

Overrepresented sequences

Sequence	Count	Percentage	Possible Source
GATCGGAAGAGCACACGTCTGAACTCCAGTCACGATCAGATCTCGTATGCC	27818	0.69545	TruSeq Adapter, Index 9 (100% over 51bp)
AGATCGGAAGAGCACACGTCTGAACTCCAGTCACGATCAGATCTCGTATGC	22437	0.560925	TruSeq Adapter, Index 9 (100% over 50bp)