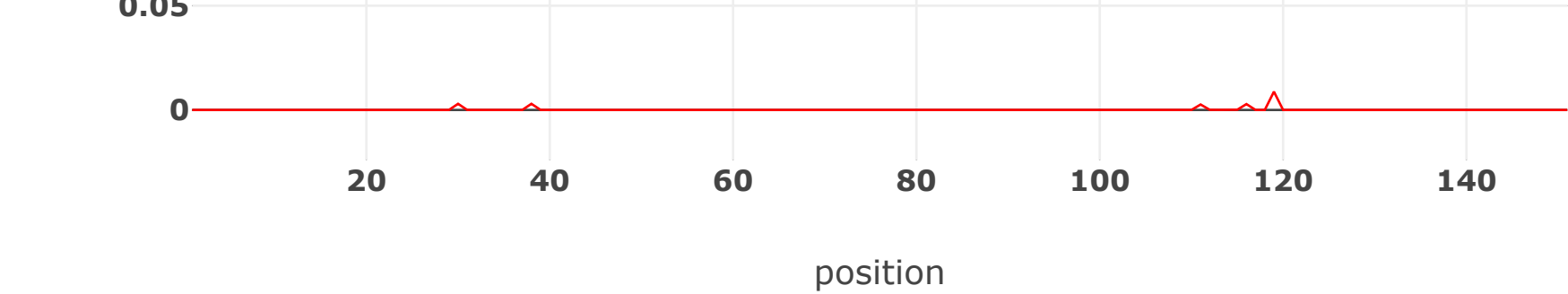
		
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GTGTGTGTGTGTGTGTGTGTGCTGTCTCTTATACACATCT	1293 (0.004124%)	
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GT	17947 (0.057244%)	
TACACACACACACACACACA	67853 (0.108212%)	
TACACATCTAGATGTGTATAAGAGACAGCACACACACACA	117 (0.000373%)	
TACACATCTAGATGTGTATAAGAGACAGGTGTGTGTGTGT	84 (0.000268%)	
TACCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	33619 (0.107231%)	
TATAAGAGACAGGTGTGTGTGTGTGTGTGTGTGTGTGT	14 (0.000045%)	
TATACACATCTAGATGTGTATAAGAGACAGCACACACACA	53 (0.000169%)	
TATACACATCTAGATGTGTATAAGAGACAGGTGTGTGTGT	47 (0.000150%)	
TATCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	16280 (0.051927%)	
TCACACACACACACACACACACACACACACACACACAC	4154 (0.013250%)	
TCCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	36679 (0.116991%)	
TCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTC	3442 (0.010979%)	
TCTCTTATACACATCTAGATGTGTATAAGAGACAGCACAC	50 (0.000159%)	
TCTCTTATACACATCTAGATGTGTATAAGAGACAGGTGTG	46 (0.000147%)	
TCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGA	2648 (0.008446%)	
TCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGC	5226 (0.016669%)	
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TCTTATACACATCTAGATGTGTATAAGAGACAGCACACAC	23 (0.000073%)	
TCTTATACACATCTAGATGTGTATAAGAGACAGGTGTGTG	26 (0.000083%)	
TGAGATGTGTATAAGAGACA	121328 (0.193494%)	
TGCCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	22560 (0.071957%)	
TGCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	27592 (0.088007%)	
TGTATAAGAGACAGGTGTGTGTGTGTGTGTGTGTGTGT	24 (0.000077%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCAC	171 (0.000545%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCAT	99 (0.000316%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCCC	66 (0.000211%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCCT	69 (0.000220%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCTC	53 (0.000169%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCTG	56 (0.000179%)	



After filtering: read2: base contents

Value of each position will be shown on mouse over.





After filtering: read2: KMER counting

Darker background means larger counts. The count will be shown on mouse over.





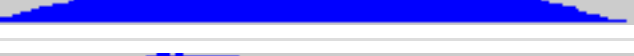
























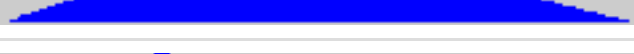




	AA	AT	AC	AG	TA	TT	TC	TG	CA	CT	CC	CG	GA	GT	GC	GG
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AAT	AATAA	AATAT	AATAC	AATAG	AATTA	AATTT	AATTC	AATTG	AATCA	AATCT	AATCC	AATCG	AATGA	AATGT	AATGC	AATGG
AAC	AACAA	AACAT	AACAC	AACAG	AACTA	AAC TT	AAC TC	AAC TG	AACCA	AACCT	AACCC	AACCG	AACGA	AACGT	AACGC	AACGG
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ATT	ATTAA	ATTAT	ATTAC	ATTAG	ATTTA	ATTTT	ATTT C	ATTT G	ATTCA	ATTCT	ATTCC	ATT CG	ATTGA	ATTGT	ATTGC	ATTGG
ATC	ATCAA	ATCAT	ATCAC	ATCAG	ATCTA	ATCTT	ATCTC	ATCTG	ATCCA	ATCCT	ATCCC	ATCCG	ATCGA	ATCGT	ATCGC	ATCGG
ATG	ATGAA	ATGAT	ATGAC	ATGAG	ATGTA	ATGTT	ATGTC	ATGTG	ATGCA	ATGCT	ATGCC	ATGCG	ATGGA	ATGGT	ATGGC	ATGGG
ACA	ACAAA	ACAAT	ACAAC	ACAAG	ACATA	ACATT	ACATC	ACATG	ACACA	ACACT	ACACC	ACACG	ACAGA	ACAGT	ACAGC	ACAGG
ACT	ACTAA	ACTAT	ACTAC	ACTAG	ACTTA	ACTTT	ACTTC	ACTTG	ACTCA	ACTCT	ACTCC	ACTCG	ACTGA	ACTGT	ACTGC	ACTGG
ACC	ACCAA	ACCAT	ACCAC	ACCAG	ACCTA	ACCTT	ACCTC	ACCTG	ACCCA	ACCCT	ACCCC	ACCCG	ACCGA	ACCGT	ACCGC	ACCGG
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TAT	TATAA	TATAT	TATAC	TATAG	TATTA	TATTT	TATTC	TATTG	TATCA	TATCT	TATCC	TATCG	TATGA	TATGT	TATGC	TATGG
TAC	TACAA	TACAT	TACAC	TACAG	TACTA	TACTT	TACTC	TACTG	TACCA	TACCT	TACCC	TACCG	TACGA	TACGT	TACGC	TACGG
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TGC	TGCAA	TGCAT	TGCAC	TGCAG	TGCTA	TGCTT	TGCTC	TGCTG	TGCCA	TGCCT	TGCCC	TGCCG	TGCGA	TGCGT	TGCGC	TGCGG
TGG	TGAAA	TGGAT	TGGAC	TGGAG	TGGTA	TGGTT	TGGTC	TGGTG	TGGCA	TGGCT	TGGCC	TGGCG	TGGGA	TGGGT	TGGGC	TGGGG
CAA	CAAAA	CAAAT	CAAAC	CAAAG	CAATA	CAATT	CAATC	CAATG	CAACA	CAACT	CAACC	CAACG	CAAGA	CAAGT	CAAGC	CAAGG
CAT	CATAA	CATAT	CATAC	CATAG	CATTA	CATTT	CATTC	CATTG	CATCA	CATCT	CATCC	CATCG	CATGA	CATGT	CATGC	CATGG
CAC	CACAA	CACAT	CACAC	CACAG	CACTA	CACTT	CACTC	CACTG	CACCA	CACCT	CACCC	CACCG	CACGA	CACGT	CACGC	CACGG
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CTA	CTAAA	CTAAT	CTAAC	CTAAG	CTATA	CTATT	CTATC	CTATG	CTACA	CTACT	CTACC	CTACG	CTAGA	CTAGT	CTAGC	CTAGG
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CTC	CTCAA	CTCAT	CTCAC	CTCAG	CTCTA	CTCTT	CTCTC	CTCTG	CTCCA	CTCCT	CTCCC	CTCCG	CTCGA	CTCGT	CTCGC	CTCGG
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CCA	CCAAA	CCAAT	CCAAC	CCAAG	CCATA	CCATT	CCATC	CCATG	CCACA	CCACT	CCACC	CCACG	CCAGA	CCAGT	CCAGC	CCAGG
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CGT	CGTAA	CGTAT	CGTAC	CGTAG	CGTTA	CGTTT	CGTTC	CGTTG	CGTCA	CGTCT	CGTCC	CGTCG	CGTGA	CGTGT	CGTGC	CGTGG
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GTT	GTTAA	GTTAT	GTTAC	GTTAG	GTTTA	GTTTT	GTTTC	GTTTG	GTTCA	GTTCT	GTTCC	GTT CG	GTTGA	GTTGT	GTTGC	GTTGG
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GGA	GGA AA	GGAAT	GGAAC	GGAAG	GGATA	GGATT	GGATC	GGATG	GGACA	GGACT	GGACC	GGACG	GGAGA	GGAGT	GGAGC	GGAGG
GGT	GGTAA	GGTAT	GGTAC	GGTAG	GGTTA	GGTTT	GGTTC	GGTTG	GGTCA	GGTCT	GGTCC	GGTCG	GGTGA	GGTGT	GGTGC	GGTGG
GGC	GGCAA	GGCAT	GGCAC	GGCAG	GGCTA	GGCTT	GGCTC	GGCTG	GGCCA	GGCCT	GGCCC	GGCCG	GGCGA	GGCGT	GGCGC	GGCGG
GGG	GGGAA	GGGAT	GGGAC	GGGAG	GGGTA	GGGTT	GGGTC	GGGTG	GGGCA	GGGCT	GGGCC	GGGCG	GGGGA	GGGGT	GGGGC	GGGGG


































After filtering: read2: overrepresented sequences

Sampling rate: 1 / 20















overrepresented sequence	count (% of bases)	distribution: cycle 1 ~ cycle 151
AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	35 (0.000112%)	
AACCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	22746 (0.072705%)	
AAGATGTGTATAAGAGACAG	302696 (0.483769%)	

[illegible]

		
CTCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	17749 (0.056733%)	
CTCTTATACACATCTAGATGTGTATAAGAGACAGCACACA	48 (0.000153%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGAA	1175 (0.003756%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGAC	3094 (0.009890%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGAG	1509 (0.004823%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGAT	3592 (0.011481%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCA	416 (0.001330%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCC	220 (0.000703%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCT	233 (0.000745%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGA	243 (0.000777%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGC	296 (0.000946%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGG	268 (0.000857%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGT	536 (0.001713%)	
CTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGTC	1591 (0.005085%)	
CTGTCTCTTATACACATCTC	314903 (0.503278%)	
CTGTCTCTTATACACATCTG	261585 (0.418065%)	
CTGTCTCTTATACACATCTT	277770 (0.443932%)	
CTTATACACATCTAGATGTGTATAAGAGACAGCACACACA	36 (0.000115%)	
GACCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	22614 (0.072283%)	
GA	5593 (0.017877%)	
GAGATGTGTATAAGAGACAG	110034 (0.175856%)	
GATGTGTATAAGAGACAGCACACACACACACACACACACA	121 (0.000387%)	
GCACACACACACACACACACACACACACACACACACAC	14317 (0.045763%)	
GCCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	45039 (0.143963%)	
GCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGA	2195 (0.007016%)	
GCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGC	4447 (0.014214%)	
GCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGG	6627 (0.021183%)	
GCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGT	1131 (0.003615%)	
GGCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	16756 (0.053559%)	
GGGCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	20301 (0.064890%)	
GG	8 (0.000026%)	
GTCTCTTATACACATCTAGATGTGTATAAGAGACAGCACA	73 (0.000233%)	
GTCTCTTATACACATCTAGATGTGTATAAGAGACAGGTGT	79 (0.000253%)	

		
GTCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	34708 (0.110941%)	
GTGCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	13003 (0.041563%)	
GTGTGCTGTCTCTTATACACATCTAGATGTGTATAAGAGA	1904 (0.006086%)	
GTGTGTGCTGTCTCTTATACACATCTAGATGTGTATAAGA	1542 (0.004929%)	
GTGTGTGTGCTGTCTCTTATACACATCTAGATGTGTATAA	2500 (0.007991%)	
GTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGA	767 (0.002452%)	
GT	21880 (0.069937%)	
TACCTGTCTCTTATACACATCTAGATGTGTATAAGAGACA	14342 (0.045843%)	
TATACACATCTAGATGTGTATAAGAGACAGCACACACACA	58 (0.000185%)	
TCACACACACACACACACACACACACACACACACACAC	11703 (0.037407%)	
TCCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	27686 (0.088495%)	
TCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTC	6206 (0.019837%)	
TCTCTTATACACATCTAGATGTGTATAAGAGACAGCACAC	52 (0.000166%)	
TCTCTTATACACATCTAGATGTGTATAAGAGACAGGTGTG	19 (0.000061%)	
TCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGA	156 (0.000499%)	
TCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGC	304 (0.000972%)	
TCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAGG	523 (0.001672%)	
TCTTATACACATCTAGATGTGTATAAGAGACAGCACACAC	31 (0.000099%)	
TGAGATGTGTATAAGAGACA	124681 (0.199265%)	
TGCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	24362 (0.077871%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCAC	230 (0.000735%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCAT	95 (0.000304%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCCC	95 (0.000304%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCCT	80 (0.000256%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGCTC	112 (0.000358%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGAG	84 (0.000268%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGCA	68 (0.000217%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGGA	68 (0.000217%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGGC	79 (0.000253%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGGG	90 (0.000288%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGGT	94 (0.000300%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGTA	131 (0.000419%)	



TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGTC	108 (0.000345%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGTG	257 (0.000821%)	
TGTCTCTTATACACATCTAGATGTGTATAAGAGACAGGTT	101 (0.000323%)	
TGTCTCTTATACACATCTCA	738 (0.001179%)	
TGTCTCTTATACACATCTCT	631 (0.001008%)	
TGTCTCTTATACACATCTGG	533 (0.000852%)	
TGTCTCTTATACACATCTGT	551 (0.000881%)	
TGTGCTGTCTCTTATACACATCTAGATGTGTATAAGAGAC	8808 (0.028154%)	
TGTGCTGTCTCTTATACACATCTAGATGTGTATAAGAG	3193 (0.010206%)	
TGTGCTGTCTCTTATACACATCTAGATGTGTATAAG	1418 (0.004532%)	
TGTGCTGTCTCTTATACACATCTAGATGTGTATA	18512 (0.059172%)	
TGTGCTGTCTGTGTGTGTGTGTGTGTGTGTGTGTGTG	29851 (0.095416%)	
TTATACACATCTAGATGTGTATAAGAGACAGCACACACAC	43 (0.000137%)	
TTCTGTCTCTTATACACATCTAGATGTGTATAAGAGACAG	14159 (0.045258%)	

```
fastp -i 10-12kb_S17_L002_R1_001.fastq.gz -I 10-12kb_S17_L002_R2_001.fastq.gz -o MP10k_F.trimmed.fq.gz -O MP10k_R.trimmed.fq.gz
-n 5 -q 30 -u 30 --length_required=100 --low_complexity_filter --complexity_threshold=20 --cut_by_quality3 --cut_by_quality5 --
cut_window_size=4 --cut_mean_quality=30 --trim_poly_g --poly_g_min_len=10 --overrepresentation_analysis --json=MP10k.json --
html=MP10k.html --report_title=MP10k --thread=8

fastp 0.19.4, at 2018-10-07 00:09:26
```