Overrepresented sequences (file.fastq.gz)

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Sequence	Percentage(%)
CGGGCAGATCGCAACTCCCAGGCATCCCGCAAAGT	8.297
AGGGCTTTCCGGTAAAAAACCAGAAAGCCTACTN	5.7
ACTCAGAGTTGTTCCTCTCCACGGAAATCTTTAGT	5.311
AGCCTACAGCACCCGGTATTCCCAGGCGGTCTCCC	4.844
AGCCTTGCCAAAGCAAGGCCTCAAAAAATTGGGTT	2.678
AGGGGCTTTCCGGTAAAAAGCCAGAAAGCCTACTN	2.193
CAGGGGAAAGCGCAGTCCCCCACTACCAC	1.975
AGCCTTGTCGGAACAAGGCCTCAAAAAATTAGCTT	1.674
GTATCAGATATTAAACTGATAAGAACAGATACTAC	1.043
TAGCTTAAGACTCAGAGTTGTTCCTCCACGGAA	1.001
ACAGCACCCGGTATTCCCAGGCGGTCTCCCATCCA	0.993
AAGGGCGCCGGGACCAACGATCACCAGCTGCCCAA	0.865
CTCGGATAGAGGACGTATCAGATATTAAACTGATA	0.752
ACGTATCAGATATTAAACTGATAAGAACAGATACT	0.745
CATTTAATATTGTCCTCGGATAGAGGACGTATC	0.645
GATAAGAACAGATACTACACTTGATCTTAGCCAAA	0.606
GAAGAAACCAGAGTATGATCTTAGCCAAA	0.6
GGGGCTTTCCGGTAAAAAACCAGAAAGCCTACTNN	0.589
ACCACTCAGACCGCGTTCTCTCCCTCTCACTCCCC	0.566
AAGGGCTTTCCGGTAAAAAACCAGAAAGCCTACT	0.562
GATCGTCGGACTGTAGAACTCTGAACGTGTAGATC	0.545
AATATTGTCCTCGGATAGAGGACGTATCAGATATT	0.444
TAAACTGATAAGAACAGATACTACACTTGATCTTN	0.443
CAGAGTTGTTCCTCTCCACGGAAATCTTTAGTAAA	0.42
TGGGCAGATCGCAACTCCCGCAAAGT	0.41
GGGGCTTTCCGGTAAAAACCAGAAAGCCTACTN	0.362
CGCCTACAGCACCCGGTATTCCCAGGCGGTCTCCC	0.333
GGCCTACAGCACCCGGTATTCCCAGGCGGTCTCCC	0.325
AGCCTTGCCTAGGCAAGACTTCAAAAAATTTAACC	0.322
TGCACCGTTCCTGGAGGTACTGCAATACCAGGTCG	0.297
CGGGGCTTTCCGGTAAAAAACCAGAAAGCCTACTN	0.277
GCTCAGAGTTGTTCCTCTCCACGGAAATCTTTAGT	0.271
CGGATAGAGGACGTATCAGATATTAAACTGATANN	0.269
GGGGCTTTCCGGTAAAAAGCCAGAAAGCCTACTNN	0.256
GAGGACGTATCAGATATTAAACTGATAAGAACNNN	0.248
TGTTCCTCTCCACGGAAATCTTTAGTAAAAGGCGA	0.244
CTGATAAGAACAGATACTACACTTGATCTTAGCCA	0.227
AATTGTCCTCGGATAGAGGACGTATCAGATATTAA	0.227
GGATCGTCGGACTGTAGAACTCTGAACGTGTNNNN	0.214
GACGTATCAGATATTAAACTGATAAGAACAGATAC	0.208
GCGCGAACGCAGTCCCCCACTACCACAAATTATGC	0.192
ATCCATTTAATATTGTCCTCGGATAGAGGACGT	0.191
AAGGGCTTTCCGGTAAAAAGCCAGAAAGCCTACT	0.191
TGCCTACAGCACCCGGTATTCCCAGGCGGTCTCCC	0.188
TCTCTCCACGGAAATCTTTAGTAAAAGGCGAANNN	0.188
CCTCAGAGTTGTTCCTCCACGGAAATCTTTAGT	0.176
AAAATTGTCCTCGGATAGAGGACGTATCAGATATT	0.168
GTTTCTCTCCACGGAAATCTTTAGTAAAAGGCGAA	0.166
CGTTCCTCCACGGAAATCTTTAGTAAAAGGCGA	0.159
TGGGGCTTTCCGGTAAAAACCAGAAAGCCTACTN	0.154
AGATATTAAACTGATAAGAACAGATACTACACTTG	0.153
CTCCACGGAAATCTTTAGTAAAAGGCGAAAGATTT	0.145
CACAAATTATGCAGTCGAGTTTCCCACATTTGGGG	0.144
GGCCTTGCCAAAGCAAGGCCTCAAAAAATTGGGTT	0.141
ACTGATAAGAACAGATACTACACTTGATCTTAGCC	0.141
GTTCCTCCACGGAAATCTTTAGTAAAAGGCGAA	0.141
GGGGCTTTCCGGTAAAAAGCCAGAAAGCCTACTN	0.138
TGTCAGACTCTTATGTTTCACTCATAAAAGGAGTG	0.136
ATTTAATATTGTCCTCGGATAGAGGACGTATCN	0.134
TGGATCGTCGGACTGTAGAACTCTGAACGTGTNNN	0.12
ATCCGGAGTGCAATGGATAAGCCTCGCCCTGGGAA	0.119
CTCCAAAAATCCATTTAATATTGTCCTCGGATN	0.117
AAGGGCTTTCCGGTAAAAAACCAGAAAGCCTACTN	0.117
ACTCAGAGTAGTTCCTCTCCACGGAAATCTTTAGT	0.116
TGATCGTCGGACTGTAGAACTCTGAACGTGTNNNN	0.116
TAGGGGAAAGCGCGAACGCACTACCAC	0.114
AGCCAACAGCACCCGGTATTCCCAGGCGGTCTCCC	0.11
CGGGGCTTTCCGGTAAAAAGCCAGAAAGCCTACTN	0.108
GGCCTTGTCGGAACAAGGCCTCAAAAAATTAGCTT	0.107
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