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✅ Basic Statistics

Measure	Value
Filename	LR13_HHH23DSXX_L3_2.fq
File type	Conventional base calls
Encoding	Sanger / Illumina 1.9
Total Sequences	8994262
Sequences flagged as poor quality	0
Sequence length	150
%GC	38

✅ Per base sequence quality

Quality scores across all bases (Sanger / Illumina 1.9 encoding)