

: C.Dson_15_02_317Ma2_index13 : 3 : AGTCAA (3_120719_BD158KACXX_1_nophix-sort-dup)

Reference organism	hg19	
Total	228,048	101bp paired
Aligned	206,040	(90.3%)
Pairs aligned	203,173	(89.1%)
Pair duplicates	300	(0.1%)
Insert size	182.2	+/- 58.4
On bait bases	17,077,067	(45.1%)
Near bait bases	7,679,192	(20.3%)
Off bait bases	13,081,387	(34.6%)
Mean bait coverage	0.3	
On target bases	17,093,054	(45.2%)
Mean target coverage	0x	
10x coverage targets	0.0%	
Zero coverage targets	75.6%	
Fold enrichment	27x	

Table 1: Summary of lane results

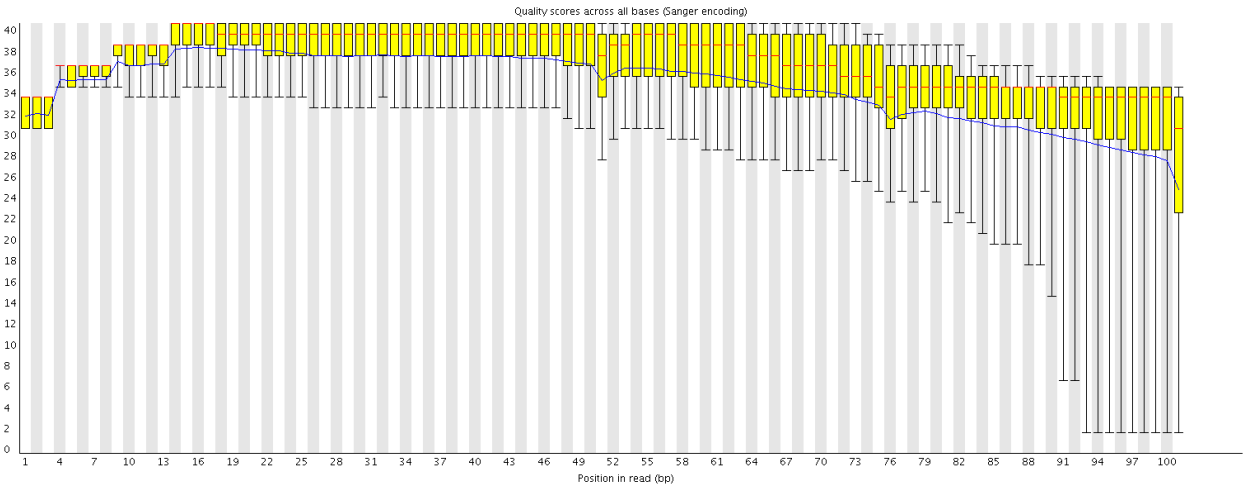


Figure 1:

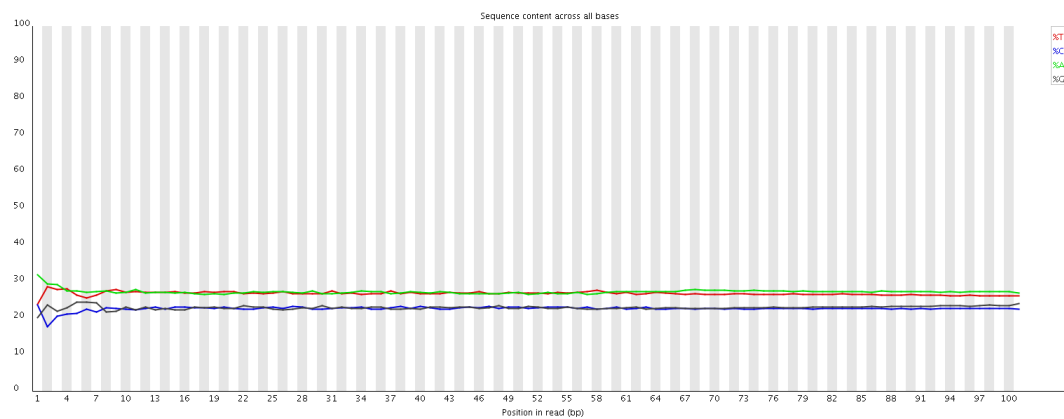


Figure 2:

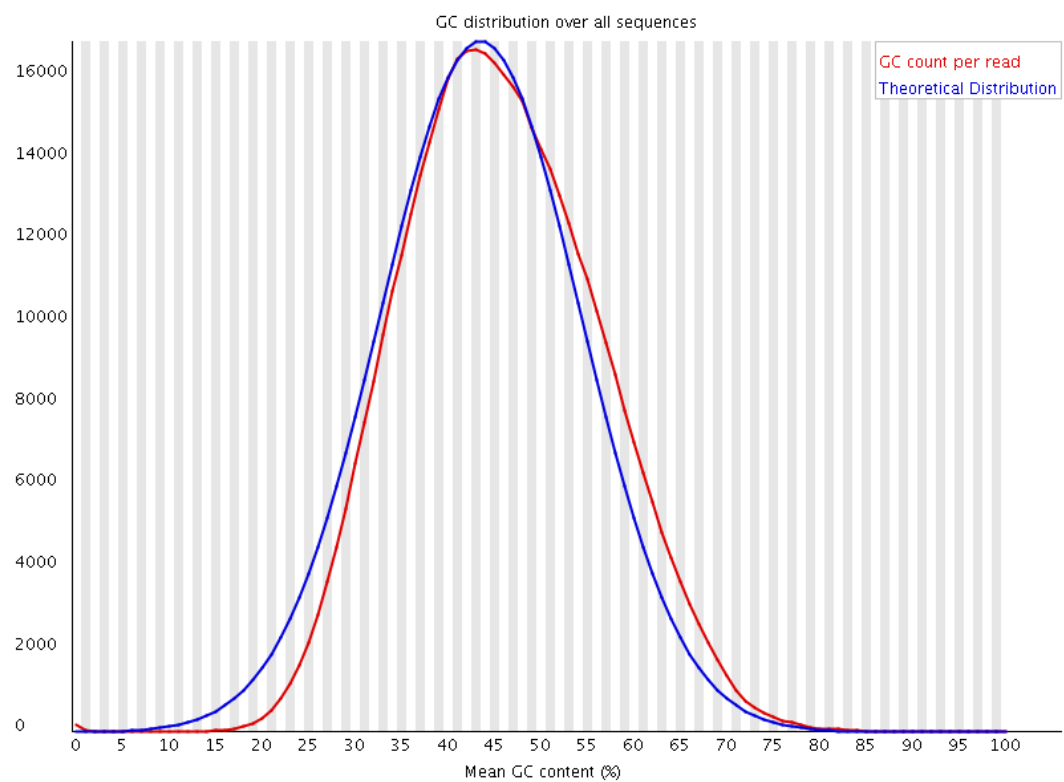


Figure 3:

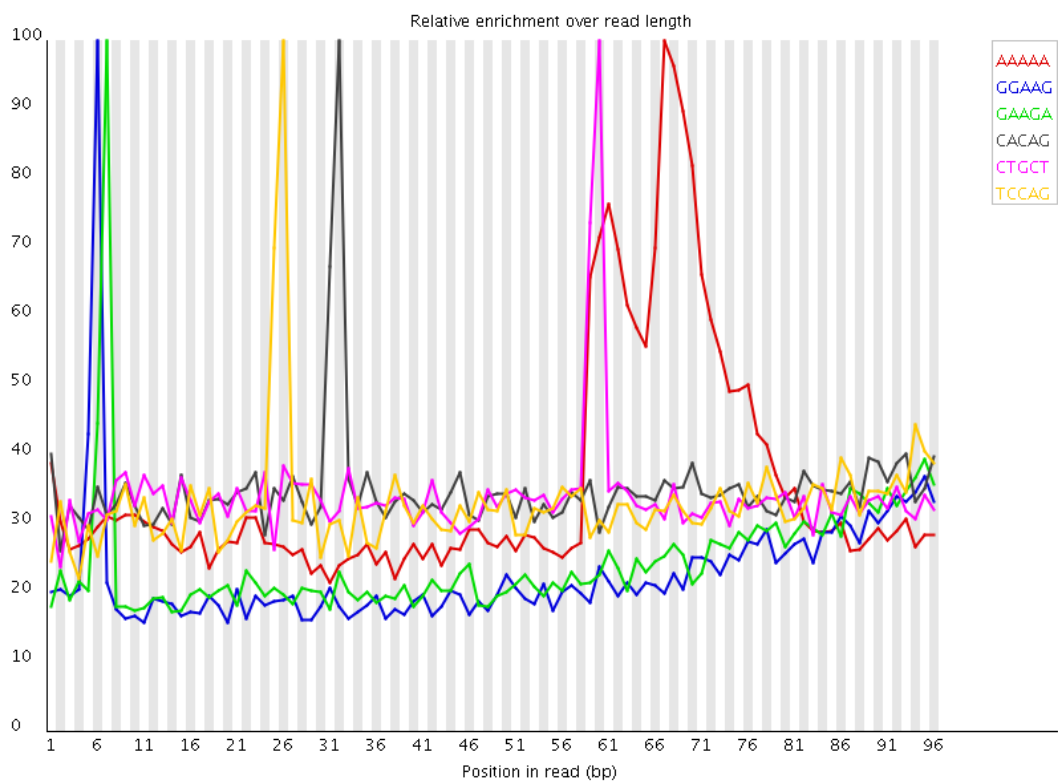


Figure 4:

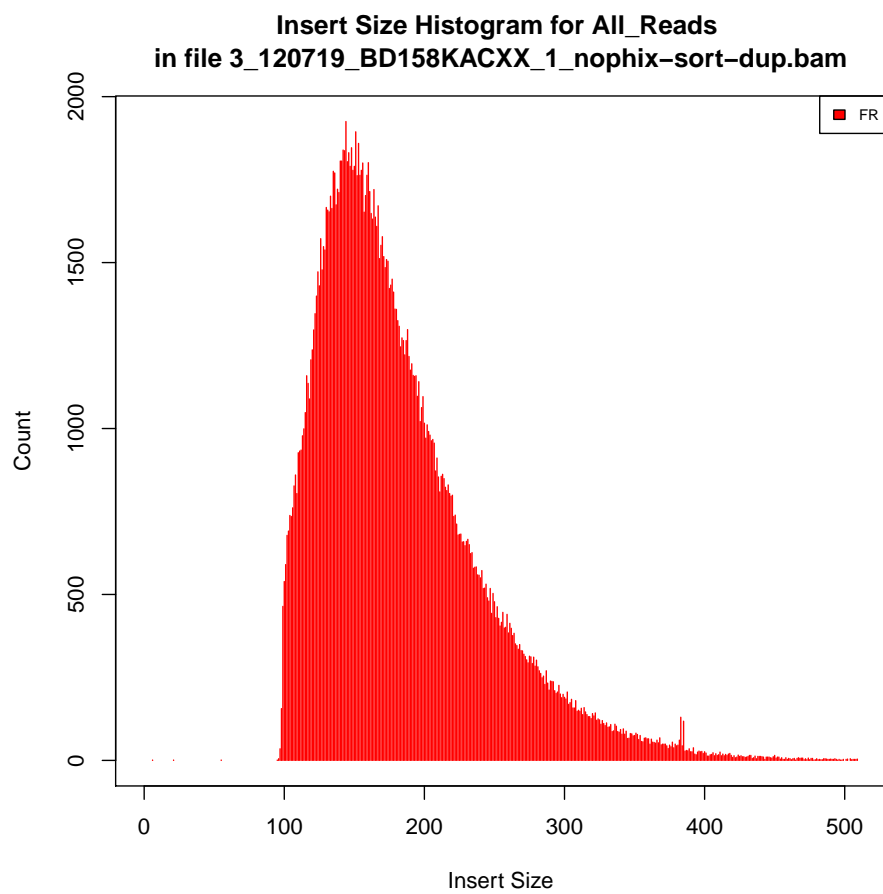


Figure 5: Distribution of paired end insert sizes