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# 1. Summary mapping report

## 1.1 Summary statistics

	Count	Average length	Total bases
Reads	12,185,418	527.52	6,428,059,816
Matched	8,119,690	518.54	4,210,388,295
Not matched	4,065,728	545.45	2,217,671,521
Contigs	232,373	1,295	300,982,316

## 1.2 General algorithm parameters

Parameter	Value
Conflict resolution	Vote (A, C, G, T)
Non specific matches	ignore

## 1.3 Reads parameters

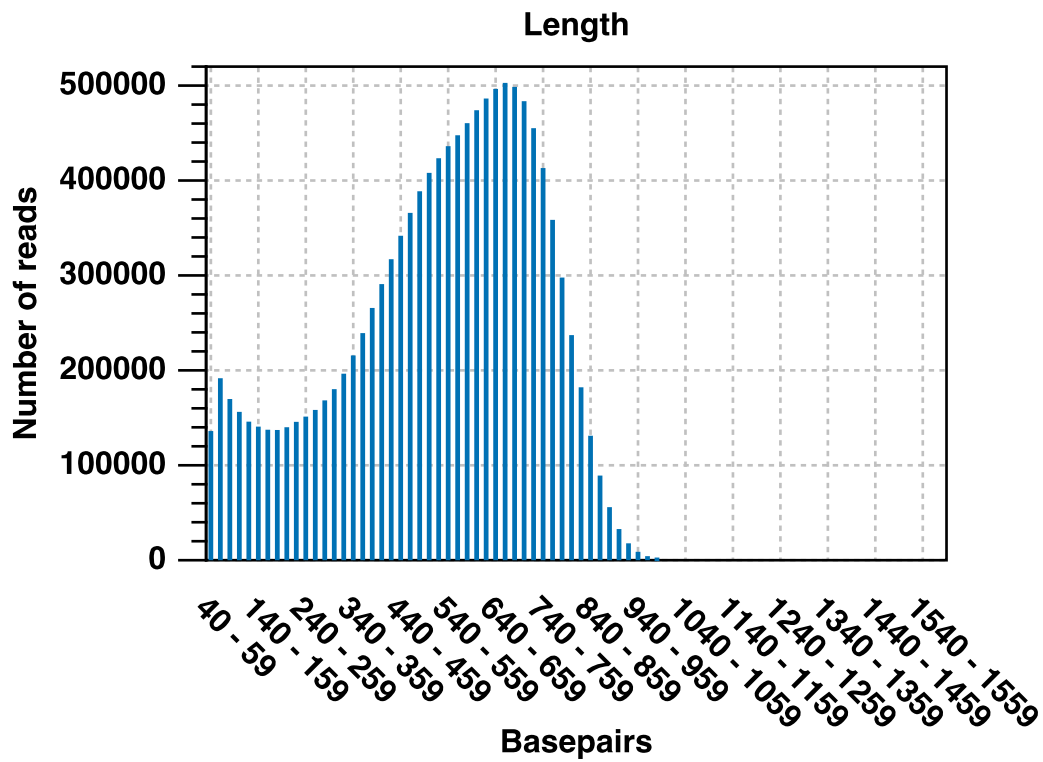
Reads	Length	Type	Parameters
G5UGEGY01 (single)	Long	Single	Similarity=0.95
G5UGEGY02 (single)	Long	Single	Similarity=0.95
G6IEU6N01 (single)	Long	Single	Similarity=0.95
G6IEU6N02 (single)	Long	Single	Similarity=0.95
G6P4JCY01 (single)	Long	Single	Similarity=0.95
G6P4JCY02 (single)	Long	Single	Similarity=0.95
G6IQQIE02 (single)	Long	Single	Similarity=0.95
G6IQQIE01 (single)	Long	Single	Similarity=0.95
G6MC6LA01 (single)	Long	Single	Similarity=0.95
G6MC6LA02 (single)	Long	Single	Similarity=0.95
G6NY9WP01 (single)	Long	Single	Similarity=0.95
G6NY9WP02 (single)	Long	Single	Similarity=0.95
G5T4LHD01 (single)	Long	Single	Similarity=0.95
G5T4LHD02 (single)	Long	Single	Similarity=0.95
G5YGYAB01 (single)	Long	Single	Similarity=0.95
G5YGYAB02 (single)	Long	Single	Similarity=0.95

Reads	Length	Type	Parameters
G5YGYJV01 (single)	Long	Single	Similarity=0.95
G5YGYJV02 (single)	Long	Single	Similarity=0.95
G5Z66BW01 (single)	Long	Single	Similarity=0.95
G5Z66BW02 (single)	Long	Single	Similarity=0.95

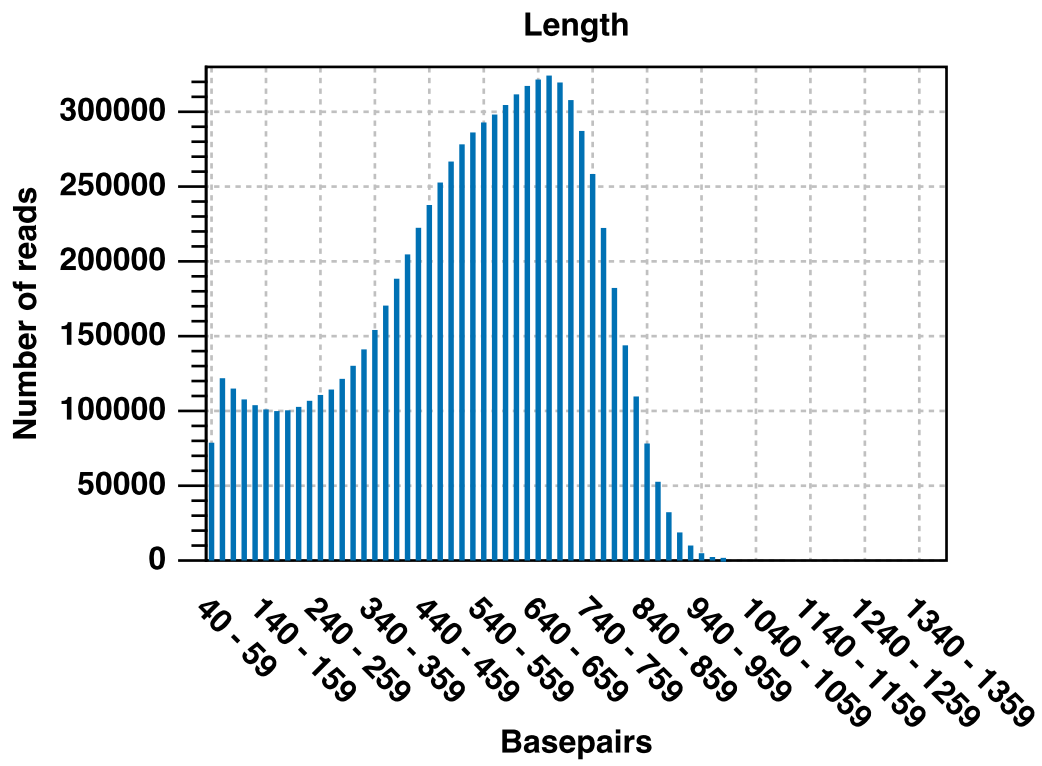
## 1.4 Quality measurement

N75	870
N50	1,604
N25	3,209

## 1.5 Distribution of read length



## 1.6 Distribution of matched read length



## 1.7 Distribution of non-matched read length

