

Before cutadapt

Overrepresented sequences

Sequence	Count	Percentage	Possible Source
GATCGGAAGAGCACACGTCTGAACTCCAGTCACTGACCAATCTCGTATGCC	28382	0.70955	TruSeq Adapter, Index 4 (100% over 51bp)
AGAGCACACGTCTGAACTCCAGTCACTGACCAATCTCGTATGCCGTCTTCT	8048	0.2012	TruSeq Adapter, Index 4 (100% over 51bp)
CTTGAATTAAGGCAACAGCAAATTCTAGGATCGTAAGTAGGATTAGAATAA	4984	0.12459999999999999	No Hit
AGATCGGAAGAGCACACGTCTGAACTCCAGTCACTGACCAATCTCGTATGC	4600	0.11499999999999999	TruSeq Adapter, Index 4 (100% over 50bp)
CTCACGCTTACCACATAGTCAACCCAAGCCCATGACCGCTGACAGGGGCCC	4492	0.11230000000000001	No Hit

After cutadapt

Overrepresented sequences

Sequence	Count	Percentage	Possible Source
AGAGCACACGTCTGAACTCCAGTCACTGACCAATCTCGTATGCCGTCTTCT	50406	0.1889822798462917	TruSeq Adapter, Index 4 (100% over 51bp)
CTTGAATTAAGGCAACAGCAAATTCTAGGATCGTAAGTAGGATTAGAATA	32410	0.1215116392853691	No Hit
CTCACGCTTACCACATAGTCAACCCAAGCCCATGACCGCTGACAGGGGCCC	28310	0.10613991077348962	No Hit