

fastp report

Summary

General

fastp version:	0.20.0 (https://github.com/OpenGene/fastp)
sequencing:	paired end (150 cycles + 150 cycles)
mean length before filtering:	148bp, 147bp
mean length after filtering:	147bp, 147bp
duplication rate:	8.521677%
Insert size peak:	239

Before filtering

total reads:	35.108356 M
total bases:	5.193082 G
Q20 bases:	5.120669 G (98.605586%)
Q30 bases:	4.969532 G (95.695227%)
GC content:	52.645179%

After filtering

total reads:	35.101238 M
total bases:	5.179270 G
Q20 bases:	5.107252 G (98.609487%)
Q30 bases:	4.956733 G (95.703304%)
GC content:	52.652993%

Filtering result

reads passed filters:	35.101238 M (99.979726%)
reads with low quality:	2 (0.000006%)
reads with too many N:	7.116000 K (0.020269%)
reads too short:	0 (0.000000%)

Adapters

Adapter or bad ligation of read1

The input has little adapter percentage (-0.238992%), probably it's trimmed before.

Sequence	Occurrences
A	45459
AG	46450
AGA	45893
AGAT	45005
AGATC	44722
AGATCG	44215
AGATCGG	44403
AGATCGGA	43185
AGATCGGAA	41891
AGATCGGAAG	41834
AGATCGGAAGA	41314
AGATCGGAAGAG	40792
AGATCGGAAGAGC	40660
AGATCGGAAGAGCA	40029
AGATCGGAAGAGCAC	38972
AGATCGGAAGAGCACA	38048
other adapter sequences	37438

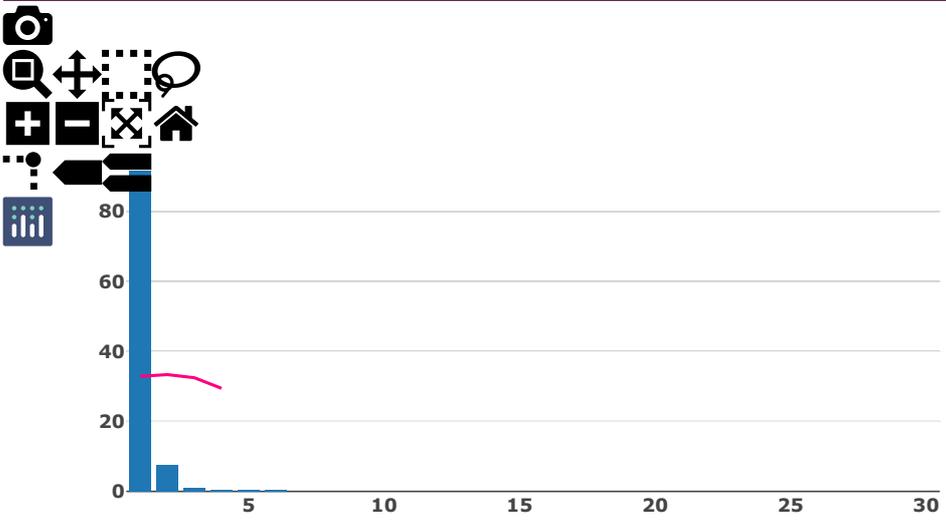
Adapter or bad ligation of read2

The input has little adapter percentage (-0.253582%), probably it's trimmed before.

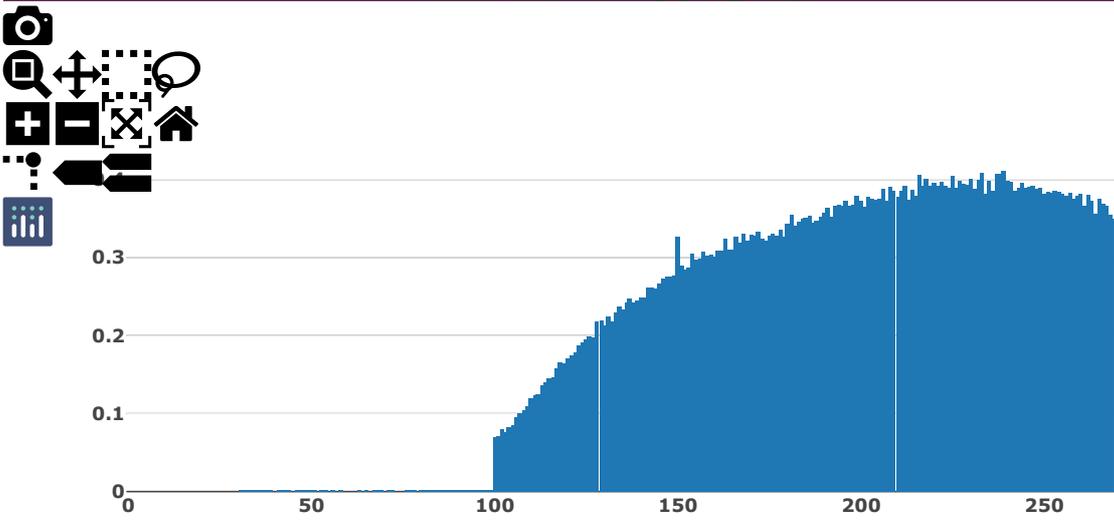
Sequence	Occurrences
A	48721

AG	47013
AGA	45799
AGAT	44734
AGATC	44756
AGATCG	44069
AGATCGG	43994
AGATCGGA	42800
AGATCGGAA	41912
AGATCGGAAG	41732
AGATCGGAAGA	40784
AGATCGGAAGAG	40502
AGATCGGAAGAGC	40283
AGATCGGAAGAGCG	41593
AGATCGGAAGAGCGT	36790
AGATCGGAAGAGCGTC	37370
other adapter sequences	69603

Duplication



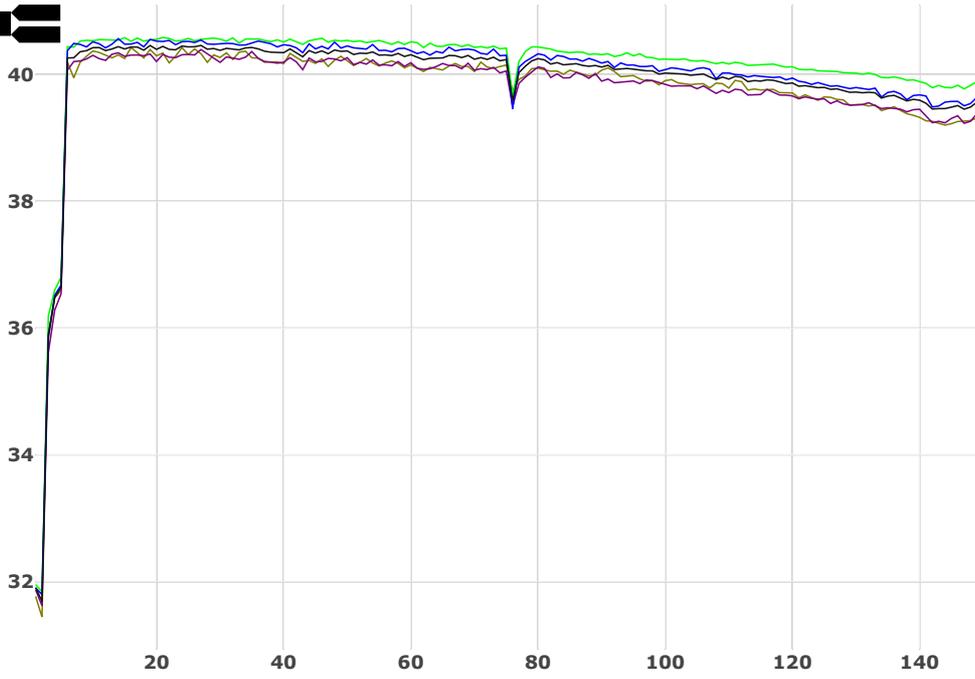
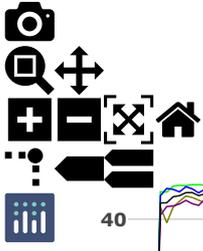
Insert size estimation



This estimation is based on paired-end overlap analysis, and there are 47.178487% reads found not overlapped. The nonoverlapped read pairs may have insert size <30 or >270, or contain too much sequencing errors to be detected as overlapped.

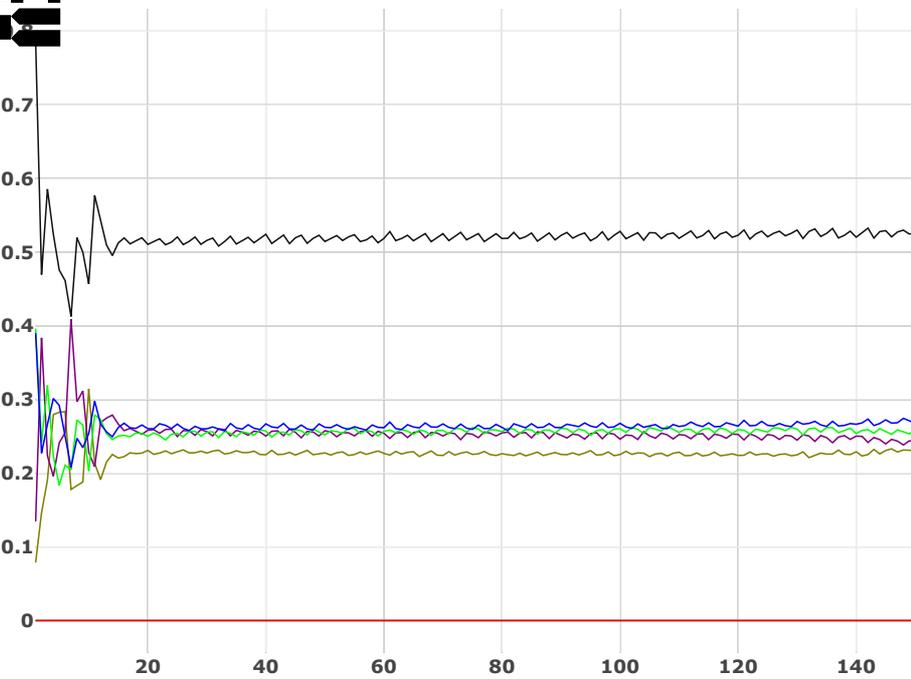
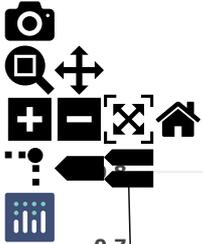
Before filtering

Before filtering: read1: quality
 Value of each position will be shown on mouse over.



Before filtering: read1: base contents

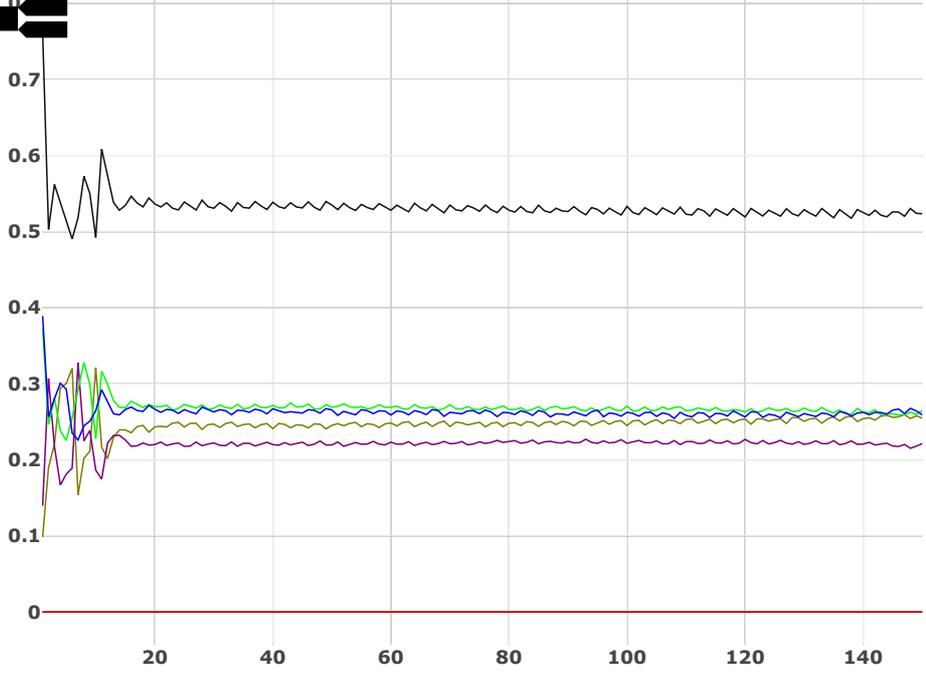
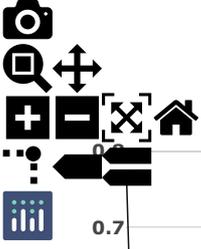
Value of each position will be shown on mouse over.



Before filtering: read1: KMER counting

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

	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	AAC TT	AAC TC	AAC TG	AAC CA	AAC CT	AAC CC	AAC CG	AAC GA	AAC GT	AAC GC	AAC GG
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	ATTT C	ATTT G	ATTT CA	ATTT CT	ATTT CC	ATTT CG	ATTT GA	ATTT GT	ATTT GC	ATTT GG
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	ATGGG	ATGGA	ATGGT	ATGGC	ATGGG
ACA	ACAAA	ACAAT	ACAAC	ACAAG	ACATA	ACATT	ACATC	ACATG	ACACA	ACACT	ACACC	ACACG	ACAGA	ACAGT	ACAGC	ACAGG



Before filtering: read2: KMER counting

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

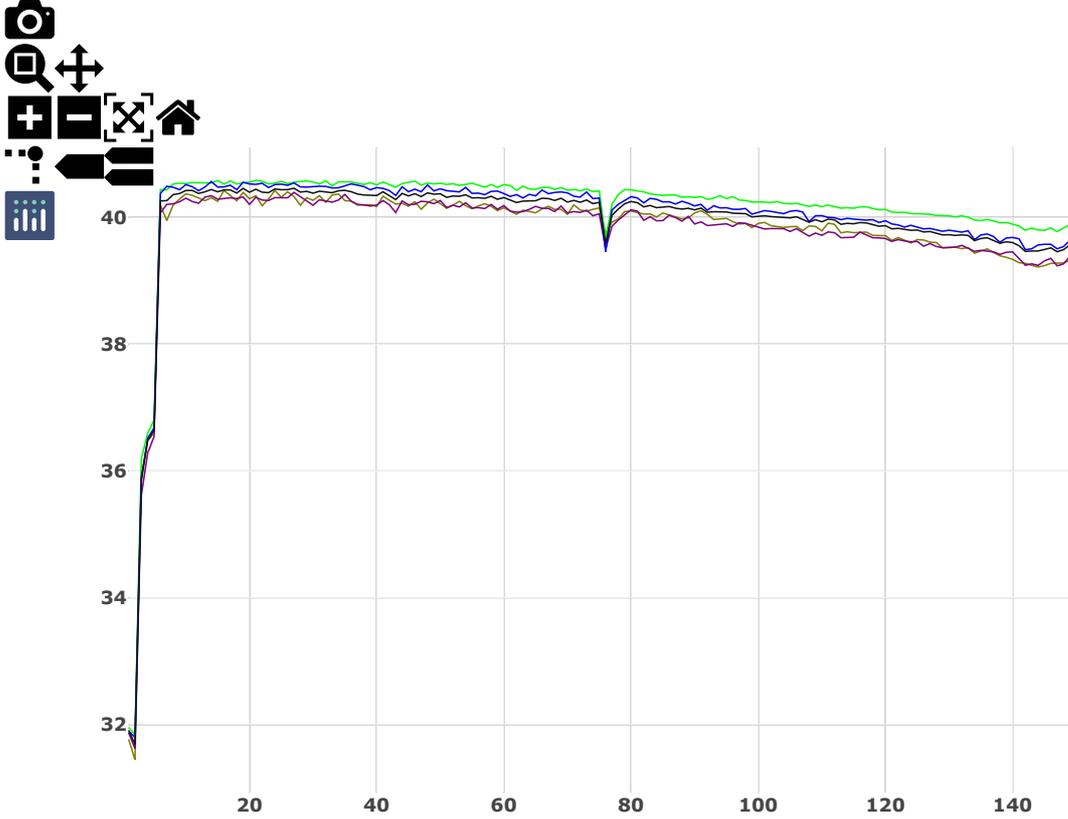
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AAC	AACA	AACAT	AACAC	AACAG	AACATA	AACATT	AACCT	AACCTG	AACCA	AACCT	AACCC	AACCG	AACGA	AACGT	AACGC	AACGG
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ATT	ATTA	ATTAT	ATTAC	ATTAG	ATTTA	ATTTT	ATTTT	ATTTG	ATTTA	ATTTT	ATTTT	ATTTG	ATTTA	ATTTG	ATTTG	ATTTG
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CCG	CCGA	CCGAT	CCGAC	CCGAG	CCGTA	CCGTT	CCGTC	CCGTG	CCGCA	CCGCT	CCGCC	CCGCG	CCGGA	CCGGT	CCGGC	CCGGG
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- A(24.67%)
- T(22.14%)
- C(26.87%)
- G(26.31%)
- N(0.001%)
- GC(53.18%)

After filtering

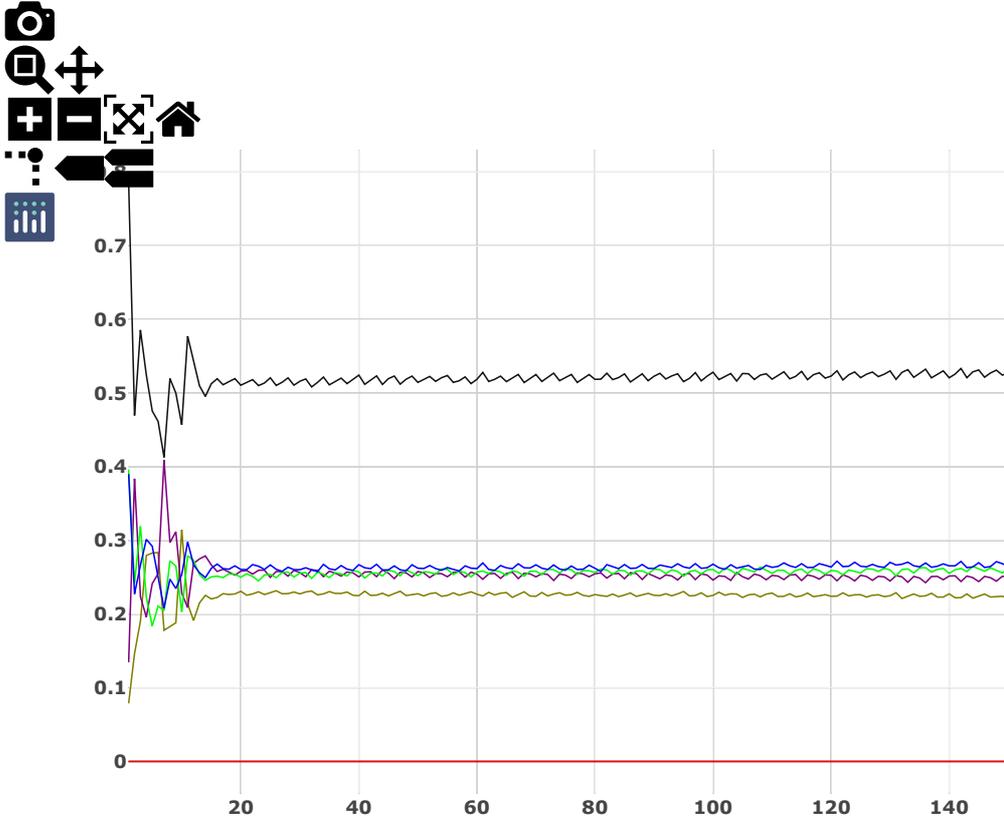
After filtering: read1: quality

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.

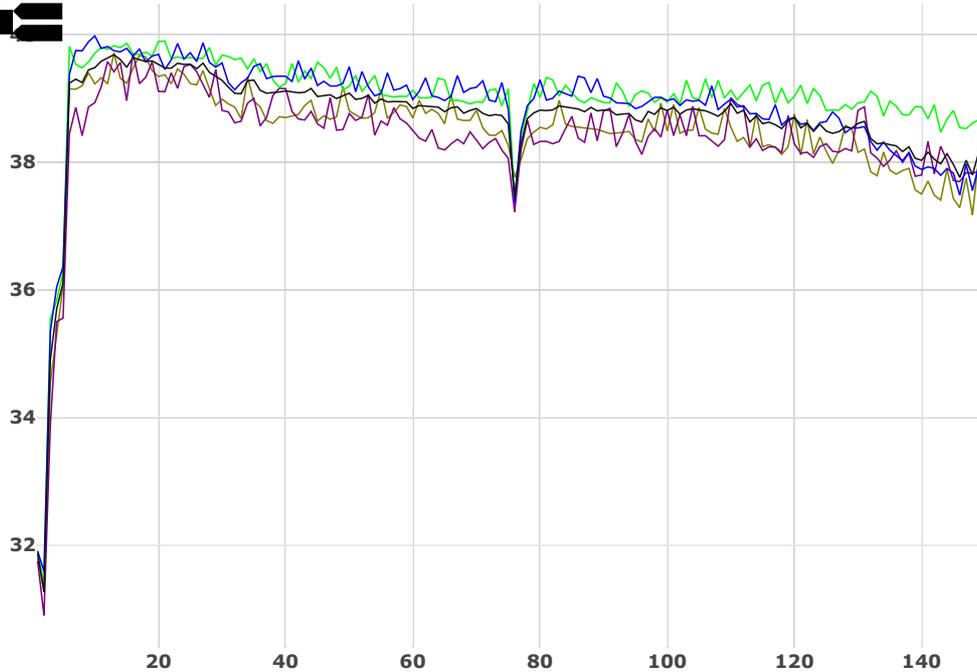
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AAT	AATAA	AATAT	AATAC	AATAG	AATTA	AATTT	AATTC	AATTG	AATCA	AATCT	AATCC	AATCG	AATGA	AATGT	AATGC	AATGG
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AAG	AAGAA	AAGAT	AAGAC	AAGAG	AAGTA	AAGTT	AAGTC	AAGTG	AAGCA	AAGCT	AAGCC	AAGCG	AAGGA	AAGGT	AAGGC	AAGGG
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ATC	ATCAA	ATCAT	ATCAC	ATCAG	ATCTA	ATCTT	ATCTC	ATCTG	ATCCA	ATCCT	ATCCC	ATCCG	ATCGA	ATCGT	ATCGC	ATCCG	ATCGG
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TTA	TTAAA	TTAAT	TTAAC	TTAAG	TTATA	TTATT	TTATC	TTATG	TTACA	TTACT	TTACC	TTACG	TTAGA	TTAGT	TTAGC	TTAGG	TTAGG
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TGG	TGAAA	TGAA	TGAA	TGAA	TGAA	TGAA	TGAA	TGAA	TGAA	TGAA	TGAA	TGAA	TGAA	TGAA	TGAA	TGAA	TGAA
CAA	CAAAA	CAAA	CAAAC	CAAAG	CAATA	CAATT	CAATC	CAATG	CAACA	CAACT	CAACC	CAACG	CAAGA	CAAGT	CAAGC	CAAGG	CAAGG
CAT	CATAA	CATAT	CATAC	CATAG	CATTA	CATTT	CATTC	CATTG	CATCA	CATCT	CATCC	CATCG	CATGA	CATGT	CATGC	CATGG	CATGG
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CAG	CAGAA	CAGAT	CAGAC	CAGAG	CAGTA	CAGTT	CAGTC	CAGTG	CAGCA	CAGCT	CAGCC	CAGCG	CAGGA	CAGGT	CAGGC	CAGGG	CAGGG
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CTT	CTAAA	CTAAT	CTAAC	CTAAG	CTATA	CTATT	CTATC	CTATG	CTACA	CTACT	CTACC	CTACG	CTAGA	CTAGT	CTAGC	CTAGG	CTAGG
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CCT	CCTAA	CCTAT	CCTAC	CCTAG	CCTTA	CCTTT	CCTTC	CCTTG	CCTCA	CCTCT	CCTCC	CCTCG	CCTGA	CCTGT	CCTGC	CCTGG	CCTGG
CCC	CCCAA	CCCA	CCCA	CCCA	CCCA	CCCA	CCCA	CCCA	CCCA	CCCA	CCCA	CCCA	CCCA	CCCA	CCCA	CCCA	CCCA
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CGT	CGTAA	CGTAT	CGTAC	CGTAG	CGTTA	CGTTT	CGTTC	CGTTG	CGTCA	CGCTT	CGCCC	CGCCG	CGCGA	CGCGT	CGCCG	CGCCG	CGCCG
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GAA	GAAAA	GAAT	GAAC	GAAG	GAATA	GAATT	GAATC	GAATG	GAACA	GAACT	GAACC	GAACG	GAAGA	GAAGT	GAAGC	GAAGG	GAAGG
GAT	GATAA	GATAT	GATAC	GATAG	GATTA	GATTT	GATTC	GATTG	GATCA	GATCT	GATCC	GATCG	GATGA	GATGT	GATGC	GATGG	GATGG
GAC	GACAA	GACAT	GACAC	GACAG	GACTA	GACTT	GACTC	GACTG	GACCA	GACCT	GACCC	GACCG	GACGA	GACGT	GACGC	GACGG	GACGG
GAG	GAGAA	GAGAT	GAGAC	GAGAG	GAGTA	GAGTT	GAGTC	GAGTG	GAGCA	GAGCT	GAGCC	GAGCG	GAGGA	GAGGT	GAGGC	GAGGG	GAGGG
GTA	GTAAA	GTAA	GTAA	GTAA	GTAA	GTAA	GTAA	GTAA	GTAA	GTAA	GTAA	GTAA	GTAA	GTAA	GTAA	GTAA	GTAA
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GTC	GTCAA	GTCAT	GTCAC	GTCAG	GCTTA	GCTTT	GCTTC	GCTTG	GCTCA	GCTCT	GCTCC	GCTCG	GCTGA	GCTGT	GCTGC	GCTGG	GCTGG
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- A(22.50%)
- T(25.37%)
- C(25.68%)
- G(26.43%)
- N(0.000%)
- GC(52.11%)

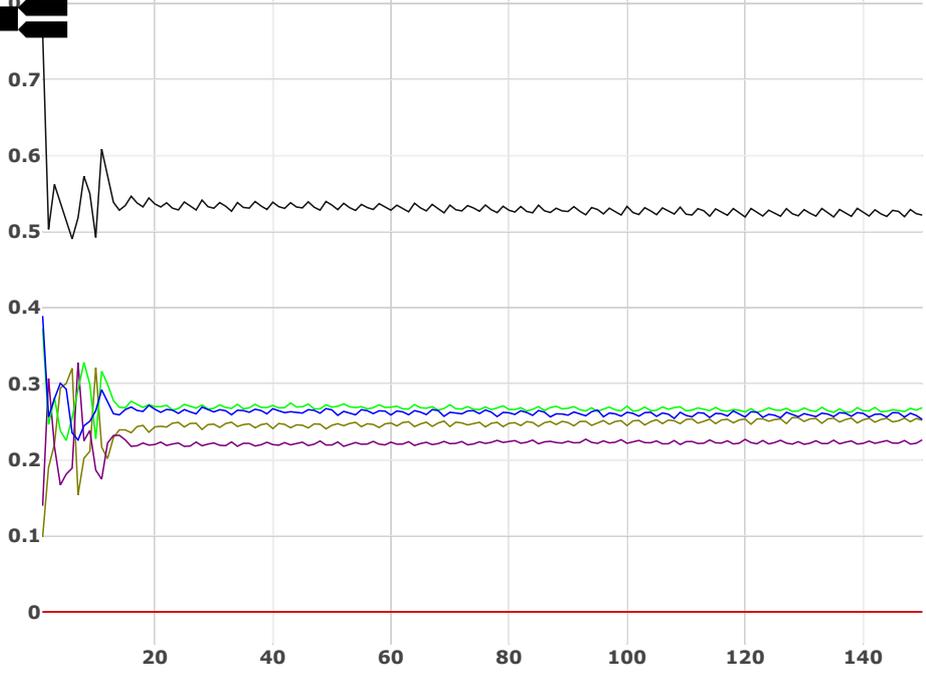
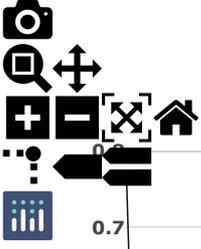
After filtering: read2: quality

Value of each position will be shown on mouse over.



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	
AAA	AAAA	AAAT	AAAC	AAAG	AAATA	AAATT	AAATC	AAATG	AAACA	AAACT	AAACC	AAACG	AAAGA	AAAGT	AAAGC	AAAGG
AAT	AATA	AATAT	AATAC	AATAG	AATTA	AATTT	AATTC	AATTG	AATCA	AATCT	AATCC	AATCG	AATGA	AATGT	AATGC	AATGG
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AAG	AAGA	AAGAT	AAGAC	AAGAG	AAGTA	AAGTT	AAGTC	AAGTG	AAGCA	AAGCT	AAGCC	AAGCG	AAGGA	AAGGT	AAGGC	AAGGG
ATA	ATAA	ATAAT	ATAAC	ATAAG	ATATA	ATATT	ATATC	ATATG	ATACA	ATACT	ATACC	ATACG	ATAGA	ATAGT	ATAGC	ATAGG
ATT	ATTA	ATTAT	ATTAC	ATTAG	ATTTA	ATTTT	ATTTT	ATTTG	ATTTA	ATTTT	ATTTCC	ATTTCC	ATTTGA	ATTTGT	ATTTGC	ATTTGG
ATC	ATCA	ATCAT	ATCAC	ATCAG	ATCTA	ATCTT	ATCTC	ATCTG	ATCCA	ATCCT	ATCCC	ATCCG	ATCGA	ATCGT	ATCGC	ATCGG
ATG	ATGA	ATGAT	ATGAC	ATGAG	ATGTA	ATGTT	ATGTC	ATGTG	ATGCA	ATGCT	ATGCC	ATGCG	ATGGA	ATGGT	ATGGC	ATGGG
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TAC	TACA	TACAT	TACAC	TACAG	TACTA	TACTT	TACTC	TACTG	TACCA	TACCT	TACCC	TACCG	TACGA	TACGT	TACGC	TACGG
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TTG	TTGA	TTGAT	TTGAC	TTGAG	TTGTA	TTGTT	TTGTC	TTGTG	TTGCA	TTGCT	TTGCC	TTGCG	TTGGA	TTGGT	TTGGC	TTGGG
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- A(24.63%)
- T(22.17%)
- C(26.90%)
- G(26.28%)
- N(0.000%)
- GC(53.19%)

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