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1. Summary de novo report

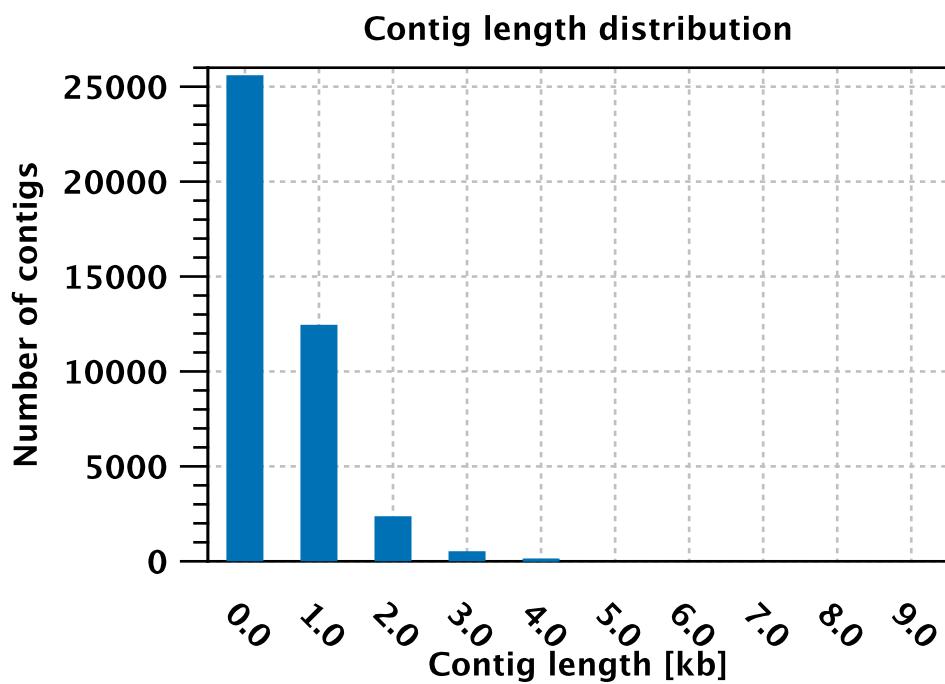
1.1 Nucleotide distribution

Nucleotide	Count	Frequency
Adenine (A)	7,512,962	29.9%
Cytosine (C)	5,073,725	20.2%
Guanine (G)	5,083,943	20.2%
Thymine (T)	7,466,906	29.7%
Any nucleotide (N)	156	0.0%

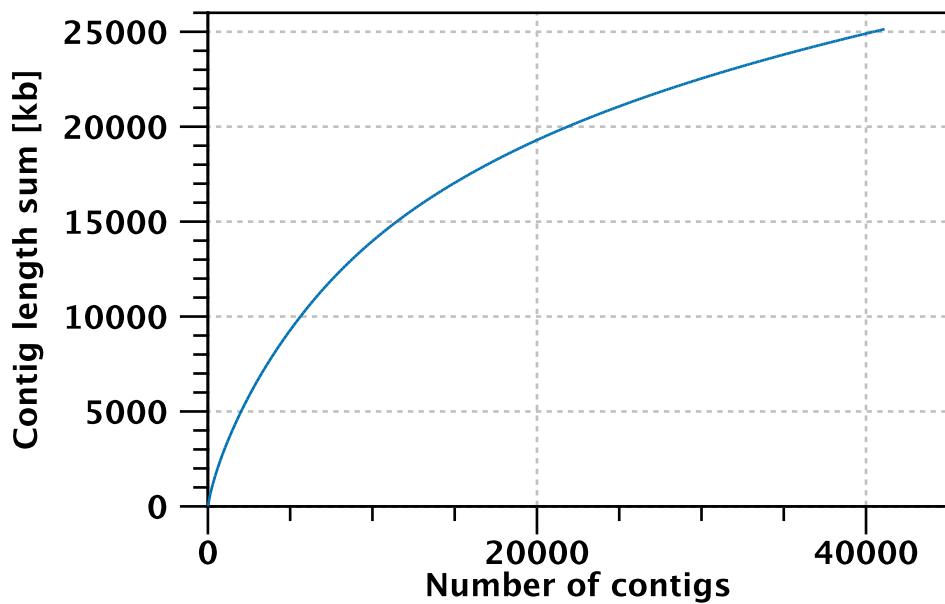
1.2 Contig measurements

N75	417
N50	863
N25	1,564
Minimum	185
Maximum	8,529
Average	611
Count	41,136

Total	25,137,692
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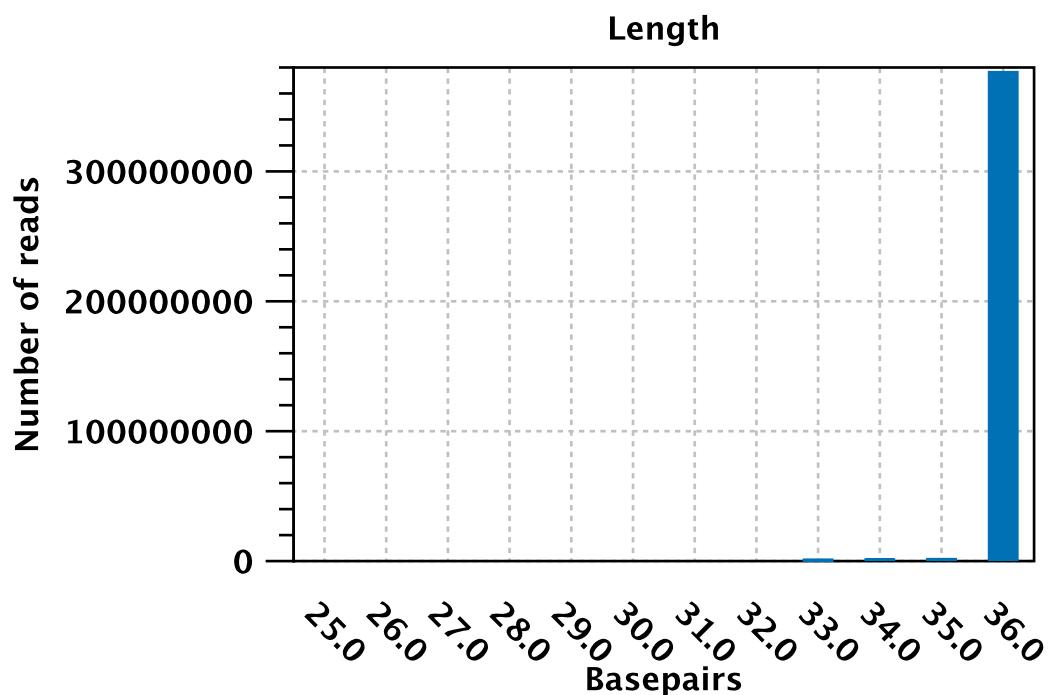
1.3 Accumulated contig lengths



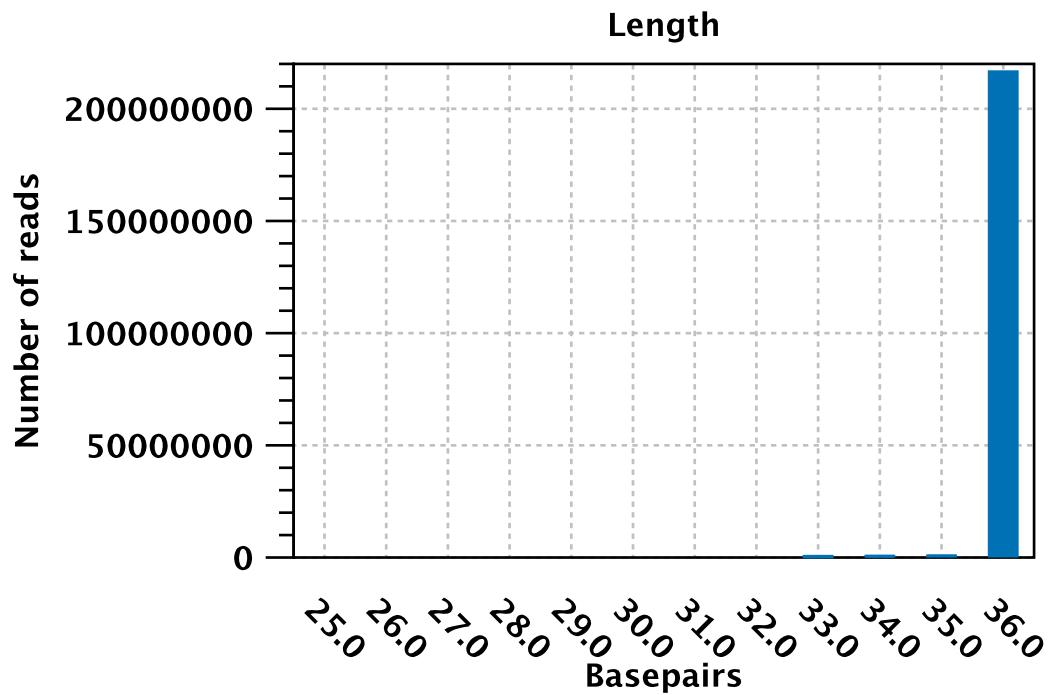
1.4 Summary statistics

	Count	Average length	Total bases
Reads	387,720,843	35.9	13,918,087,430
Matched	222,773,113	35.91	7,999,005,359
Not matched	164,947,730	35.88	5,919,082,071
Contigs	41,136	611	25,137,692

1.5 Distribution of read length



1.6 Distribution of matched read length



1.7 Distribution of non-matched read length

