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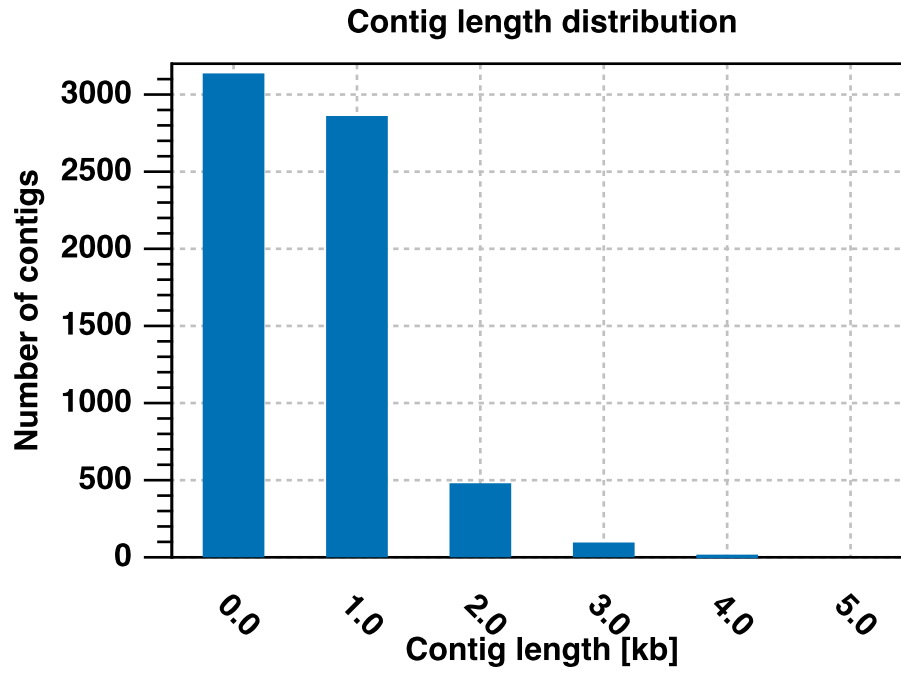
1. Summary de novo report

1.1 Nucleotide distribution

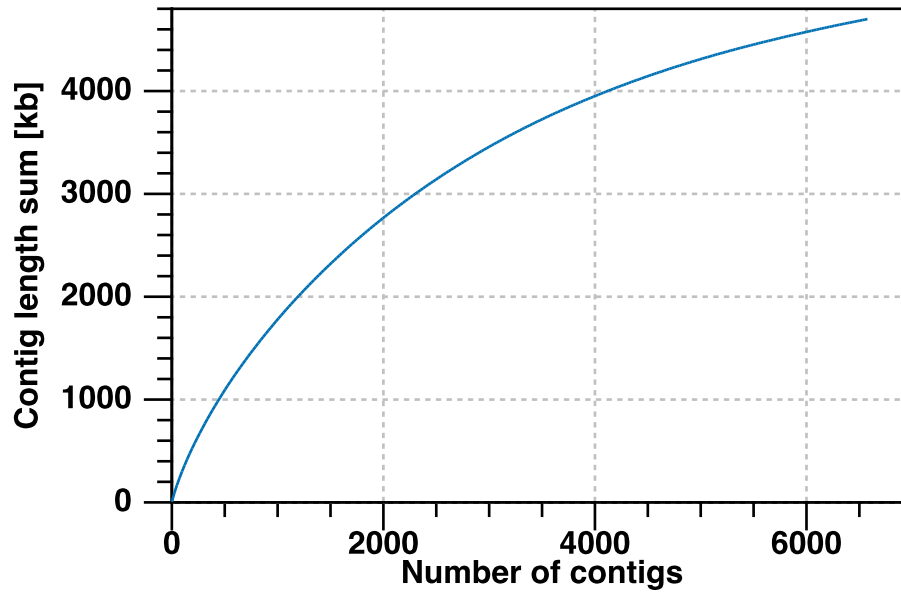
Nucleotide	Count	Frequency
Adenine (A)	1,259,943	26.8%
Cytosine (C)	1,087,941	23.1%
Guanine (G)	1,086,042	23.1%
Thymine (T)	1,268,074	27.0%

1.2 Contig measurements

N75	558
N50	967
N25	1,542
Minimum	198
Maximum	4,657
Average	714
Count	6,588



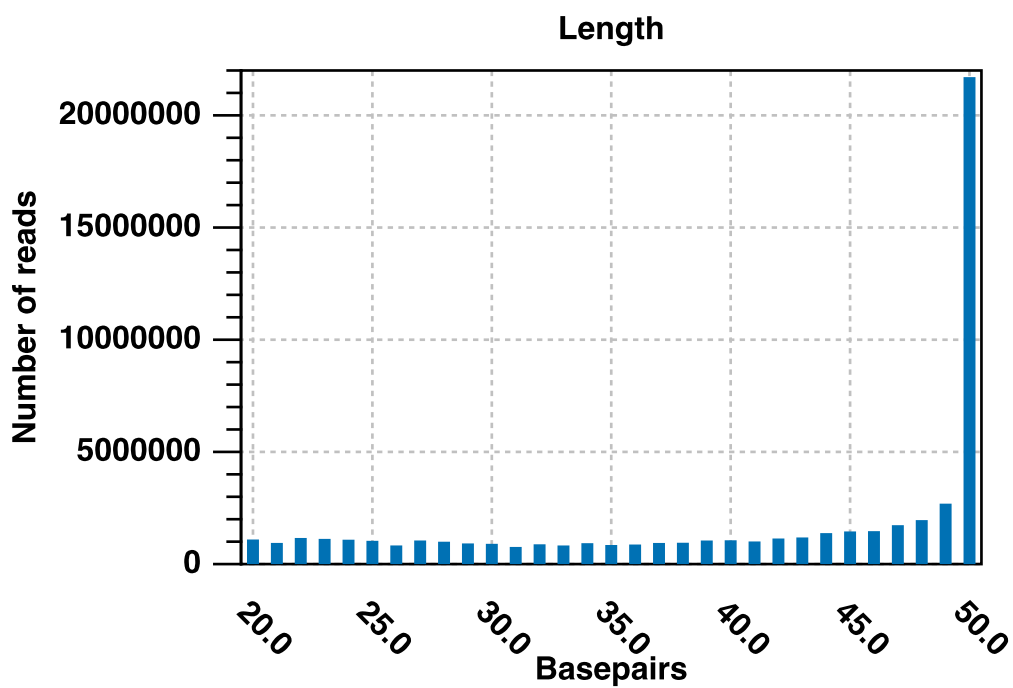
1.3 Accumulated contig lengths



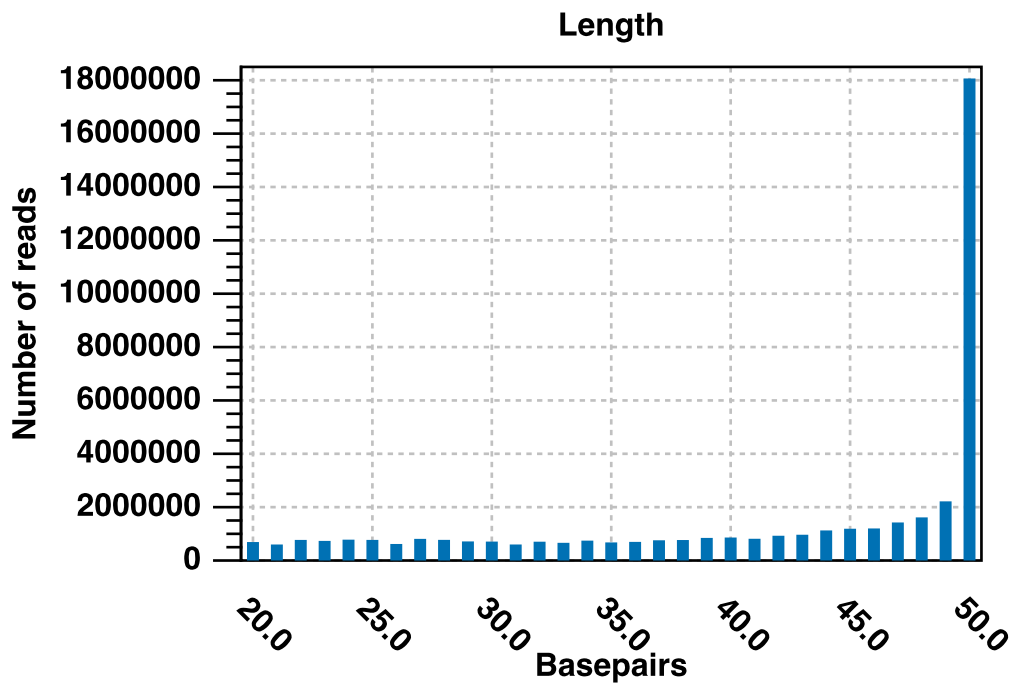
1.4 Summary statistics

	Count	Average length	Total bases
Reads	56,024,538	41.6	2,330,379,926
Matched	44,869,510	42.15	1,891,319,516
Not matched	11,155,028	39.36	439,060,410
Contigs	6,588	713	4,702,000

1.5 Distribution of read length



1.6 Distribution of matched read length



1.7 Distribution of non-matched read length

